



CENTRE D'ÉCOLOGIE
FONCTIONNELLE
& ÉVOLUTIVE



Adaptation to changing environments

Models of moving optimum

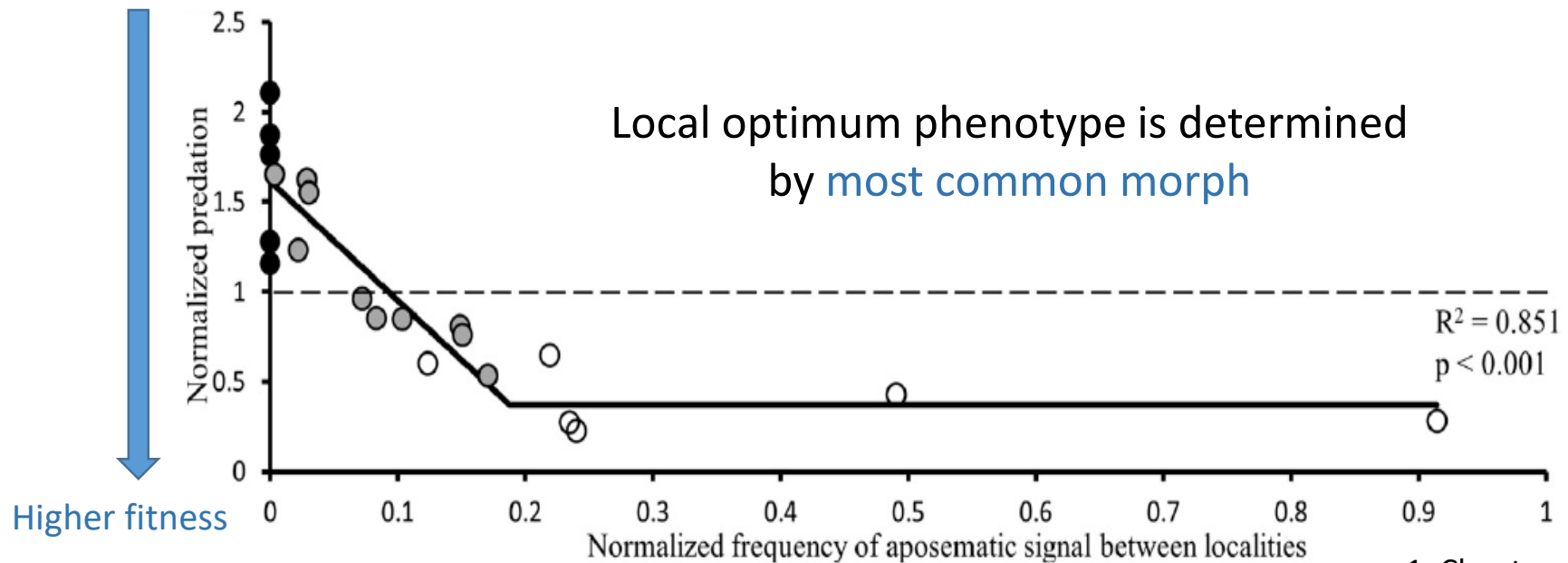
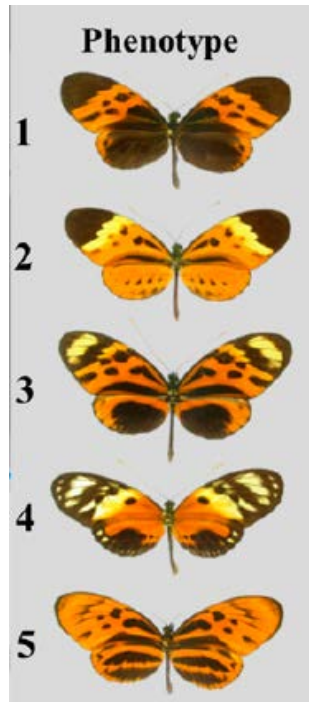
Luis-Miguel Chevin, CEFE CNRS, Montpellier, France.

Drivers of adaptive evolution

- When do organisms need to adapt by natural selection?

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 - Ecological: Resource competition causing negative frequency dependence, mimicry of warning signals causing positive FD¹



1: Chouteau et al (2016 PNAS)

Drivers of adaptive evolution

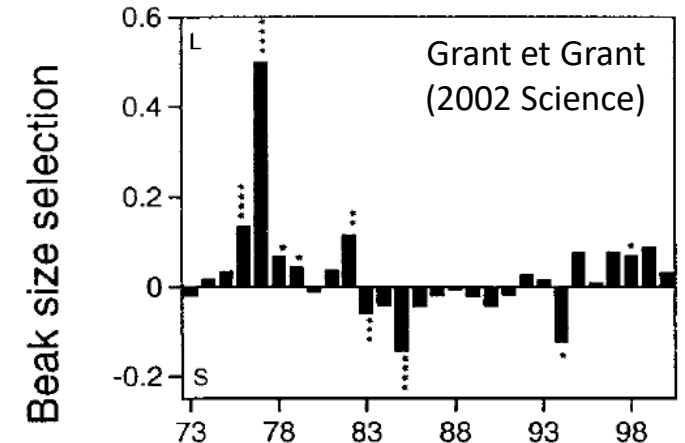
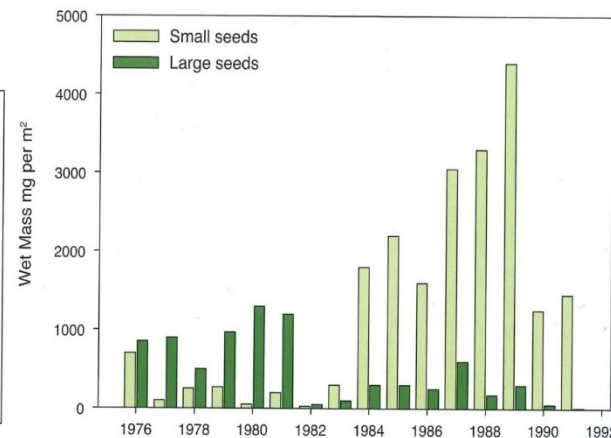
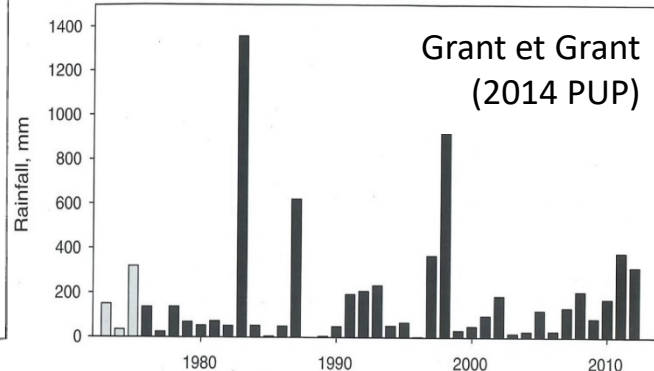
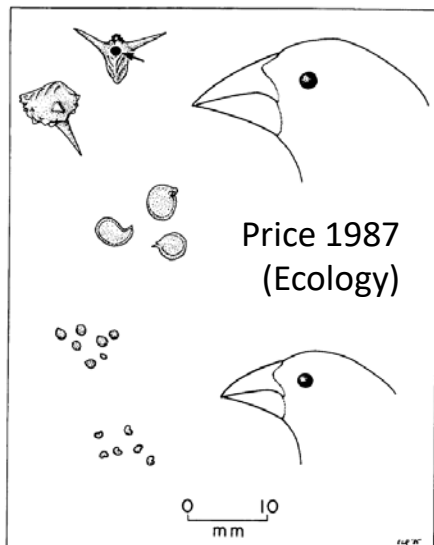
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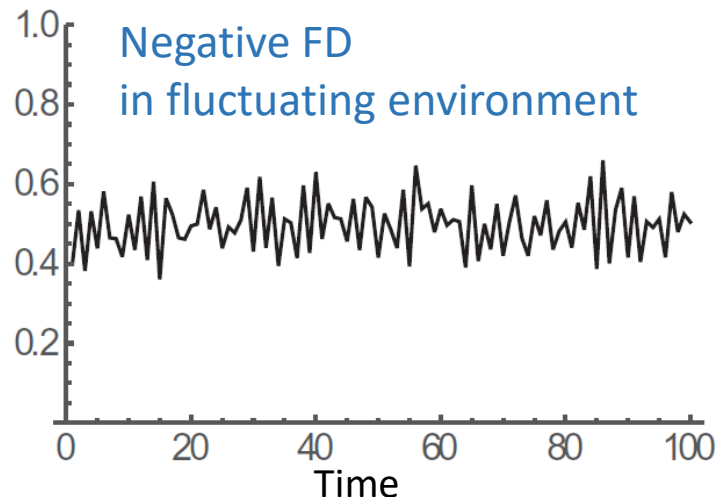
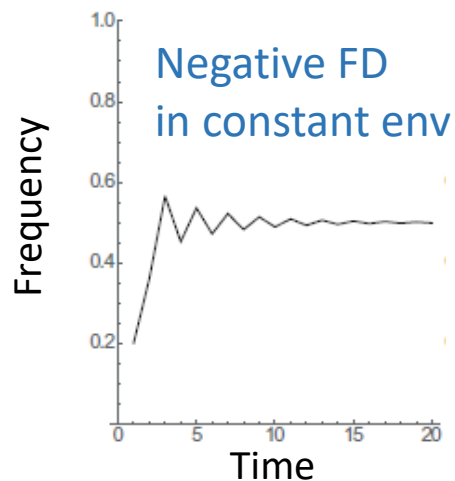
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 - **External forcing:** A **changing environment** modifies which phenotype is optimal
- Even when internal feedbacks exist, long-term evolutionary dynamics may be sustained by environmental variation³



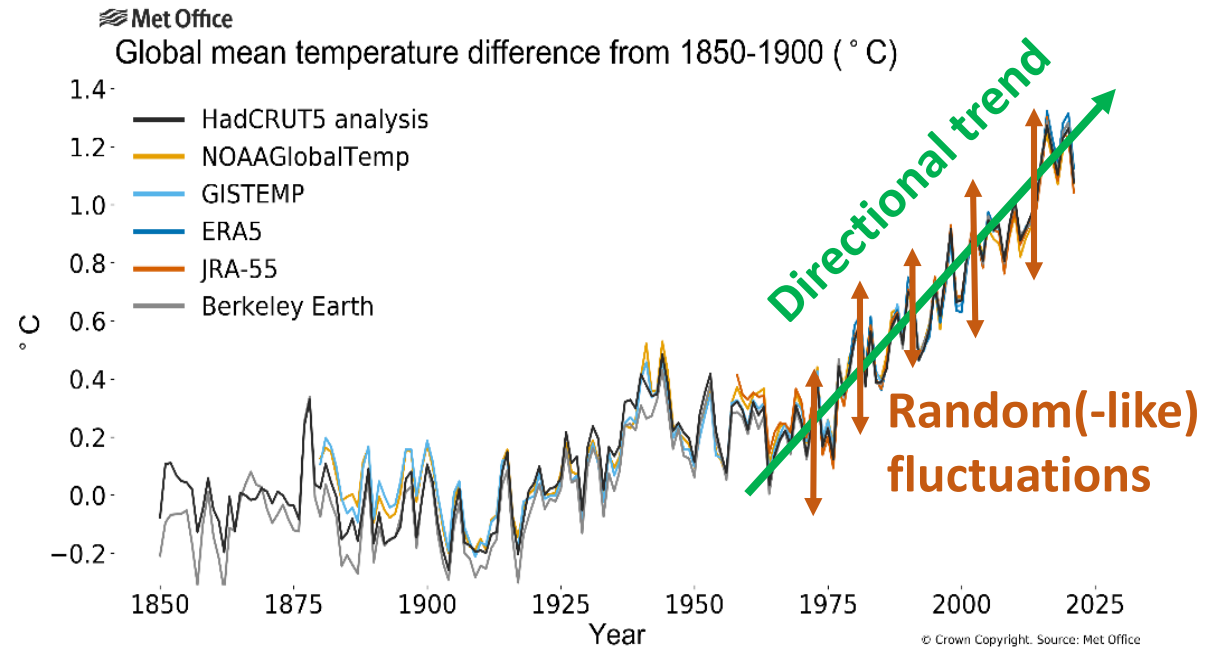
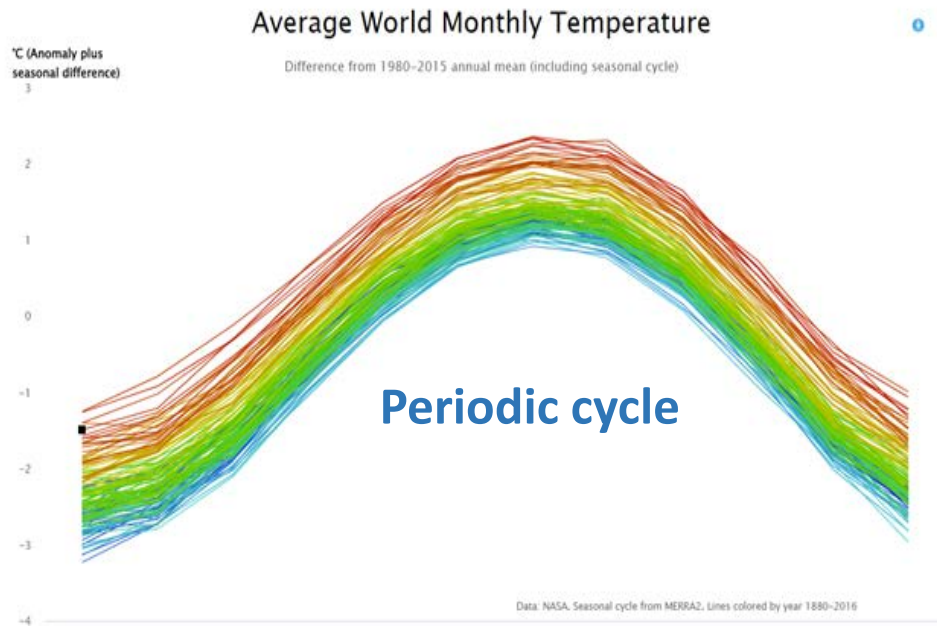
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2: Burt & Trivers (2008)

3: Chevin et al (2022 Evolution Letters)

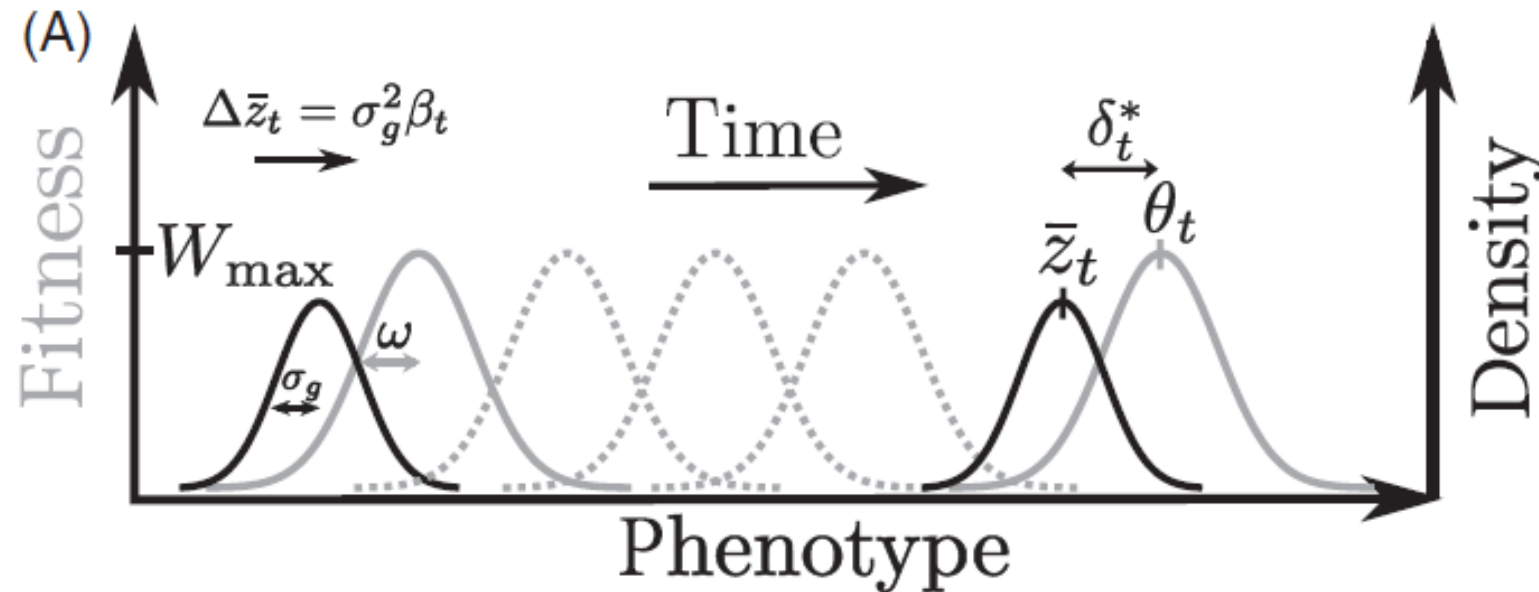
Patterns of environmental change

- Natural systems are characterized by different types of environmental changes



Evidence for moving optimum

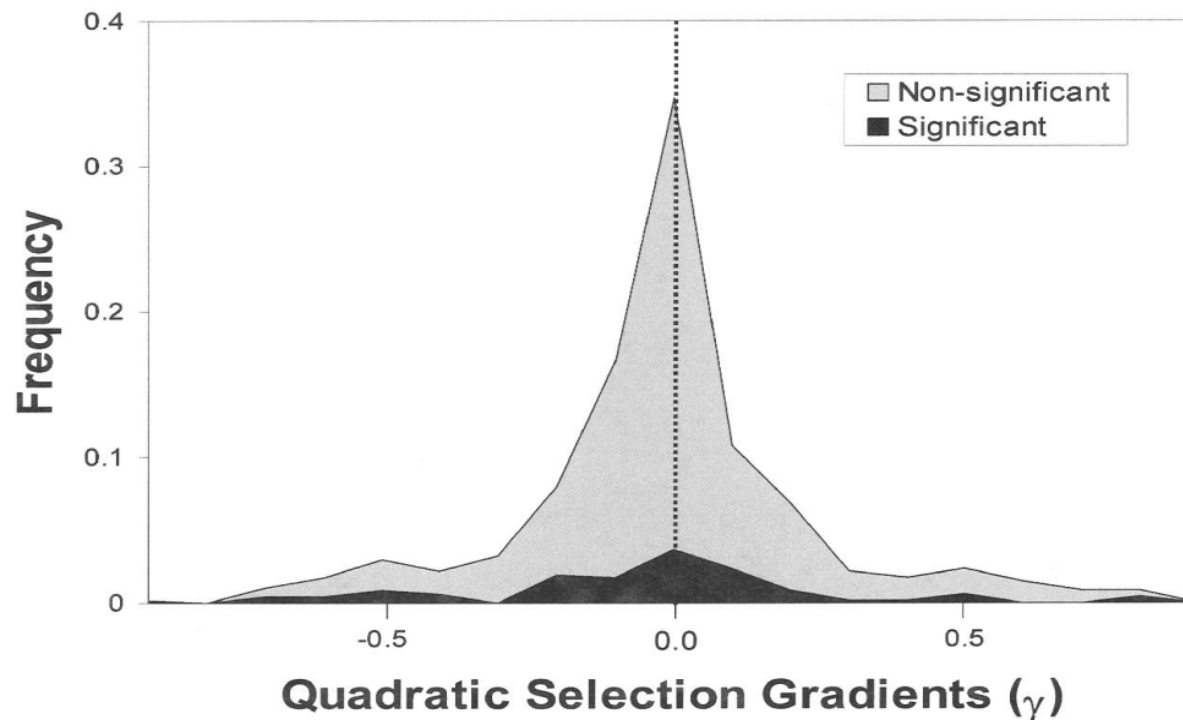
- In evolutionary theory, adaptation to changing environments often modeled as evolutionary tracking of a **moving optimum** for phenotypic traits¹.



- Makes logical sense, but **how well supported** empirically?

Evidence for moving optimum

- Direct evidence: (1) Phenotypic selection analysis.
 - Quadratic selection gradient¹ $\gamma = \text{Cov} \left[(z - \bar{z})^2, \frac{W}{\bar{W}} \right] / \sigma_z^4$
Mean curvature of fitness landscape, $\gamma < 0$ generally interpreted as stabilizing selection
 - Meta-analysis found as many $\gamma > 0$ as $\gamma < 0$, interpreted as lack of evidence for stab. selection²

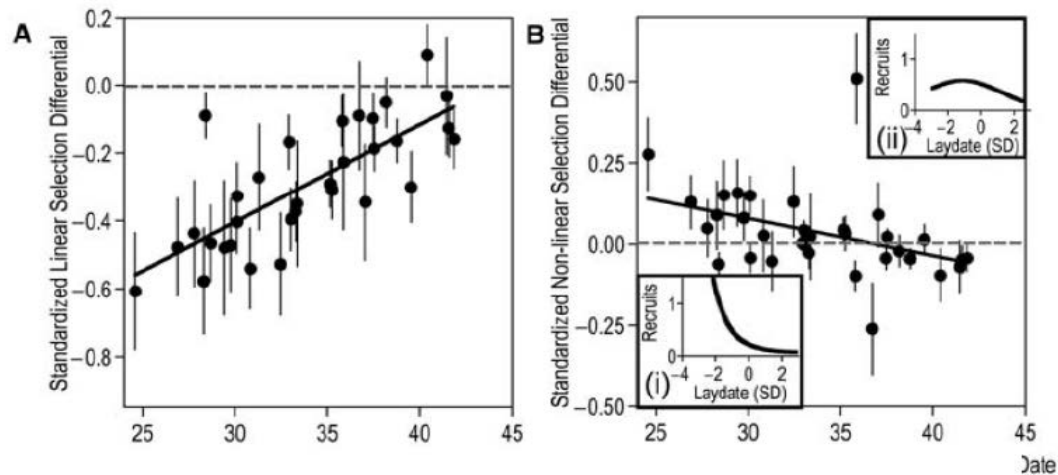


1: Lande & Arnold (1983 Evolution);
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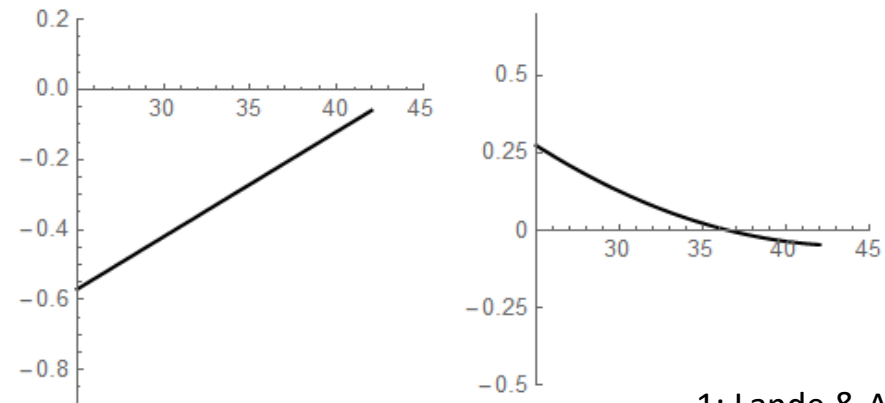
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Measured directional and quadratic gradients
(Charmantier et al 2008 Science)



Assuming a moving optimum
with constant width



1: Lande & Arnold (1983 Evolution);
2: Kingsolver et al (2001 Am Nat)

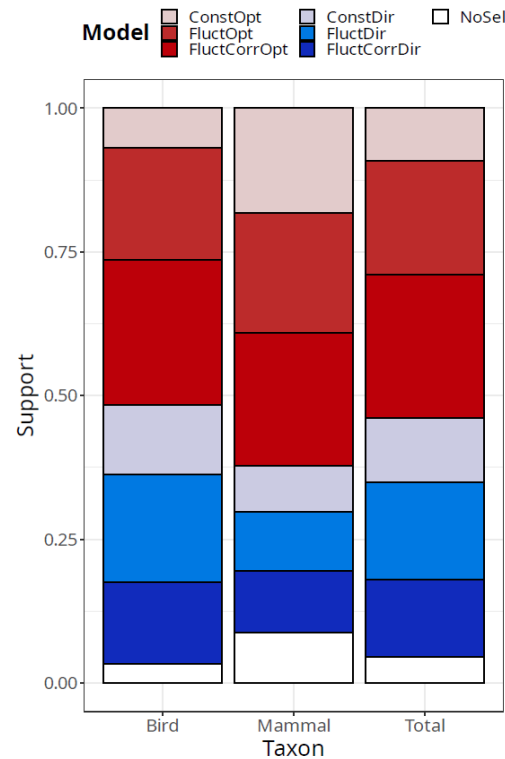
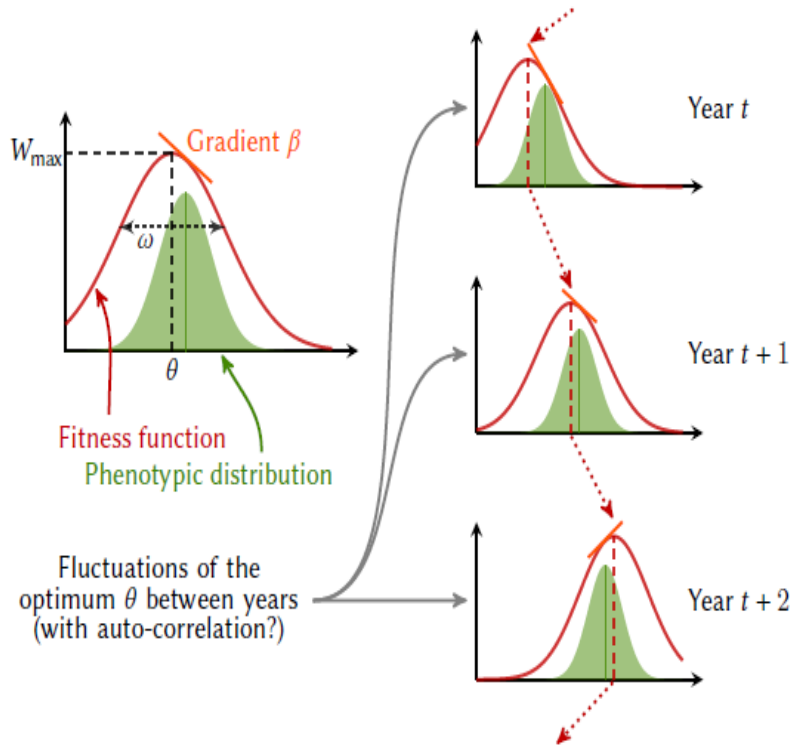
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 - Meta-analysis found as many $\gamma > 0$ as $\gamma < 0$, interpreted as lack of evidence for stab. selection²
 - But with a Gaussian peak, $\gamma > 0$ when mean phenotype sufficiently deviates from optimum
 - What appears in theoretical predictions for changes in mean and variance under selection is the strength of stabilizing selection $S = \frac{1}{V_s}$ not $\gamma = \frac{\beta^2 - S}{\sigma_z^4} = -S \frac{1 - S(z - \theta)^2}{\sigma_z^4}$
So why not estimate that directly?
 - Can be done using log link in GLM³ (eg Poisson regression, relevant for fecundity selection) or directly fitting the Gaussian peak in explicit framework (eg Stan)⁴

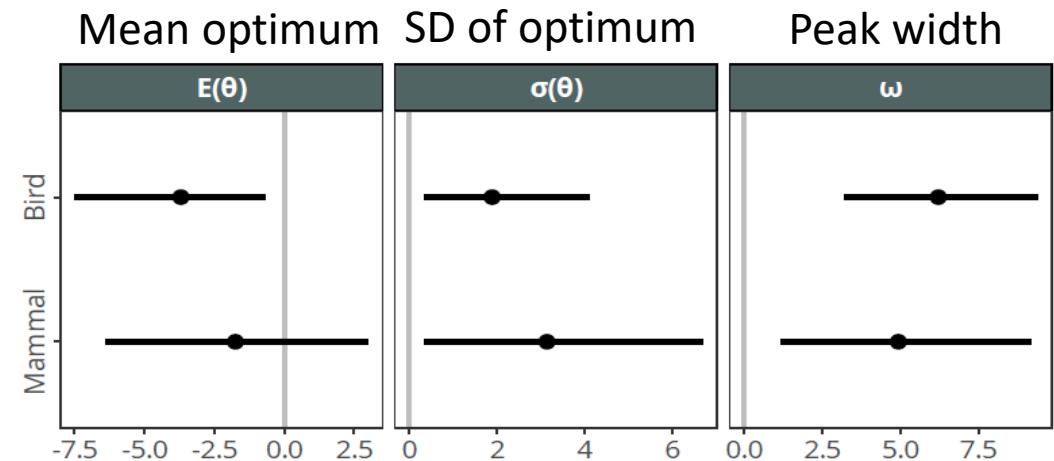
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Estimating fluctuating selection as movements of Gaussian fitness peak for breeding time across birds and mammals in the wild¹: 39 populations, 21 species, 9 to 63 yrs (average 33.2 years)



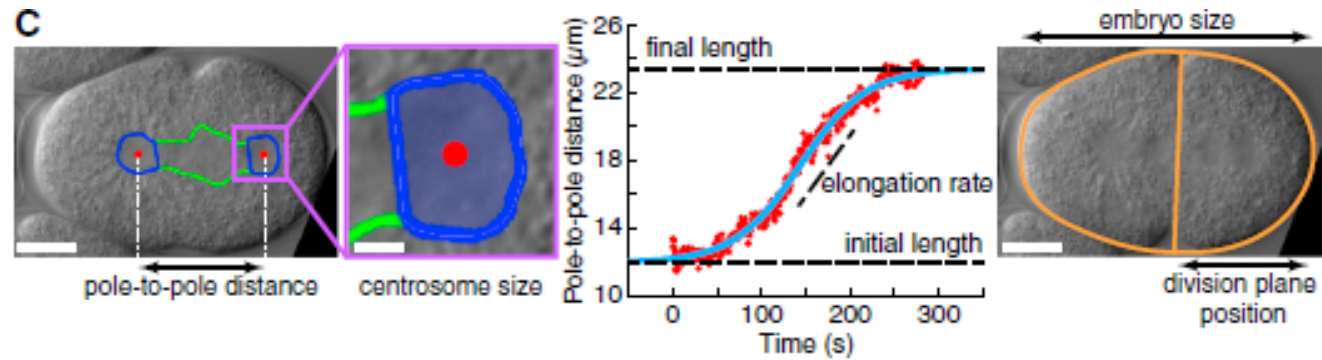
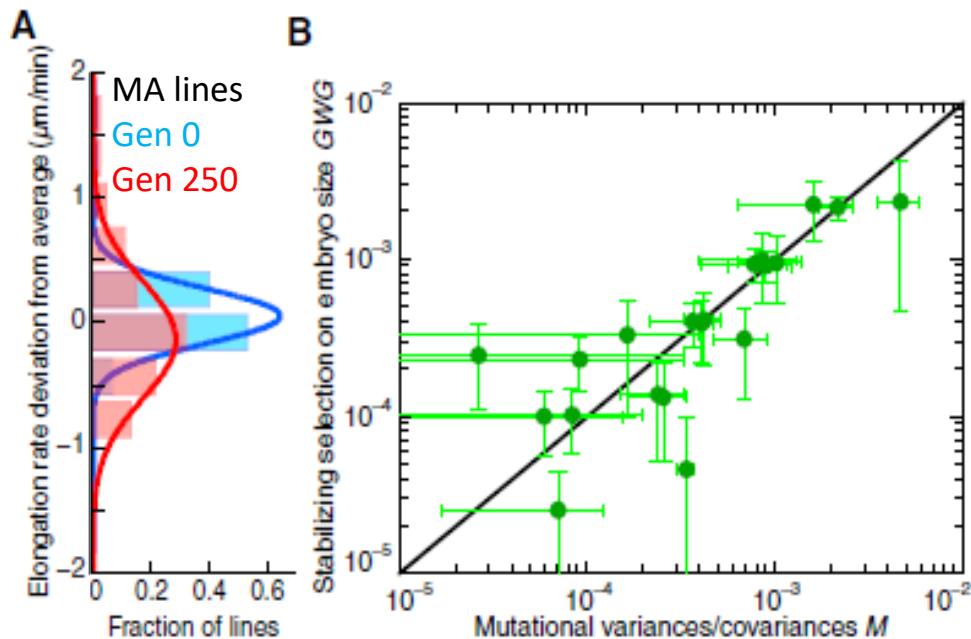
Models with optimum



1: de Villemereuil et al (2020 PNAS)

Evidence for moving optimum

- (Semi)-Direct evidence: (2) Comparison of mutational to standing genetic variance
- High-throughput measurement of many spindle traits in *C. elegans* embryos

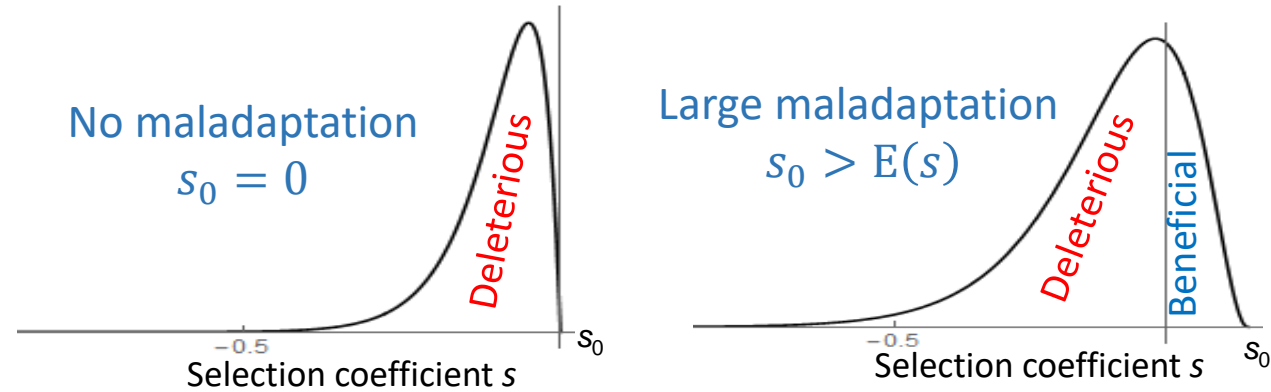
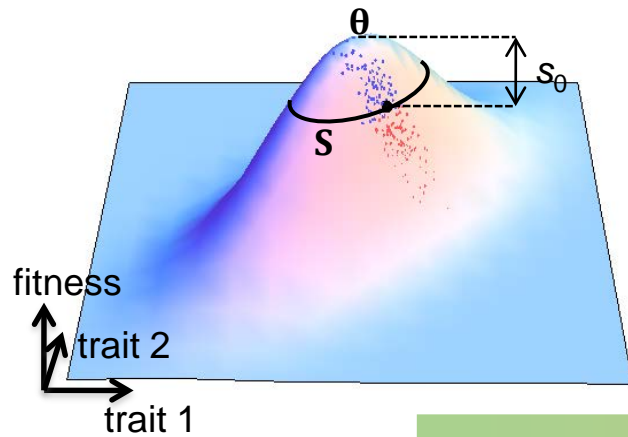


- Comparing MA lines to natural isolates: Standing variation of all traits (y-axis) well predicted by their mutational (co)variances (x-axis), but only **after accounting for stabilizing (& correlational) selection**

Evidence for moving optimum

- Indirect evidence: (3) Distribution of fitness effects across environments

Theory (Martin & Lenormand 2006 Evolution):



Reverse displaced gamma

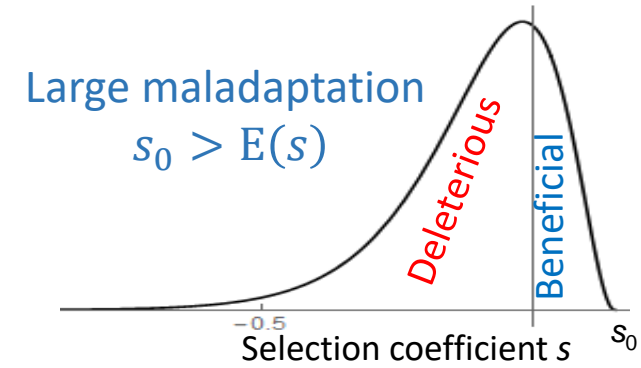
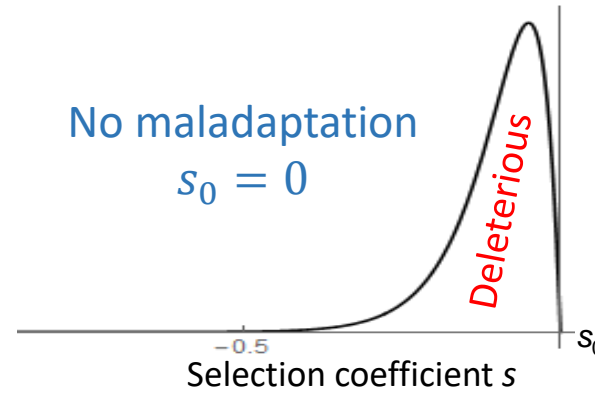
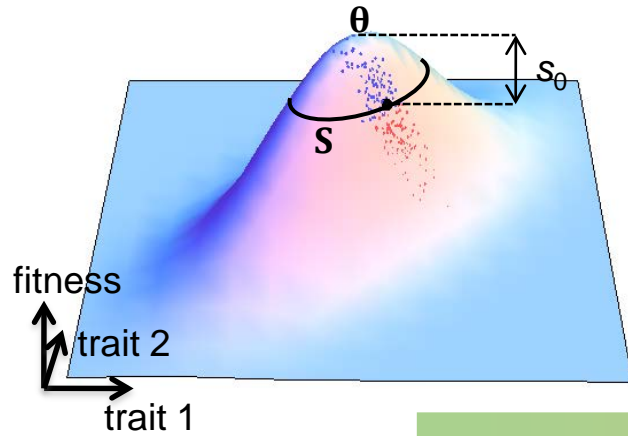
Increasing stress

- Larger variance of s
- Higher proportion of beneficials
- Same mean $E(s) = -\text{tr}(\mathbf{SM})/2$

Evidence for moving optimum

- Indirect evidence: (3) Distribution of fitness effects across environments

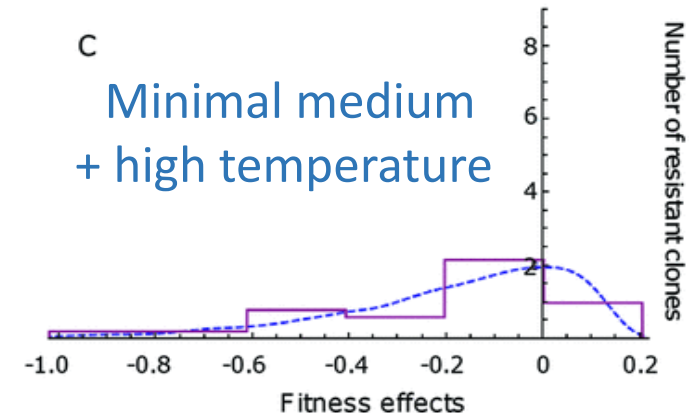
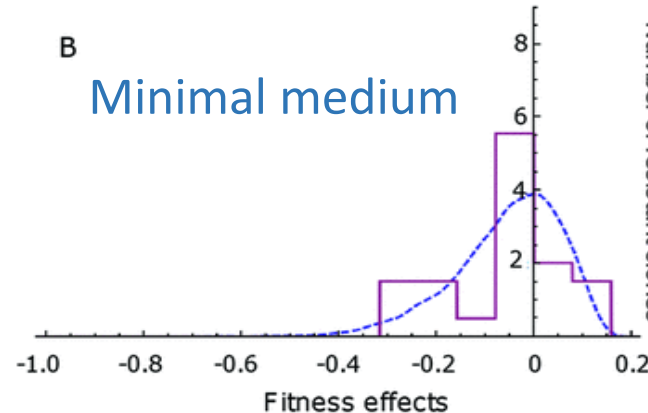
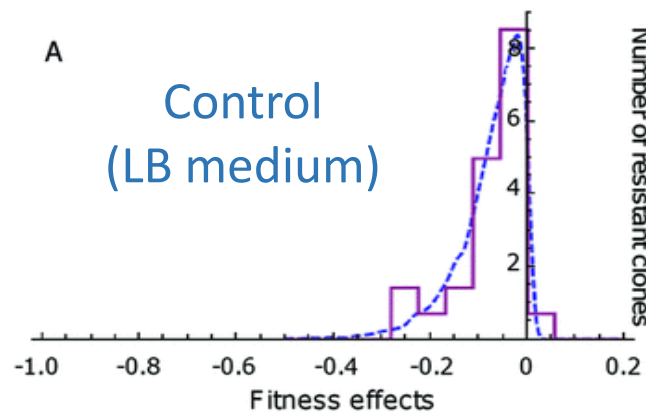
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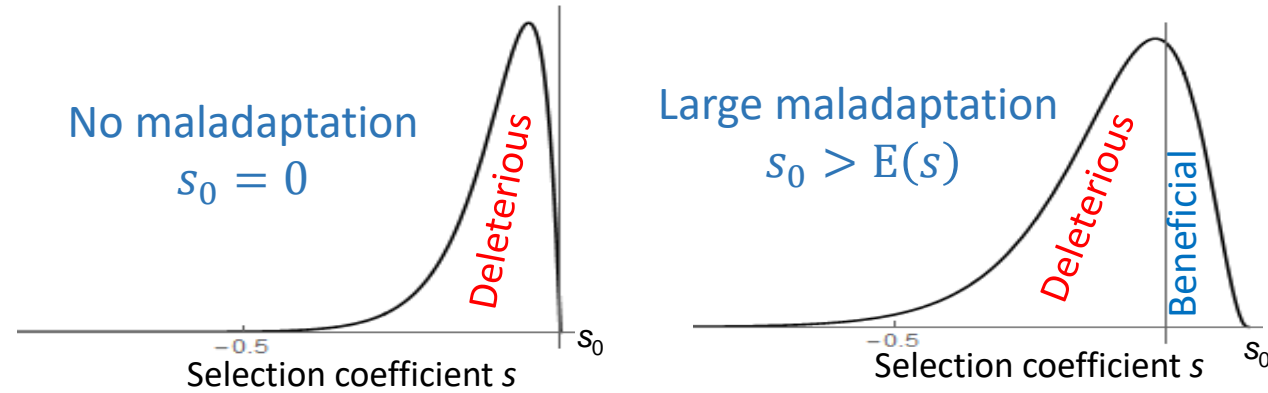
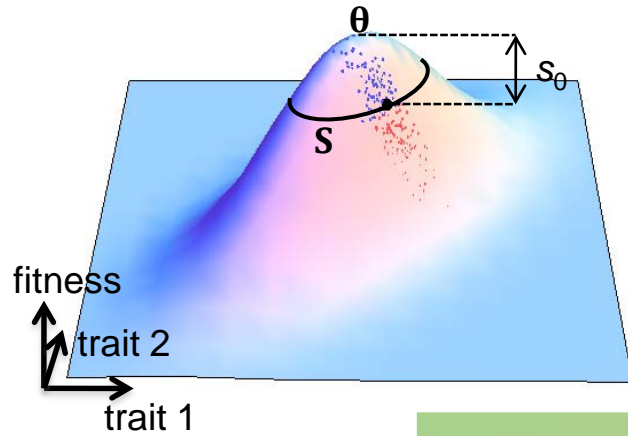
Experiment 1 (Trindade et al 2012 Evolution): Confirmed + estimated all parameters from DFE predicted by FGM



Evidence for moving optimum

- Indirect evidence: (3) Distribution of fitness effects across environments

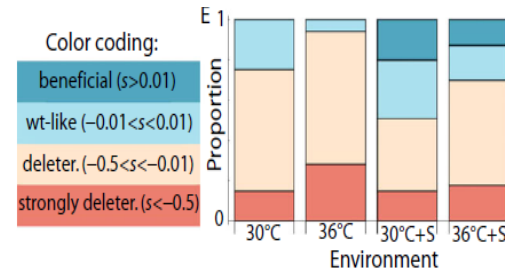
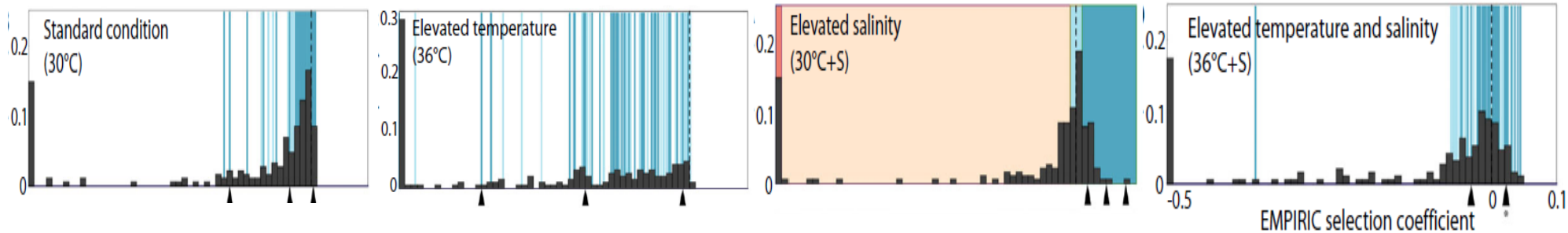
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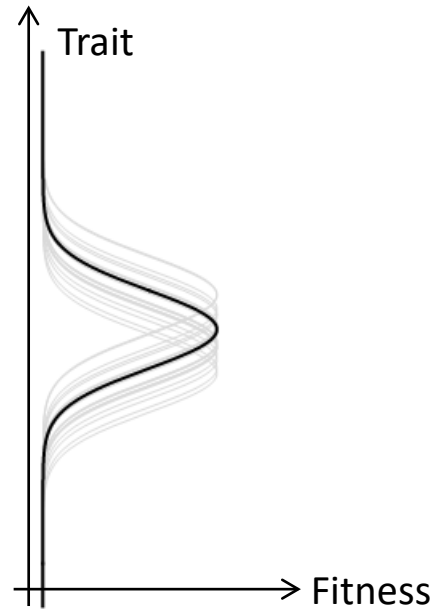
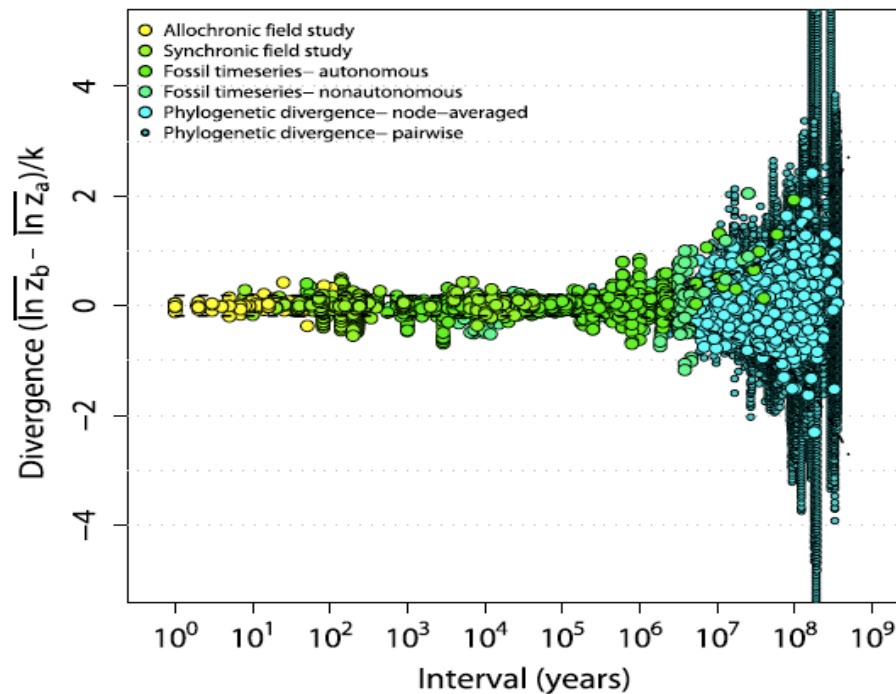
Increasing stress

Experiment 2 (Hietpas et al 2013 Evolution): Confirmed + estimated s_0 from FGM



Evidence for moving optimum

- Indirect evidence: (4) Paradox of stasis
 - **Rates of evolution** across timescales¹:
Fast over short times, stasis in the long run, then burst after 10^6 yrs.
 - Compared to evolutionary QG theory¹: Not consistent with drift, but consistent with **stationary fluctuations of an optimum phenotype + rare strong shifts**.

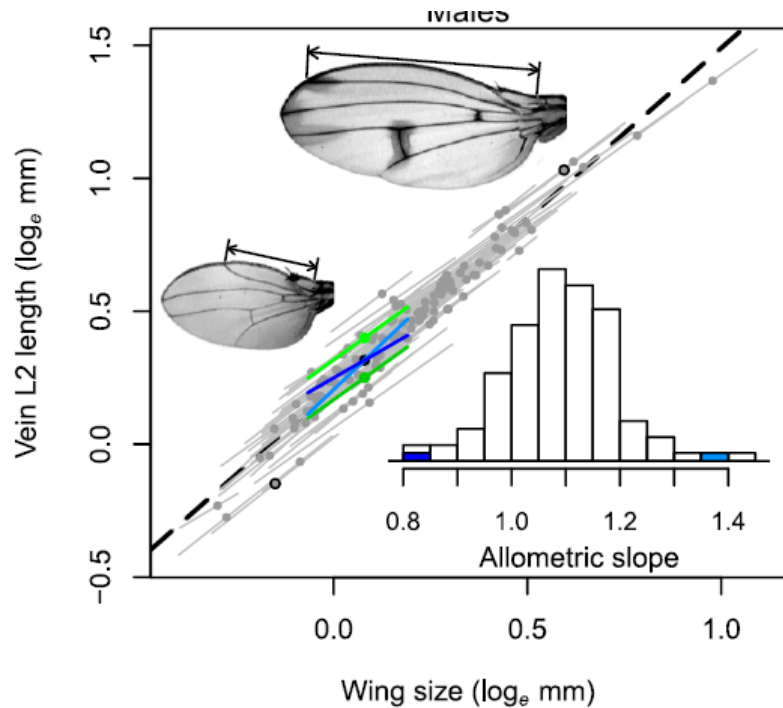


1: Estes & Arnold (2007 Am Nat);
Uyeda et al (2012 PNAS)

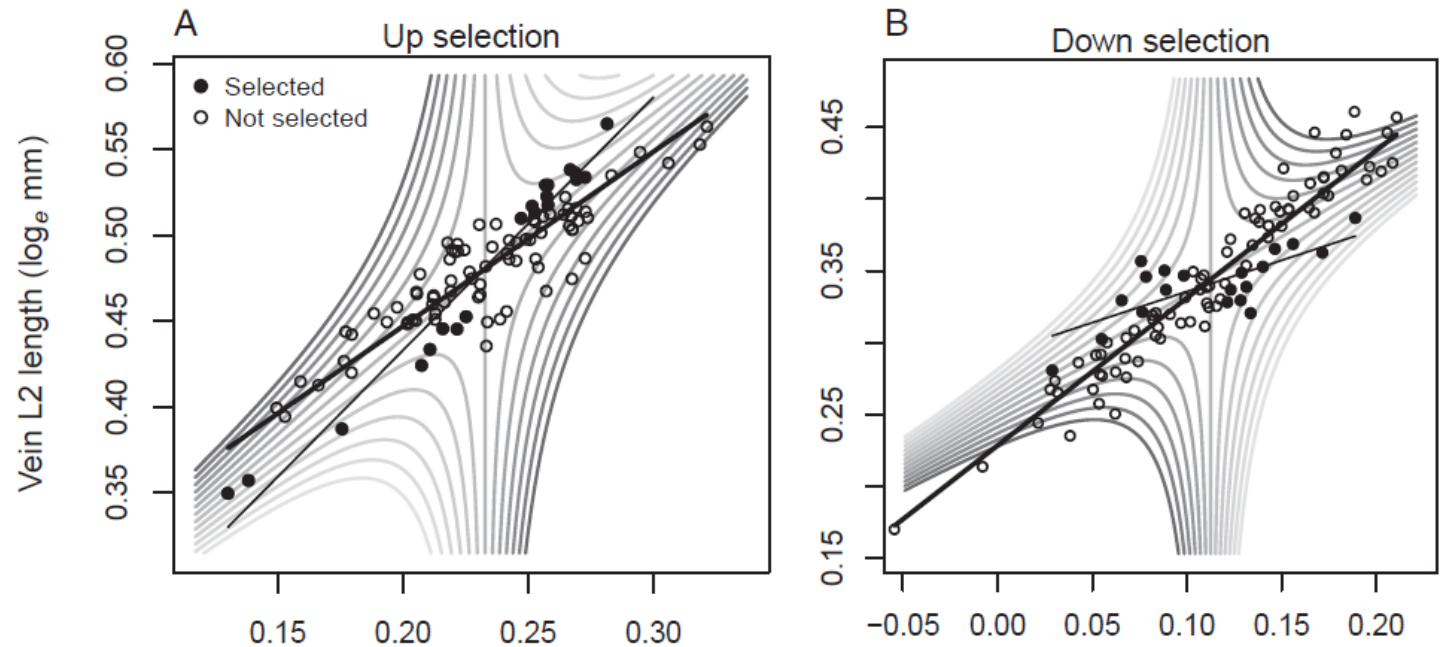
Evidence for moving optimum

- Indirect evidence: (5) Reversion of selection responses

Strongly conserved allometric relationship (111 species)



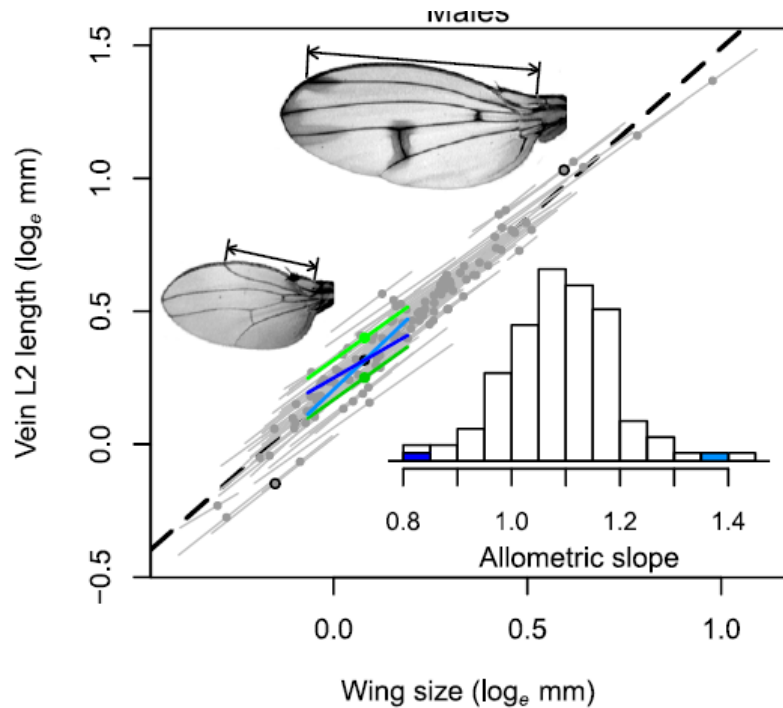
+ and - selection on allometric intercept and slope
(using correlational disruptive selection)



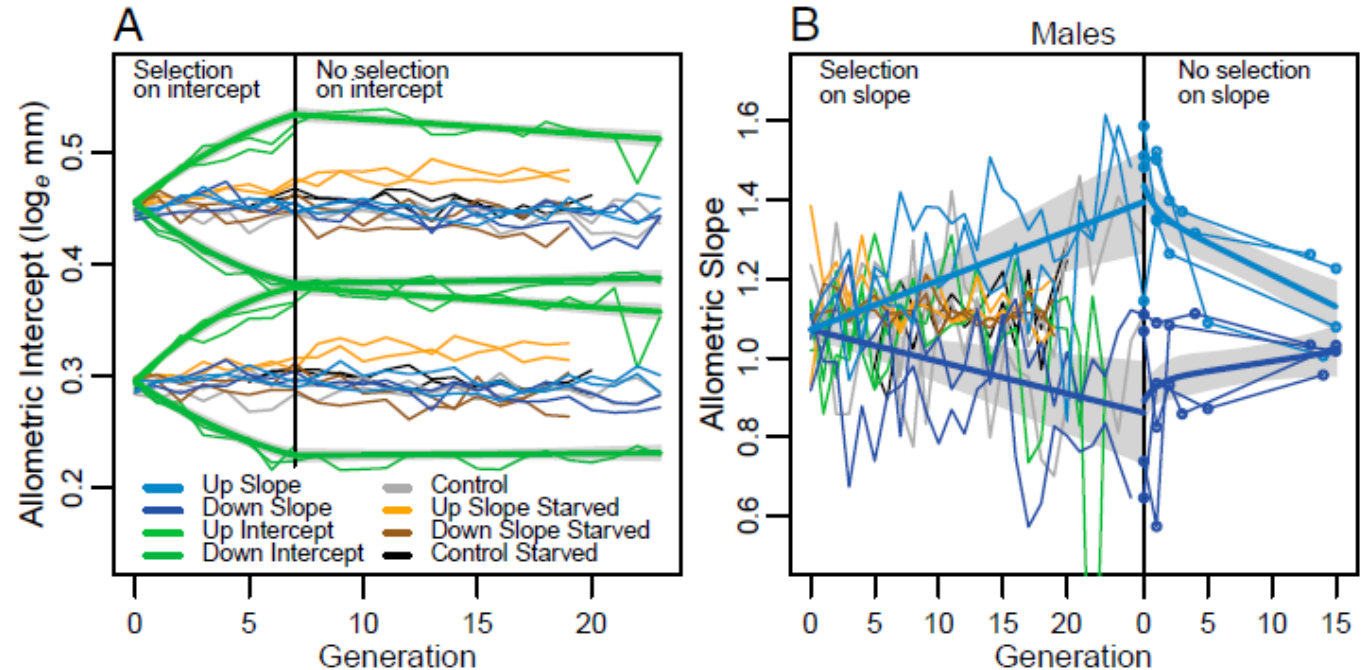
Evidence for moving optimum

- Indirect evidence: (5) Reversion of selection responses

Strongly conserved allometric relationship (111 species)



Rapid response, but **reverts after selection is relaxed**

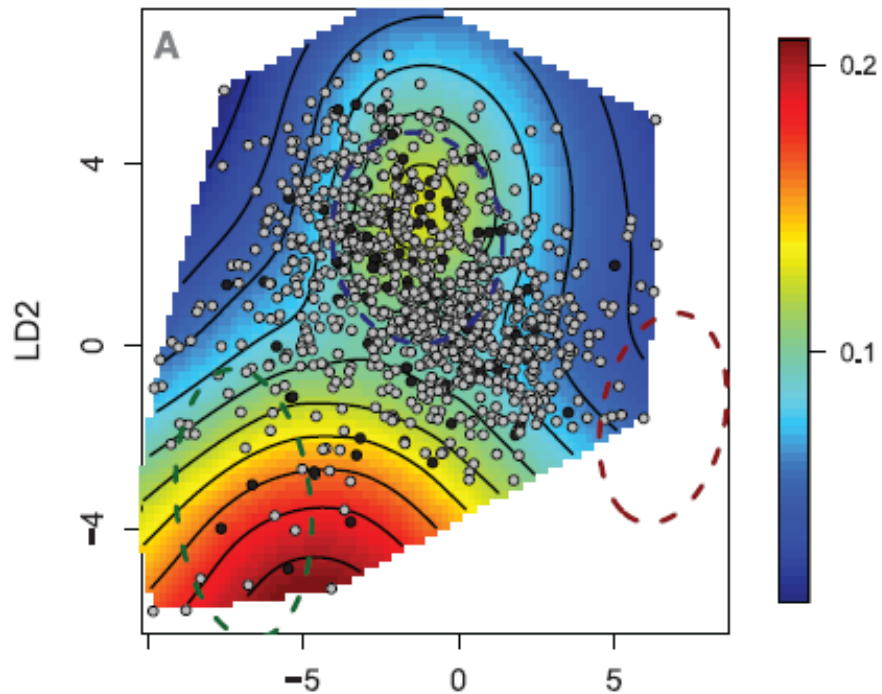


Could be due to break up of LD,
but here could only account for 15-20% of response

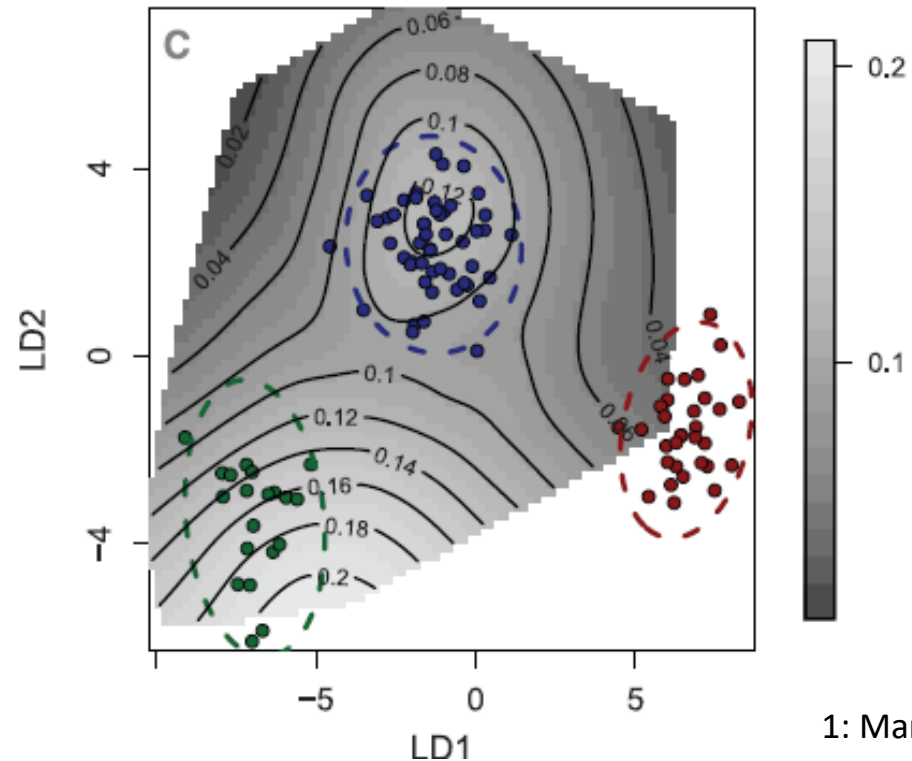
Evidence for moving optimum

- Indirect evidence: (6) Ecological speciation
 - F2 hybrids between forming species, in nascent adaptive radiation in *pupfishes*

Recombinants reveal
underlying fitness landscape



Original species **lie on**
fitness peaks



Evidence for moving optimum

Combining these converging lines of evidence:

- (1) (Fluctuating) phenotypic selection analysis
 - (2) Comparison of mutational to standing genetic variance
 - (3) Distribution of fitness effects across environments
 - (4) Paradox of stasis
 - (5) Reversion of selection responses
 - (6) Ecological speciation
- + others (eg fitness cost of artificial selection in natural environments¹, ...)

Stabilizing selection seems overall well supported.

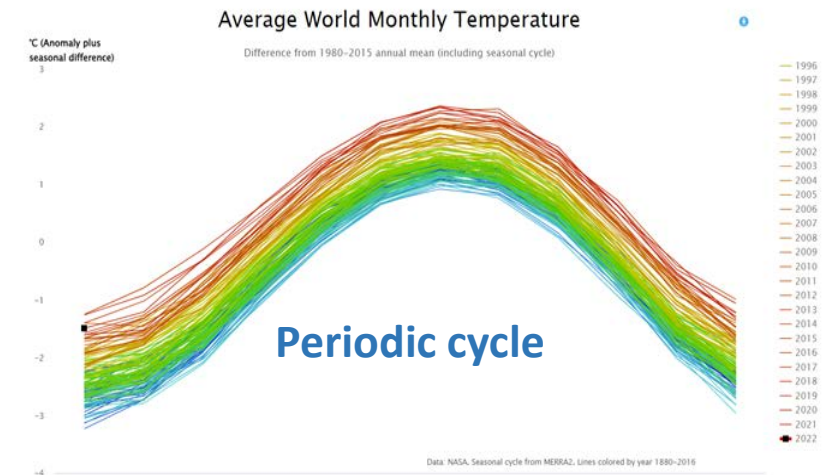
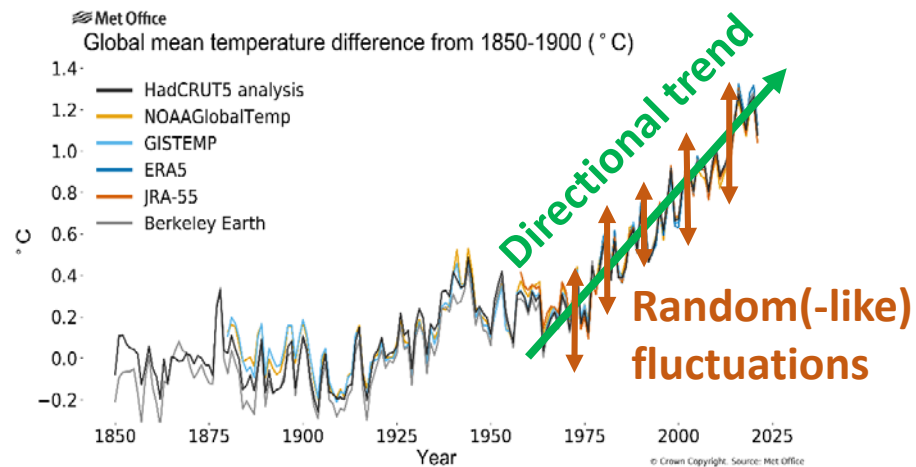
However specific shape of fitness peak may deviate from that usually assumed, especially far from optimum.

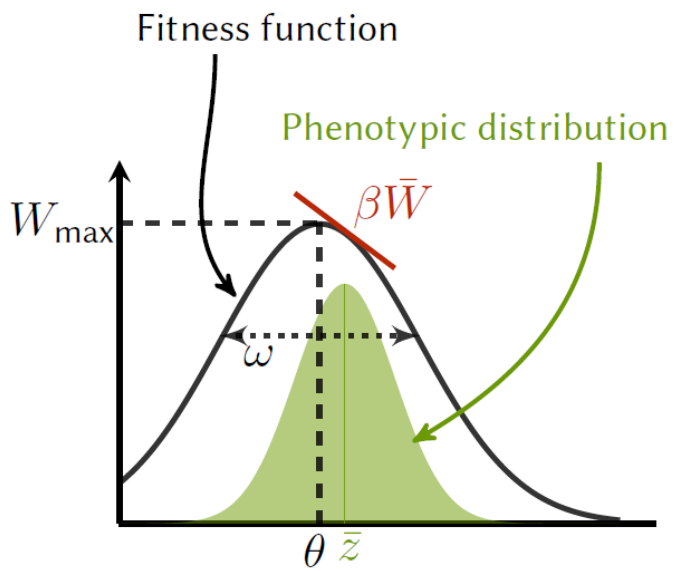
Goal and overview of the lecture

How can **moving optimum models** help understand and predict **adaptation to changing environments?**

Foreword: Moving optimum model

1. Adaptation to **directional environmental change**
2. Adaptation to **cycling environments**
3. Adaptation to **stochastic environmental fluctuations**





Gaussian fitness peak

- Any phenotype-fitness map with an optimum can be approximated as Gaussian (2nd order Taylor series on log scale)

$$W(z) = W_{\max} \exp\left(-\frac{(z - \theta)^2}{2 \omega^2}\right)$$

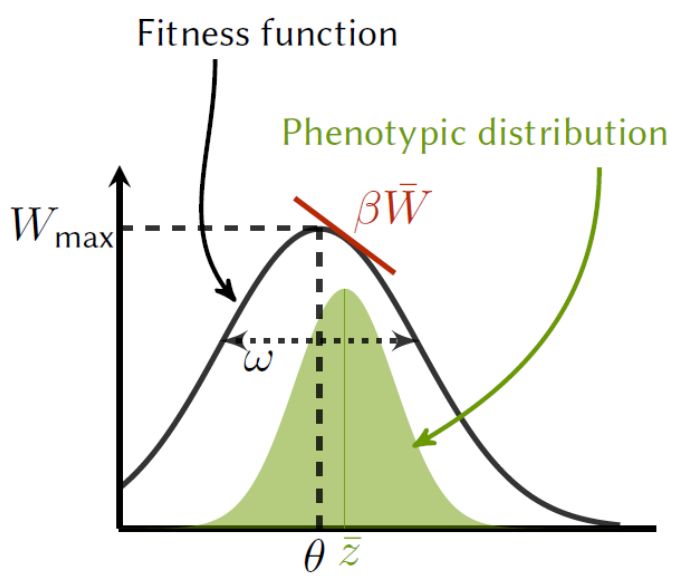
Optimum phenotype θ , width of fitness peak ω

- If trait z is a normally distributed (polygenic+residual variation), then **mean fitness is also Gaussian with respect to mean phenotype**

$$\bar{W} = \int_{-\infty}^{\infty} p(z)W(z)dz \propto \exp\left(-\frac{S(\bar{z} - \theta)^2}{2}\right)$$

$$S = \frac{1}{V_S} = \frac{1}{(\omega^2 + \sigma_z^2)}$$

is the strength of stabilizing selection



Gaussian fitness peak

- The **mean mismatch with optimum $x = \bar{z} - \theta$** drives evolutionary dynamics

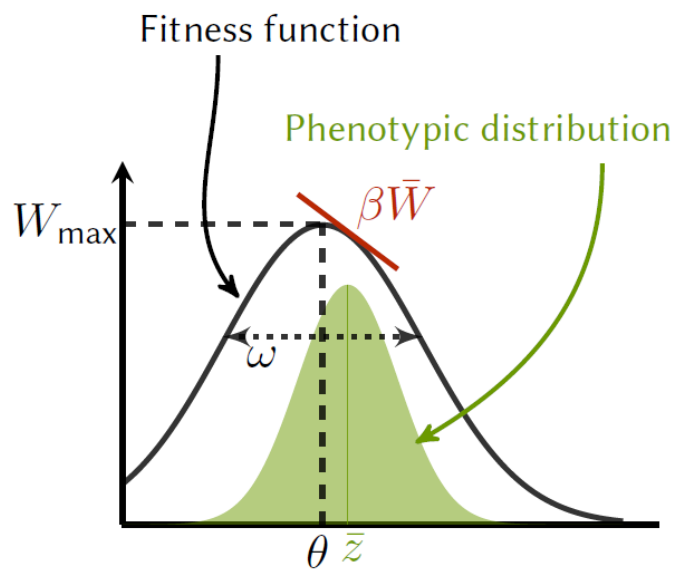
➤ Change in frequency p of a mutation with effect α on the trait in haploid population:

$$\Delta \ln \left(\frac{p}{1-p} \right) = \ln \left(\frac{W(\bar{z} + \alpha)}{W(\bar{z})} \right) = -\frac{S}{2} [(\mathbf{x} + \alpha)^2 - \mathbf{x}^2] = -\frac{S}{2} [\alpha^2 + 2\alpha\mathbf{x}]$$

→ **linear in mismatch.**

➤ For a normally distributed trait, directional gradient (selection on mean phenotype) is¹:

$$\beta = \frac{\partial \ln \bar{W}}{\partial \bar{z}} = -S\mathbf{x} \quad \rightarrow \text{linear in mismatch}$$

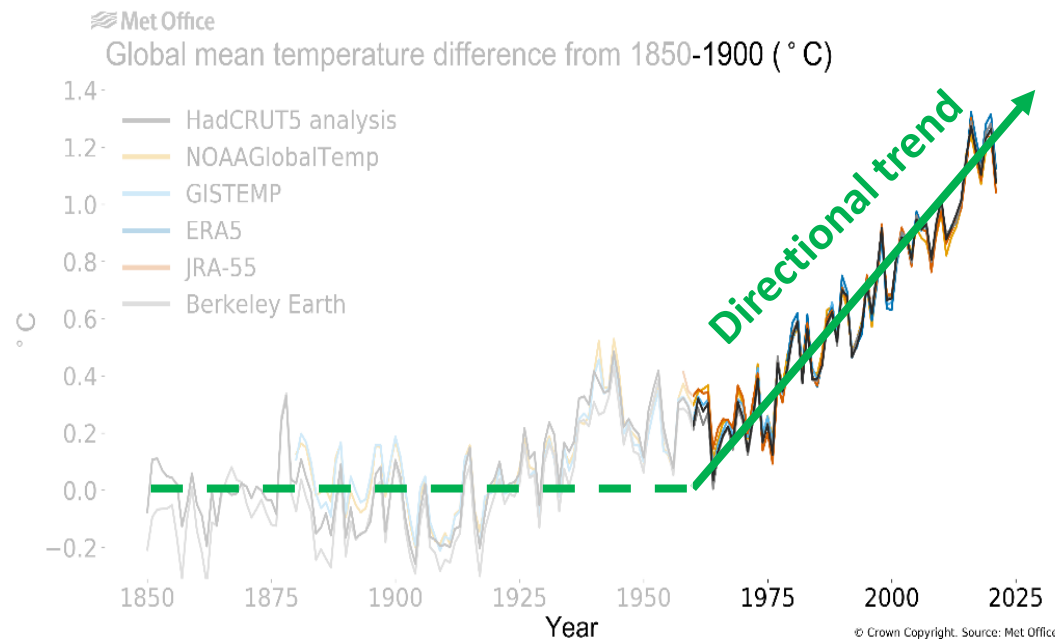


Gaussian fitness peak

- Response to selection also depends on additive genetic variance:
 $\Delta \bar{z} = G\beta = -GSx \rightarrow$ Linear restoring force reducing deviations from optimum x
- For a given deviation x , faster evolution if larger **adaptive potential SG** ,
 i.e. narrow fitness peak x large additive genetic variance.
- When genetic variance can be approximated as constant, simple dynamical system allowing analytical progress under relevant types of environmental change.

Directional environmental change

- Abrupt directional change (environmental shift) addressed in previous lectures (in adaptive walk¹ and polygenic² regimes)
- More gradual tendencies (e.g. global warming) can be modeled as **steady change at constant speed v** , preceded by a constant environment



Directional environmental change

- Low mutation regime (origin-fixation process):
Adaptive walk by sequential fixation in otherwise monomorphic population.
- Reminder: Under sudden shift, populations start far from optimum
→ large effect mutations can fix in early steps¹

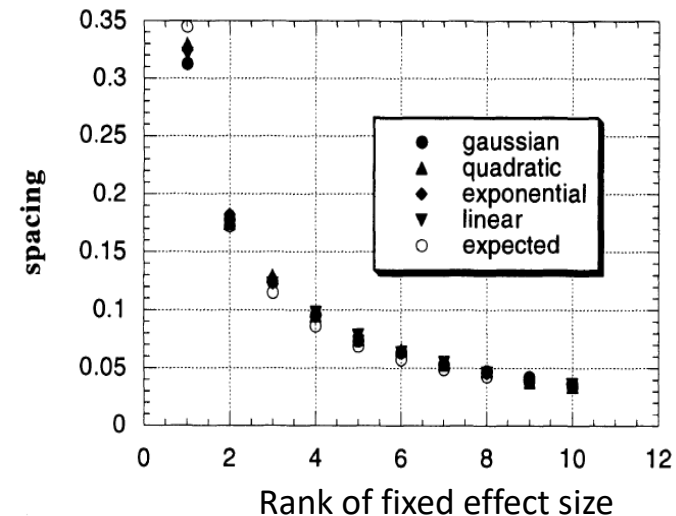
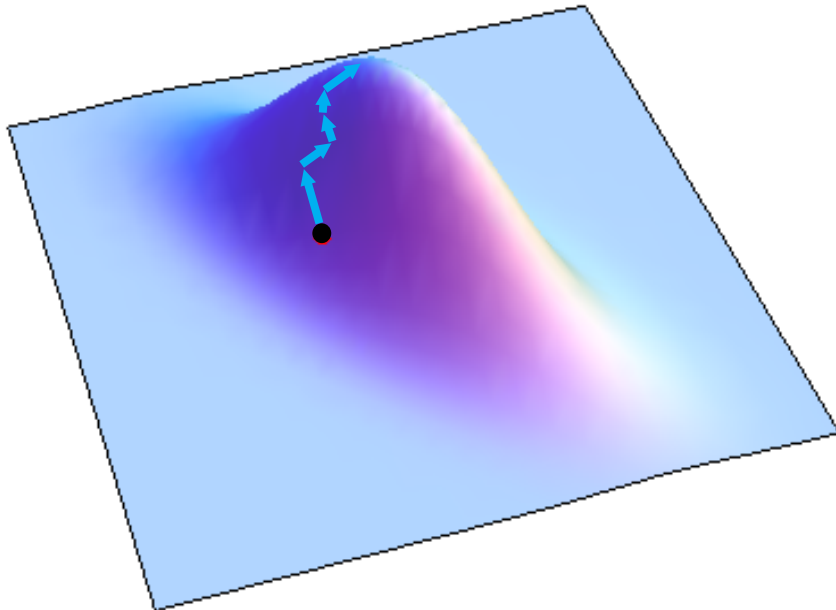
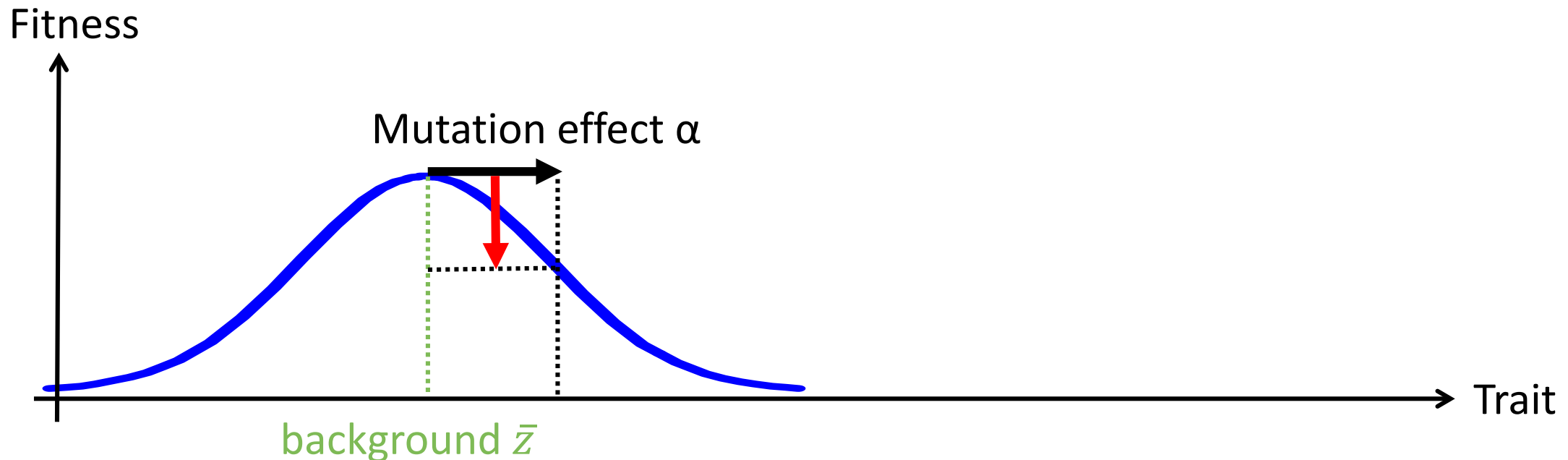


FIG. 8. Spacings between largest and next-to-largest, etc. factors fixed during adaptation.

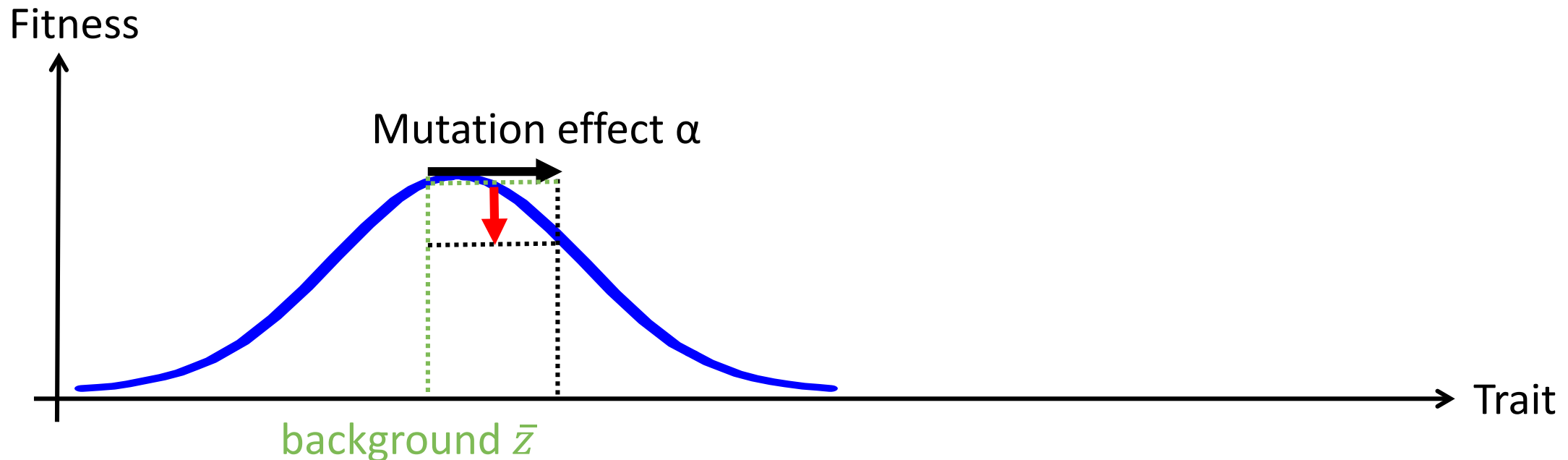
Directional environmental change

- Low mutation regime (origin-fixation process):
Adaptive walk by sequential fixation in otherwise monomorphic population.
- Under **gradual trend**, **selection coefficients change as optimum moves**
→ A mutation first needs to **become beneficial**



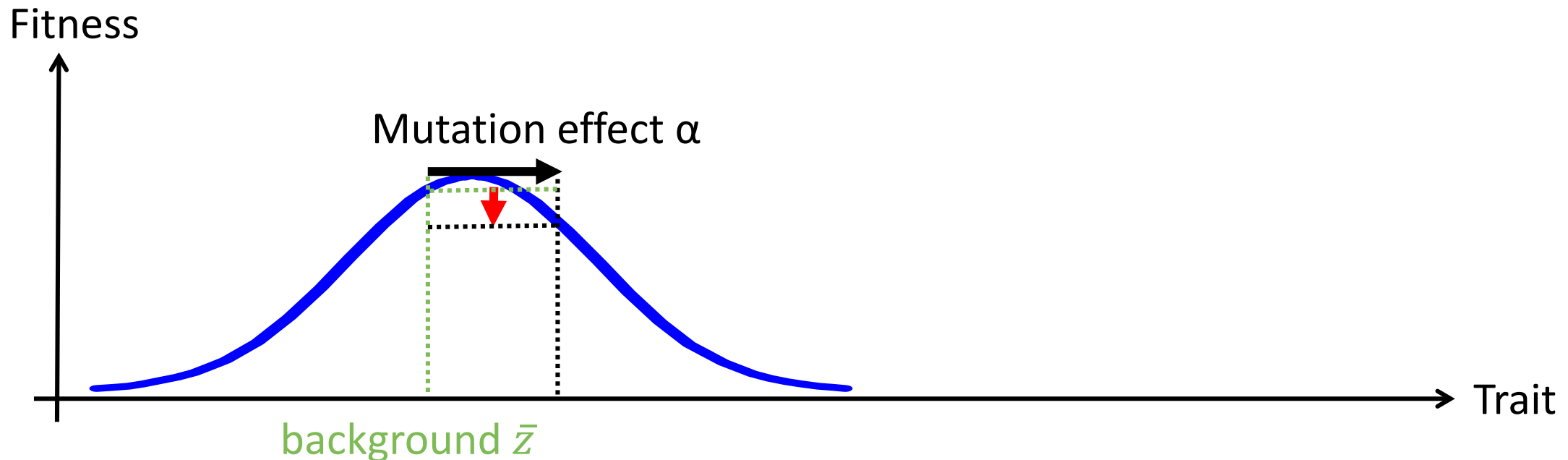
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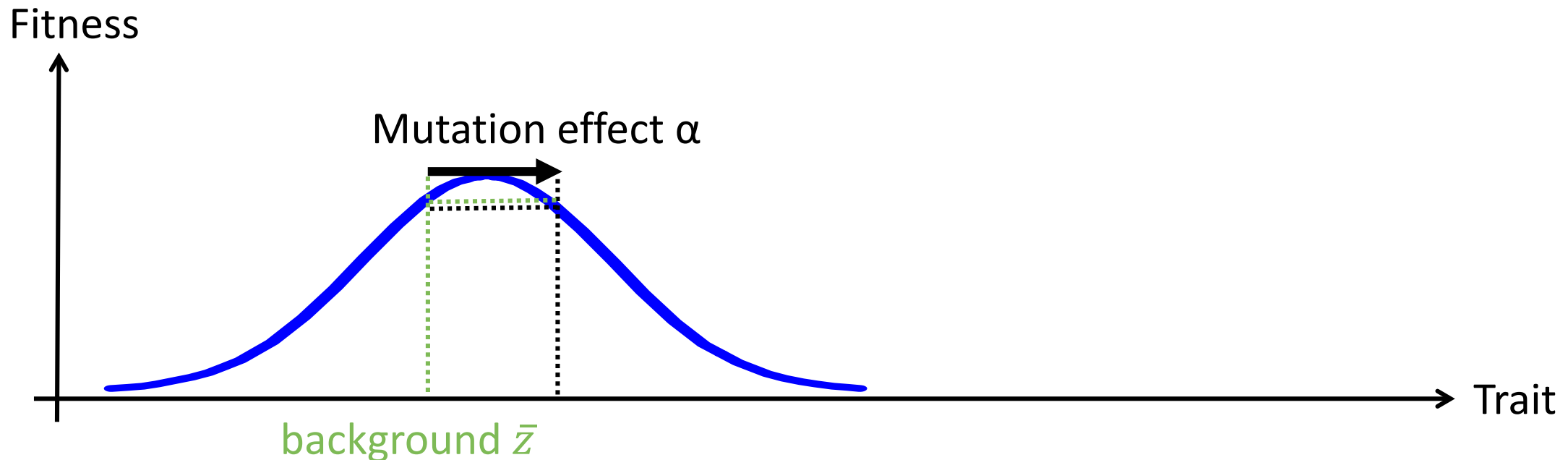
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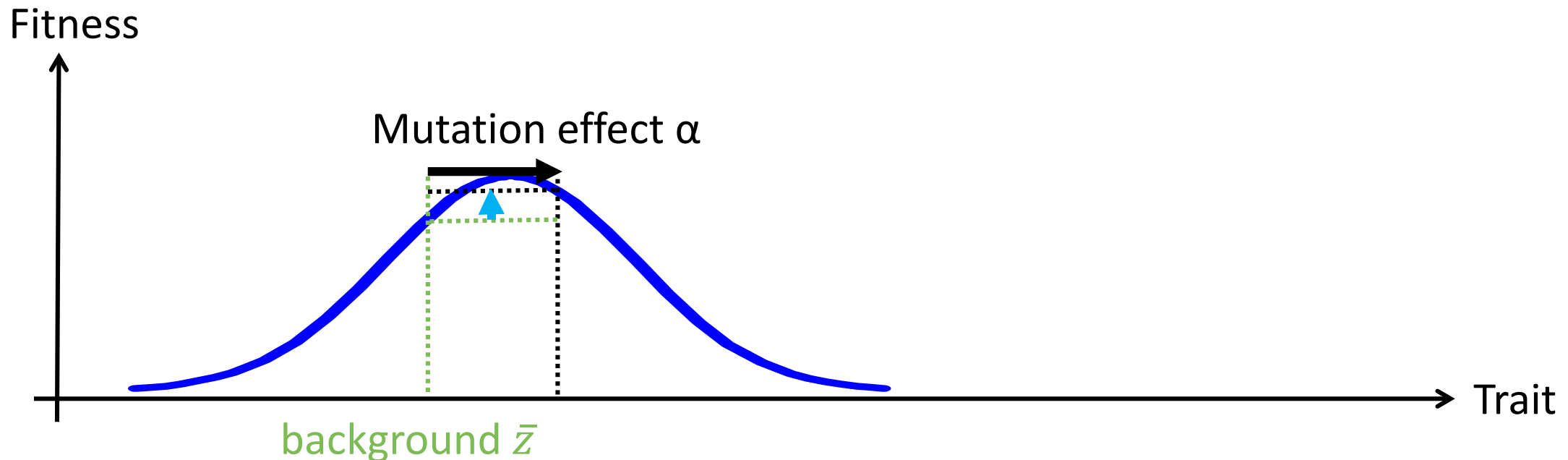
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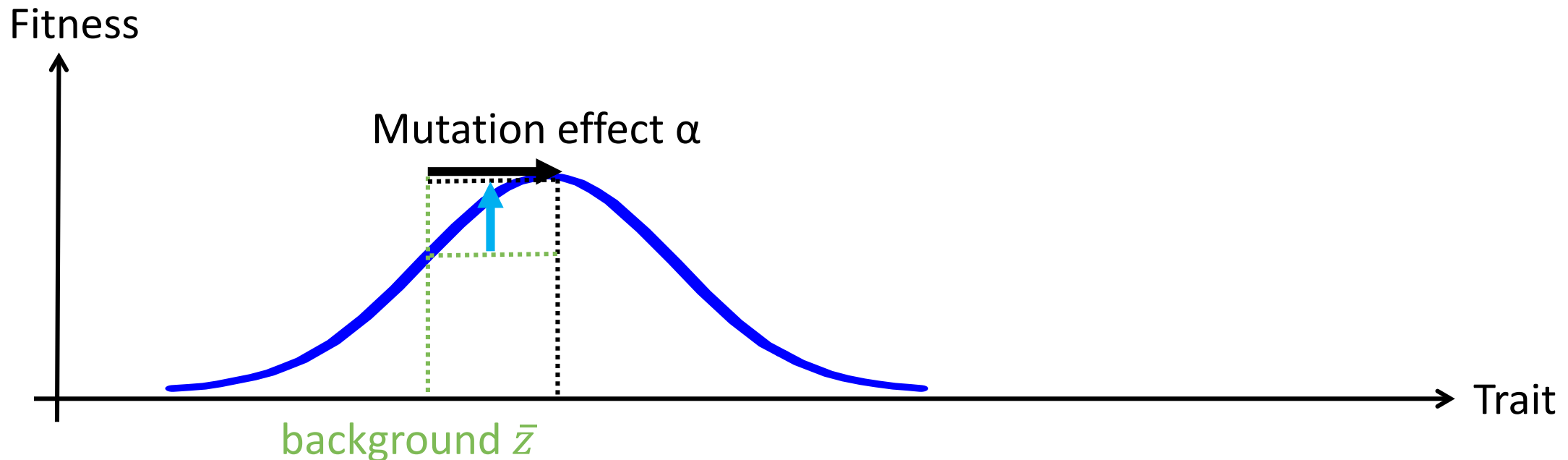
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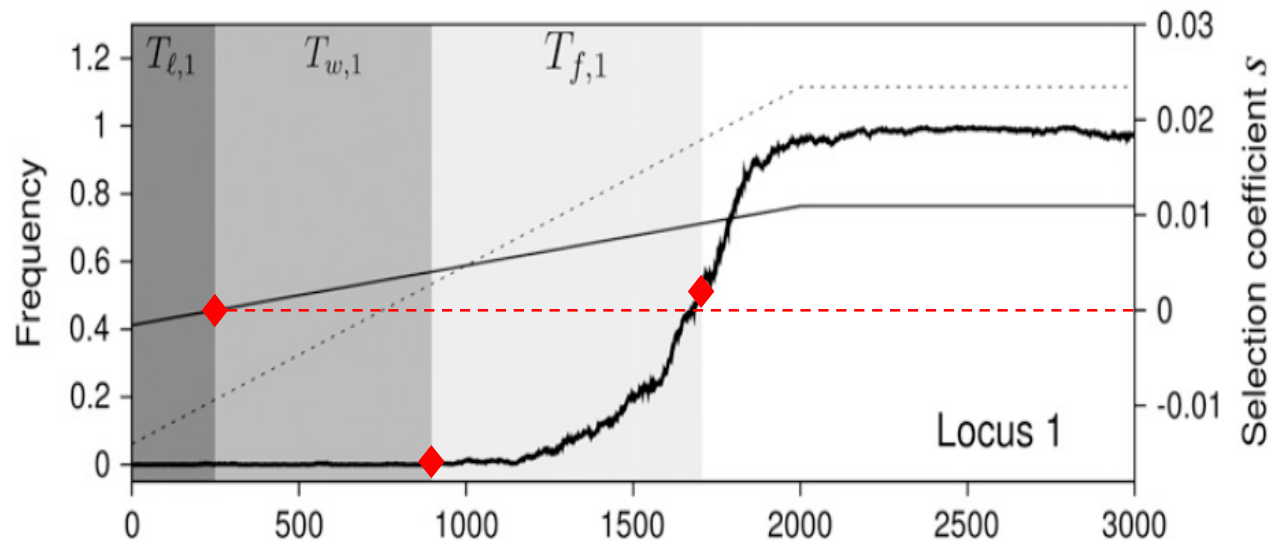
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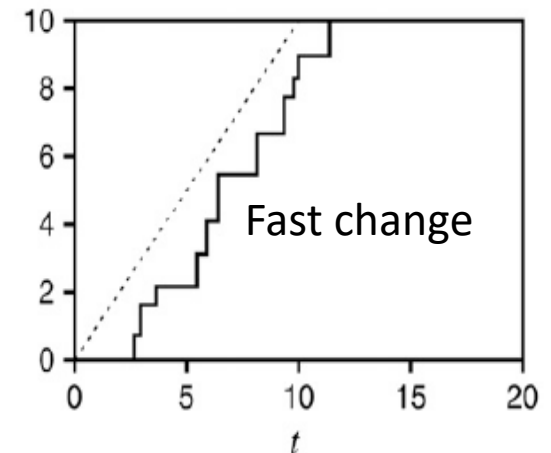
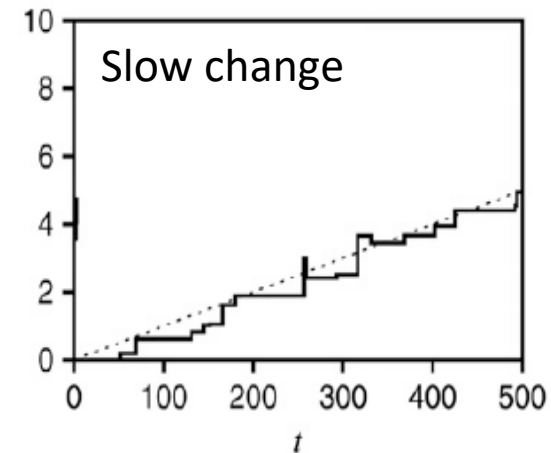
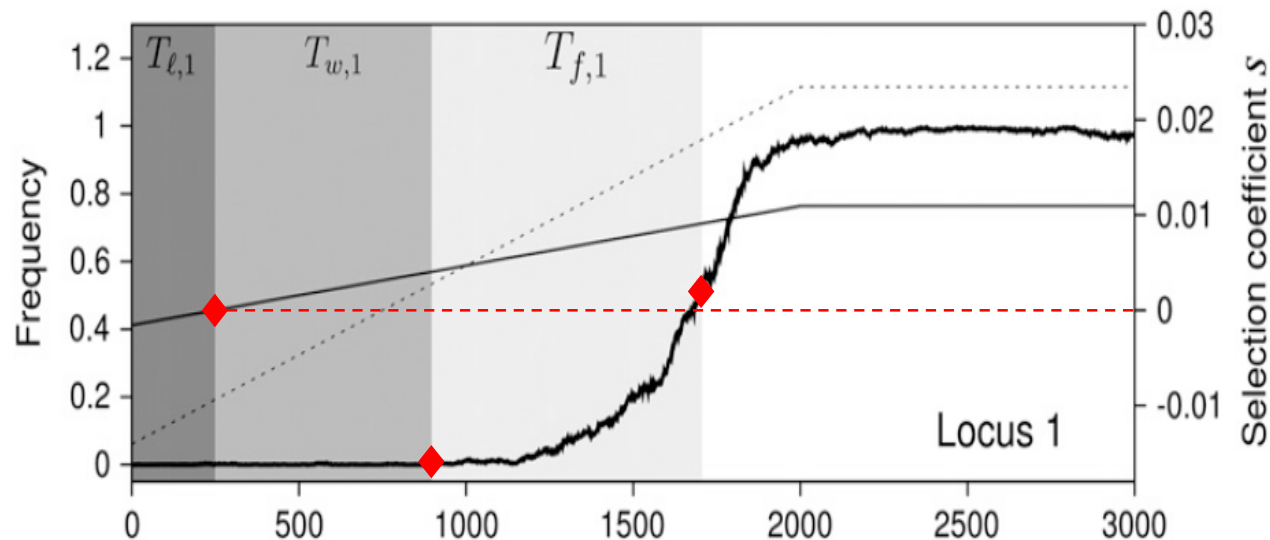
Directional environmental change

- Low mutation regime (origin-fixation process):
Adaptive walk by sequential fixation in otherwise monomorphic population.
- Under **gradual trend, selection coefficients change as optimum moves**
→ A mutation first needs to **become beneficial** (duration T_ℓ),
then **escape drift** (T_w), then **reach high frequency** (T_f)¹.



Directional environmental change

- Low mutation regime (origin-fixation process): Adaptive walk by sequential fixation in otherwise monomorphic population.
- Under **gradual trend, selection coefficients change as optimum moves**
→ A mutation first needs to **become beneficial** (duration T_ℓ), then **escape drift** (T_w), then **reach high frequency** (T_f)¹.
- Slower environmental changes are dominated by T_ℓ , **favoring small steps because they become beneficial earlier**



Directional environmental change

- Low mutation regime (origin-fixation process):
Adaptive walk by sequential fixation in otherwise monomorphic population.
- Under **gradual trend**, **selection coefficients change as optimum moves**
- A **single composite parameter** determines the genetics of adaptation¹:

$$\gamma = \frac{v}{NUS\sigma_{\alpha}^3}$$

v : Speed of environmental change

N : population size

U : genomic mutation rate

S : Strength of stabilizing selection

σ_{α} : SD of mutation phenotypic effects

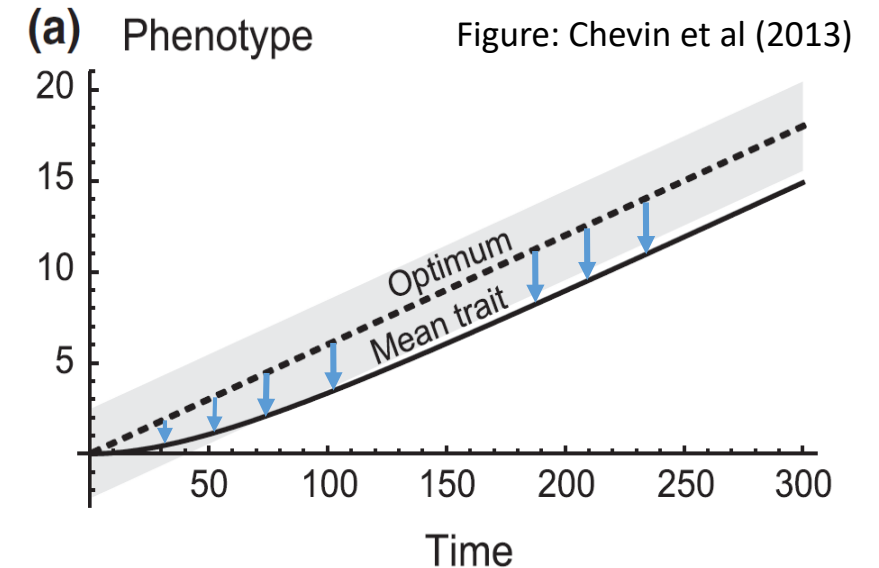
} Ecology

} Adaptive potential

- **Large γ** : Environment changes fast relative the adaptive potential.
Adaptation is **genetically limited**, mutations of large effects can fix (cf Orr 1998)
- **Small γ** : Environment changes slowly relative the adaptive potential.
Adaptation is **environmentally limited**, mutations of small effects mostly fix

Directional environmental change

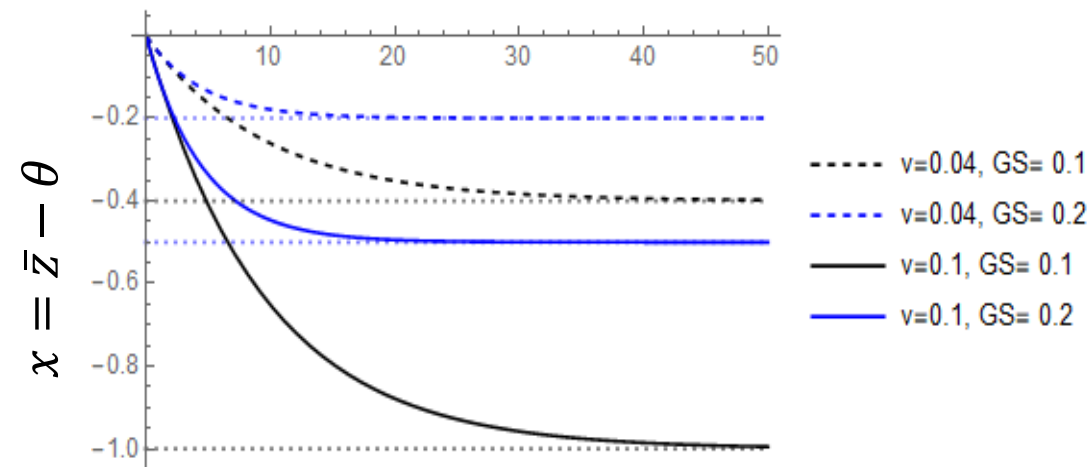
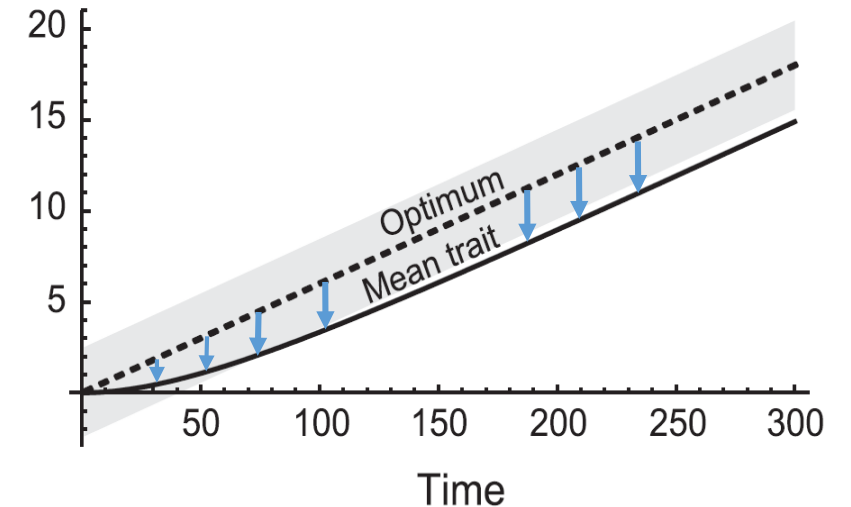
- Highly polymorphic regime:
- Distance to optimum $x = \bar{z} - \theta$ initially increases as phenotype lags behind optimum.
- This increases the strength of directional selection and response.



Directional environmental change

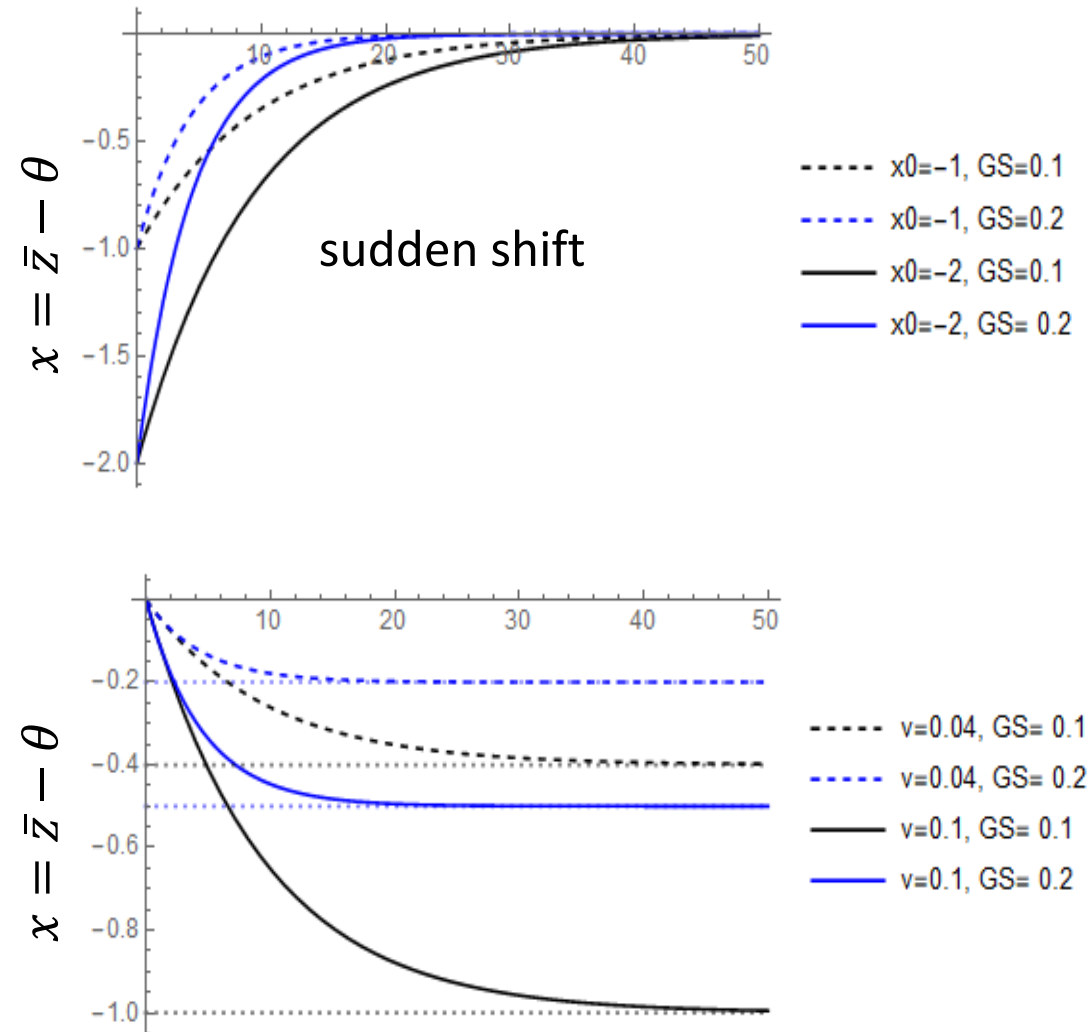
- Highly polymorphic regime:
- Distance to optimum $x = \bar{z} - \theta$ initially increases as phenotype lags behind optimum.
- This increases the strength of directional selection and response.
- Lag eventually equilibrates, with mean phenotype evolving at same speed as optimum: $\Delta\bar{z} = -GSx_{eq} = v$
- Equilibrium lag is thus $x_{eq} = -\frac{v}{SG}$
 → larger with **fast environmental change** and **low adaptive potential**

(a) Phenotype Figure: Chevin et al (2013)



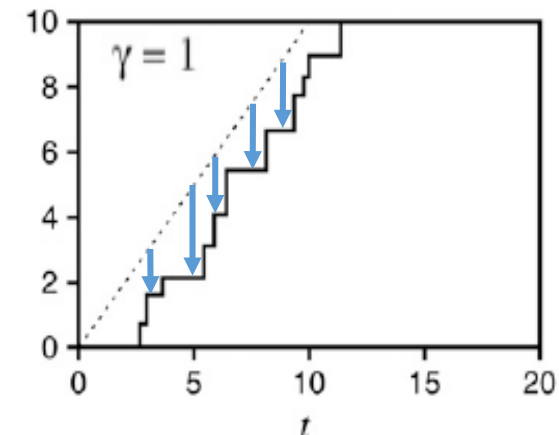
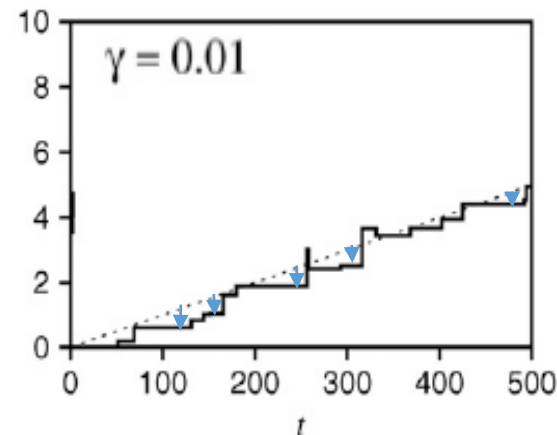
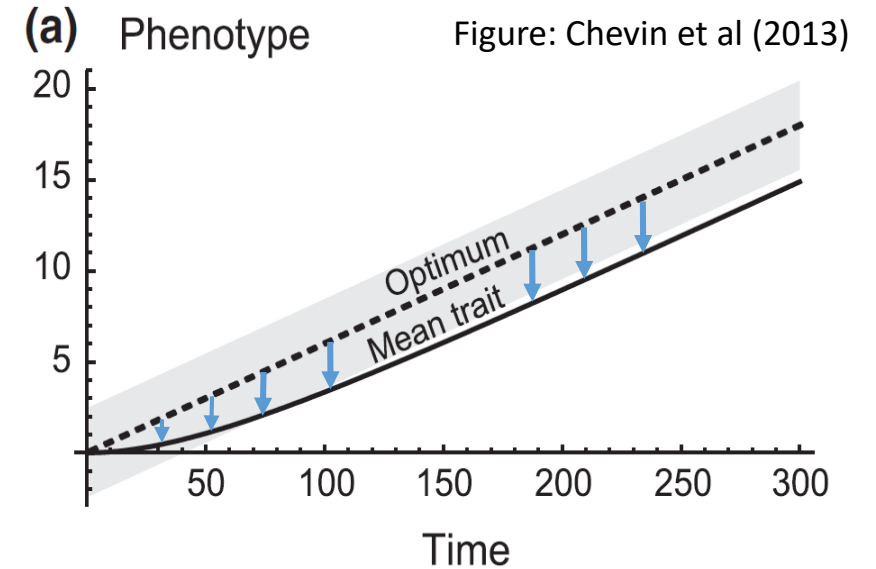
Directional environmental change

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- Reciprocal dynamics to sudden shift



Directional environmental change

- Highly polymorphic regime:
- Distance to optimum $x = \bar{z} - \theta$ initially increases as phenotype lags behind optimum.
- This increases the strength of directional selection and response.
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- Equilibrium lag is thus $x_{eq} = -\frac{v}{SG}$
 → larger with **fast environmental change** and **low adaptive potential**
- Analog to adaptive walk regime: $\gamma = \frac{v}{NUS\sigma_\alpha^3}$

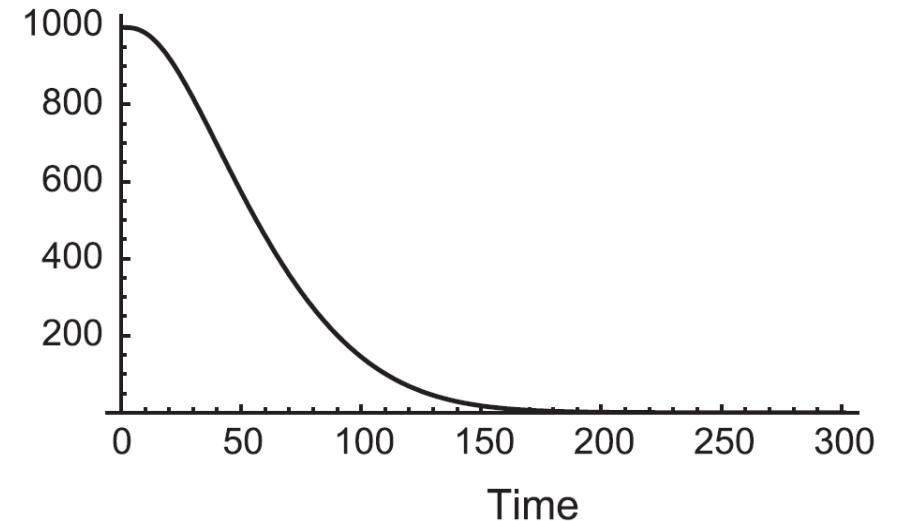
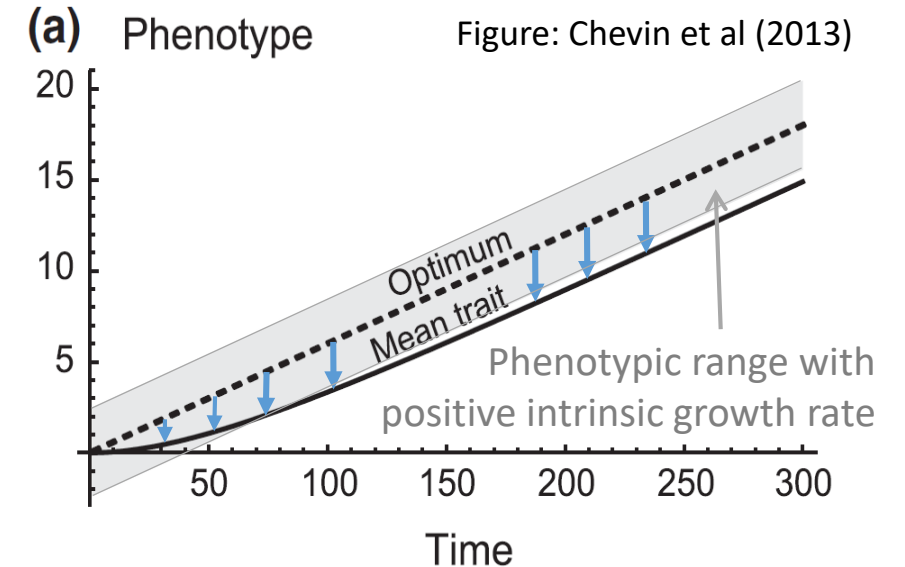


Directional environmental change

- Highly polymorphic regime:
- Maximum reduction in population growth rate caused by maladaptation:
Lag load $L = \frac{S}{2} x_{eq}^2 = \frac{v^2}{2SG^2}$
- The **critical rate of environmental change** at which $r_{max} - L = 0$ is $v_c = \sqrt{2r_{max}SG}$
- Narrower fitness peak (larger S) causes larger fitness drop for a given mismatch x , but also faster evolutionary reduction of x . The latter dominates, increasing v_c .
- But may be violated with other fitness function²

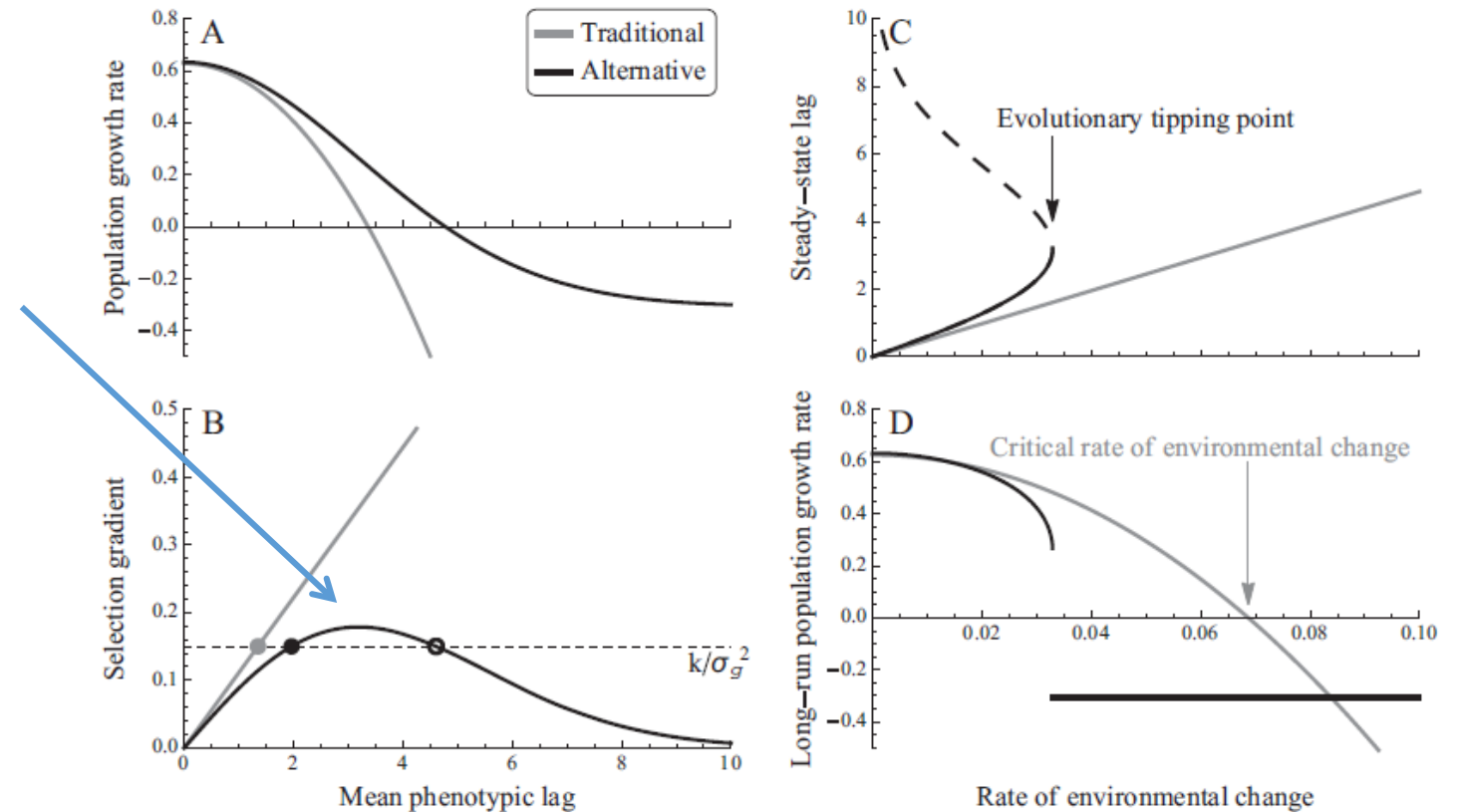
1: Pease et al 1989; Lynch et al 1991; Lynch & Lande 1993

2: Osmond et al (2017); Klausmeier et al (2020)



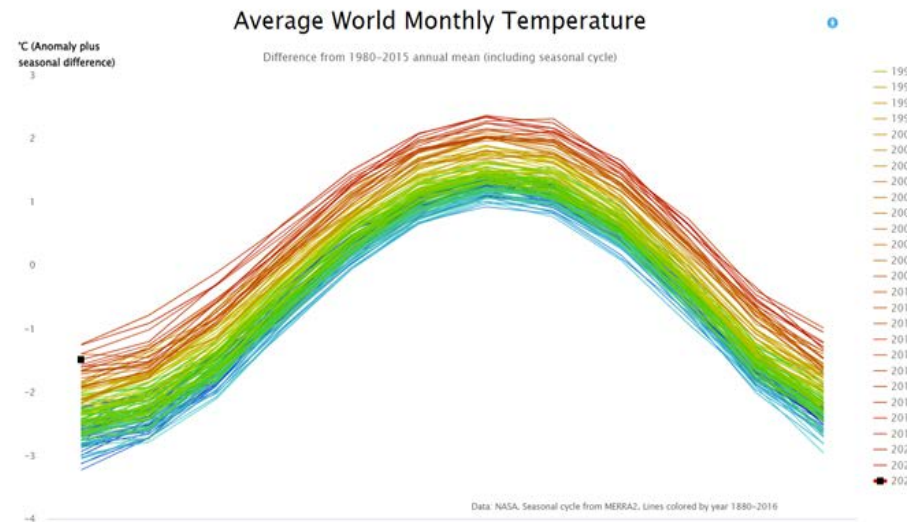
Directional environmental change

- Fitness function where strength of selection β does not increase monotonically with maladaptation
- Maximum selection gradient = **Tipping point** for rate of evolution
- Larger lags lead to ever-increasing maladaptation: **Extinction vortex**
- Even transient increase in lag may be impossible to recover from: **hysteresis**



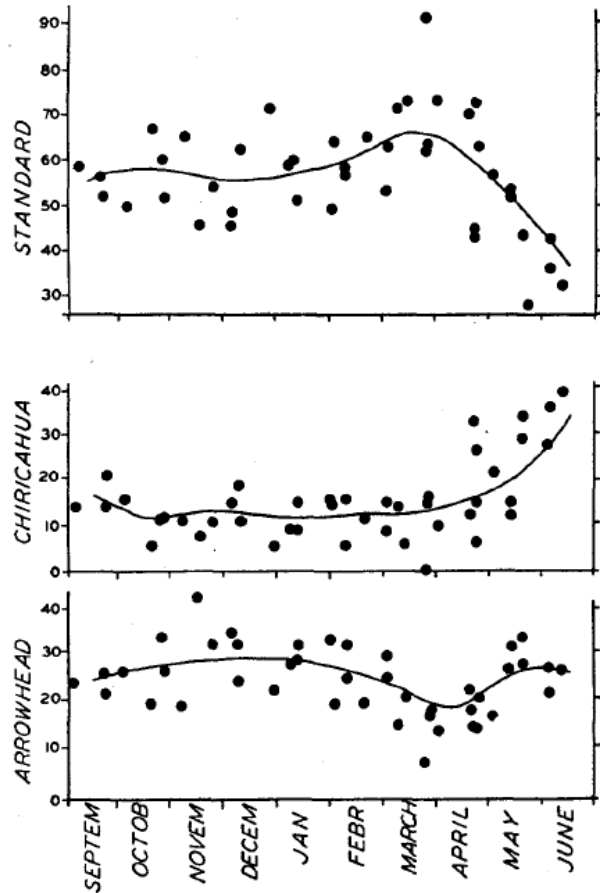
Cycling environment

- Seasonality occurs on evolutionary timescales for short lived species.
- Such organisms usually have large population sizes, thus high adaptive potential.
- Other cycles occur with larger periods (El Niño, North Atlantic oscillation...), and could be tracked by more long-lived organisms

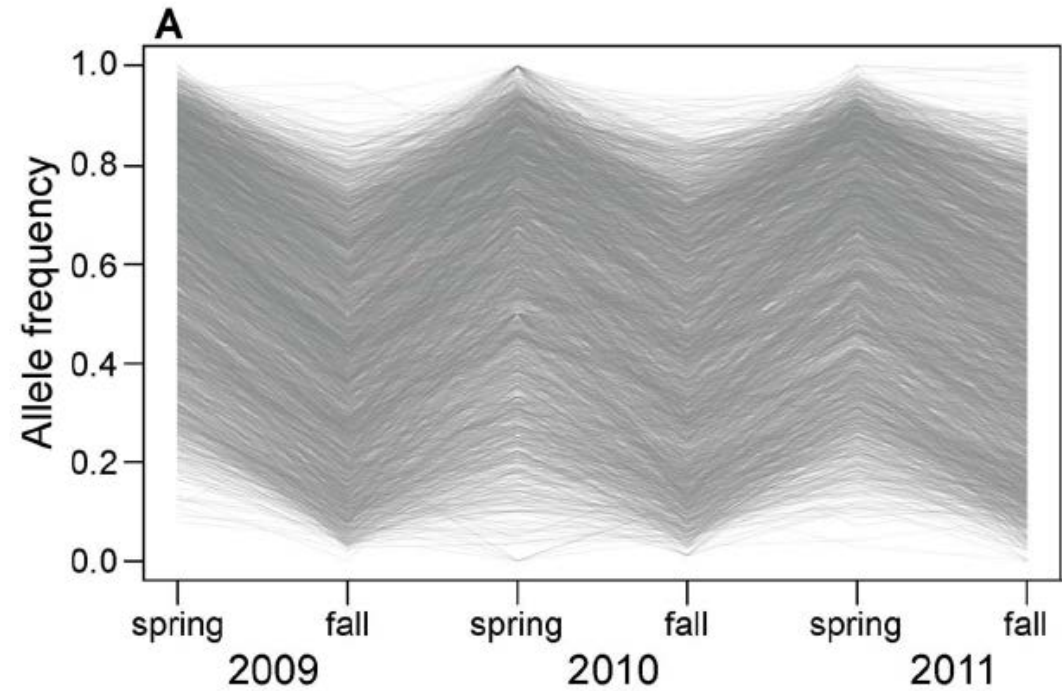


Cycling environment

Chromosomal inversions in *Drosophila*
(Dobzhansky 1943 Genetics)



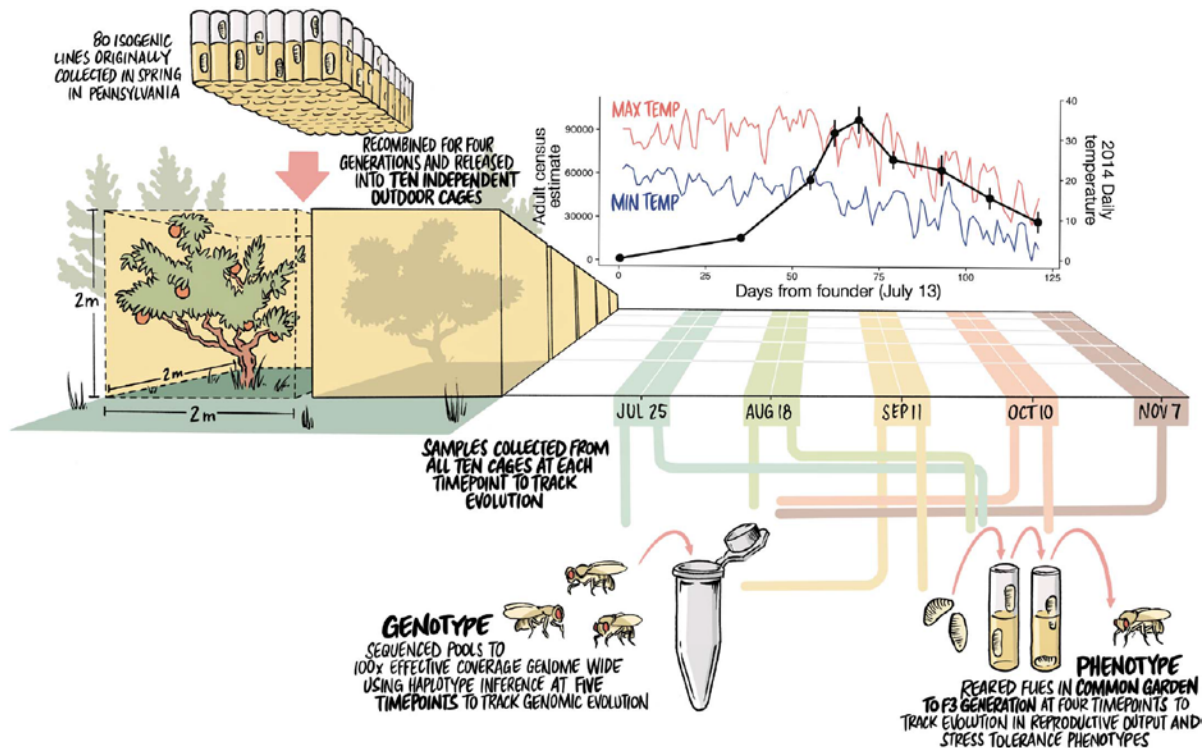
1750 SNPs throughout genome
(Bergland et al 2017 PLoS Genetics)



Cycling environment

Direct observation of adaptive tracking on ecological time scales in *Drosophila* *Science* 375, 1246 (2022)

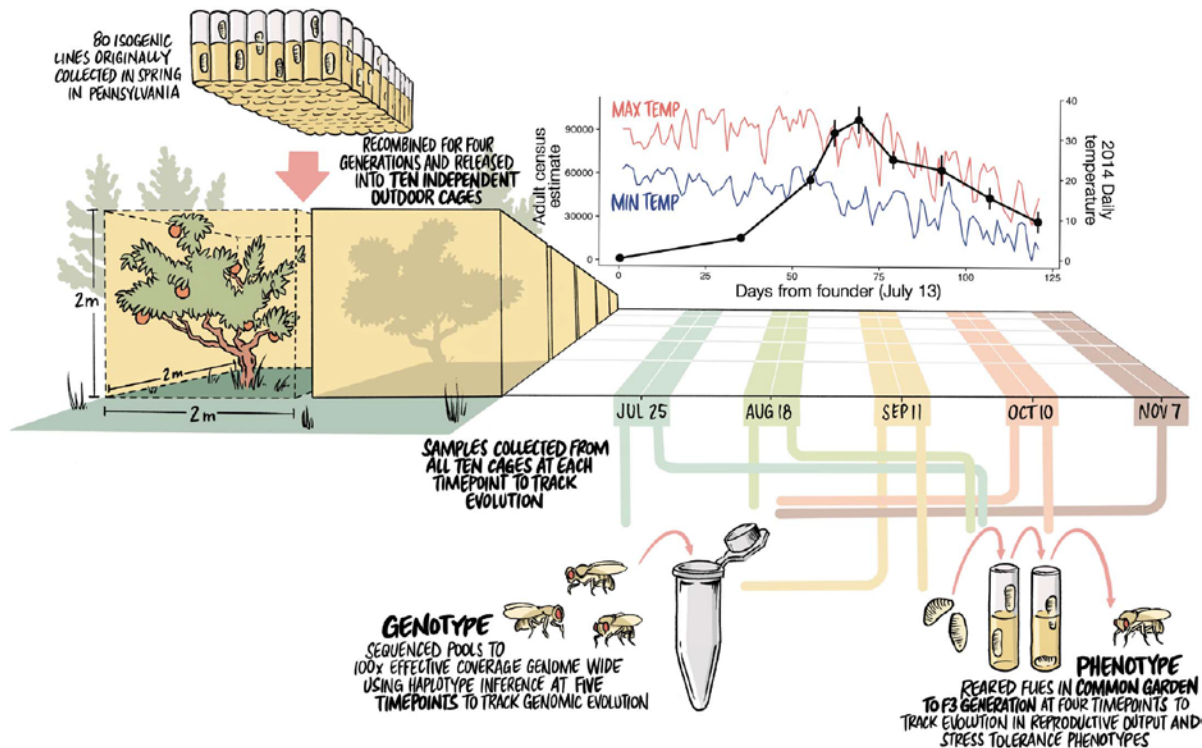
Seth M. Rudman^{1,2*}†, Sharon I. Greenblum^{3,4*}†, Subhash Rajpurohit^{1,5}†, Nicolas J. Betancourt¹, Jinjoo Hanna¹, Susanne Tilk³, Tuya Yokoyama³, Dmitri A. Petrov³, Paul Schmidt^{1*}



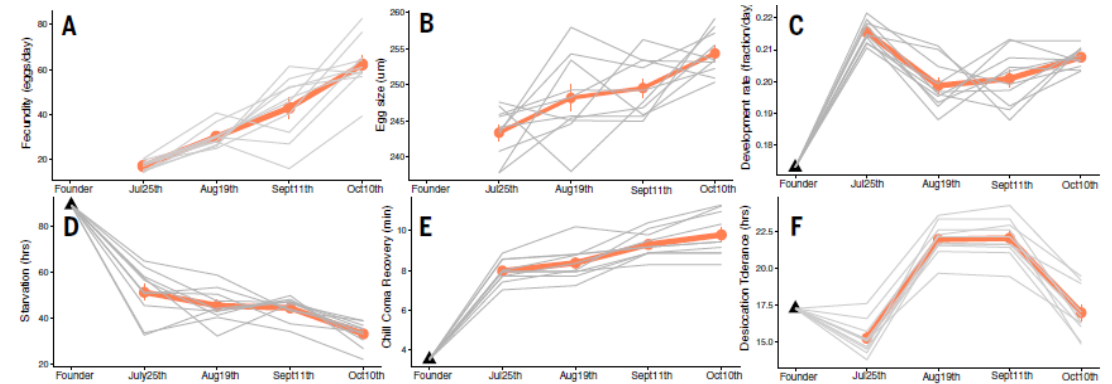
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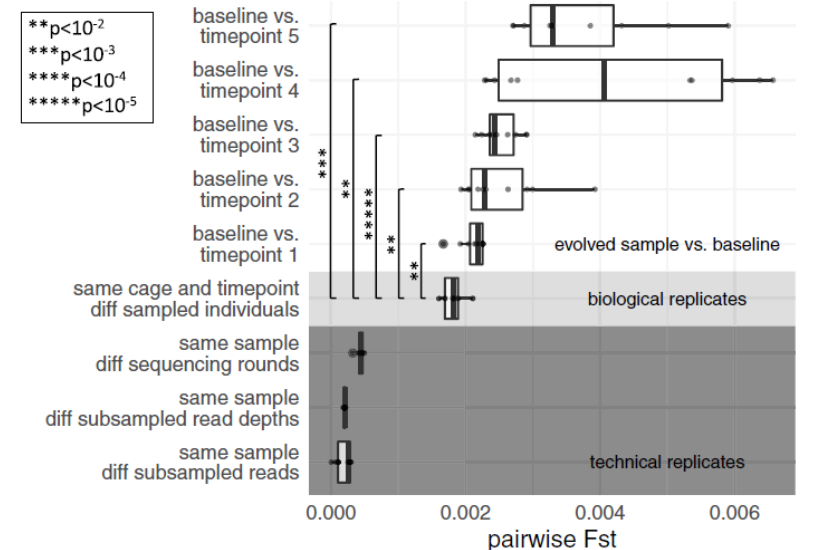
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Parallel phenotypic changes



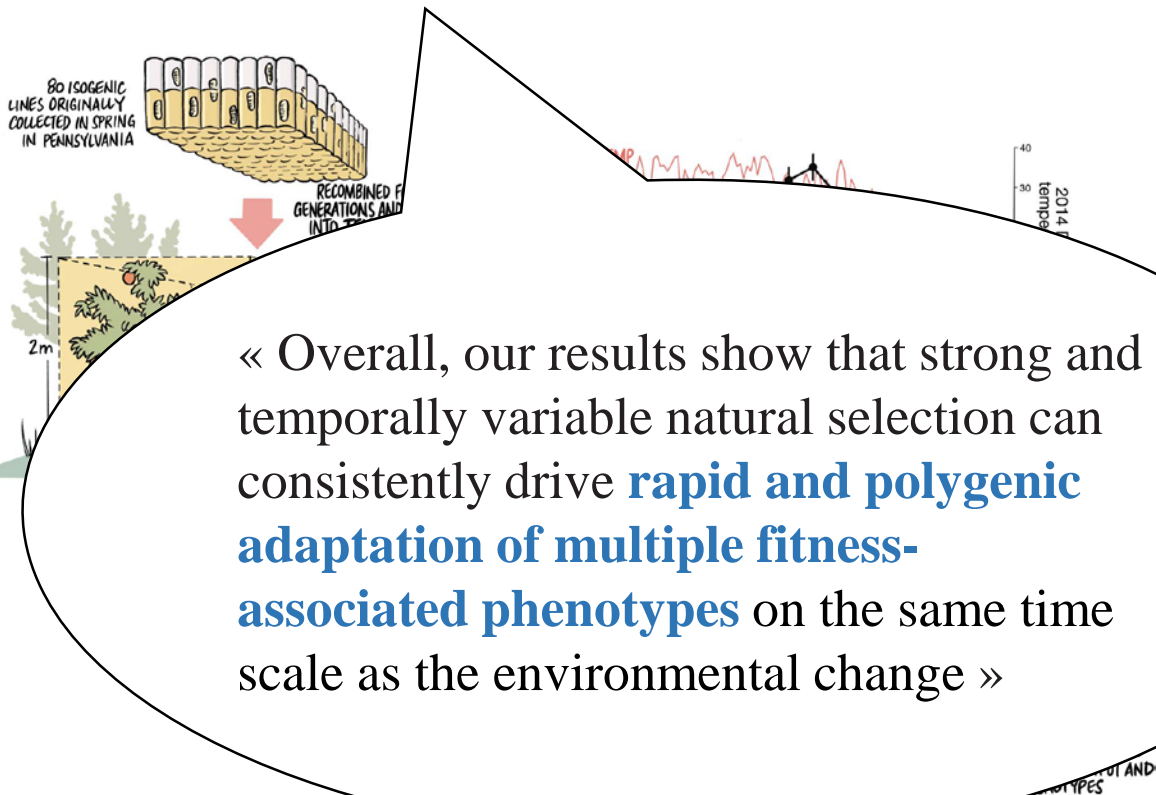
Largely repeatable genetic differentiation over time



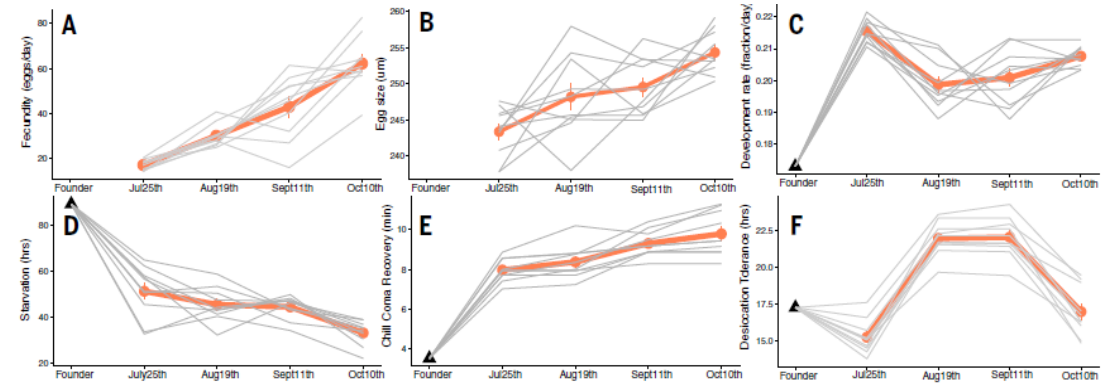
Cycling environment

Direct observation of adaptive tracking on ecological time scales in *Drosophila* *Science* **375**, 1246 (2022)

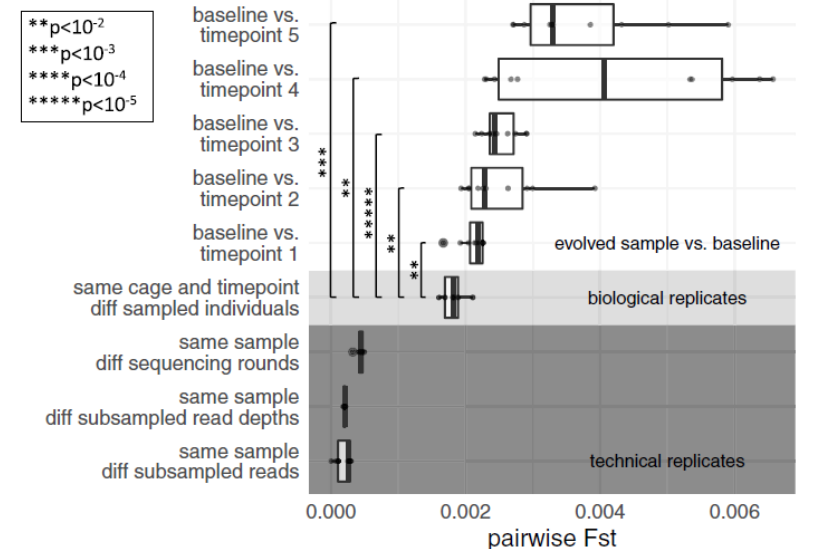
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Parallel phenotypic changes

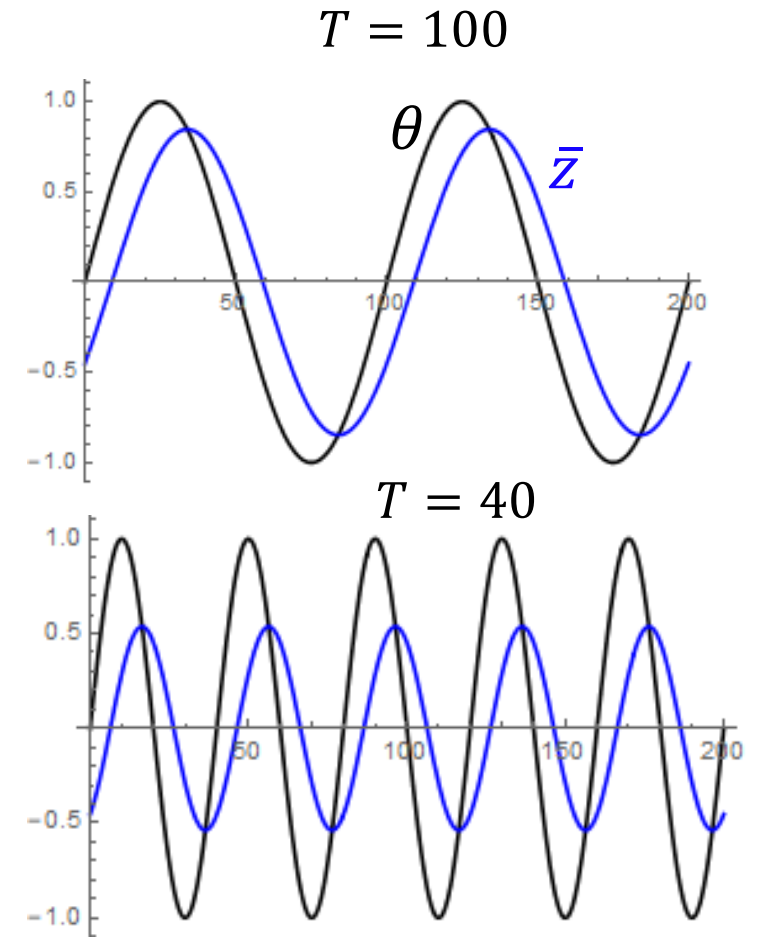


Largely repeatable genetic differentiation over time



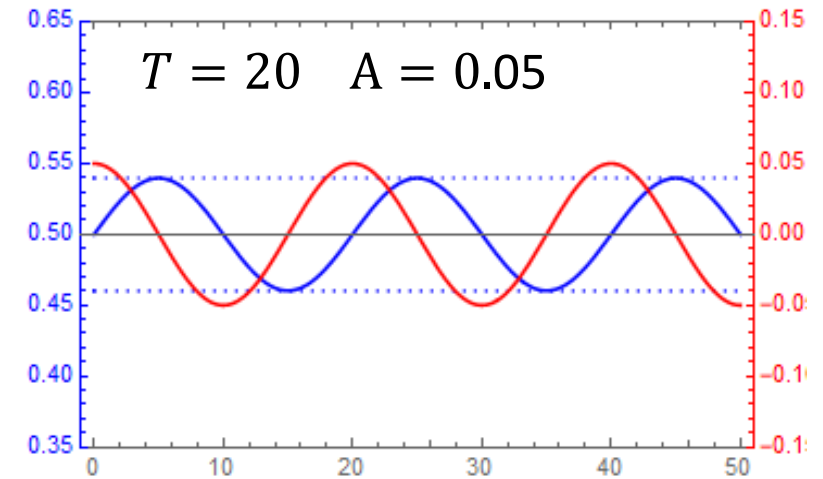
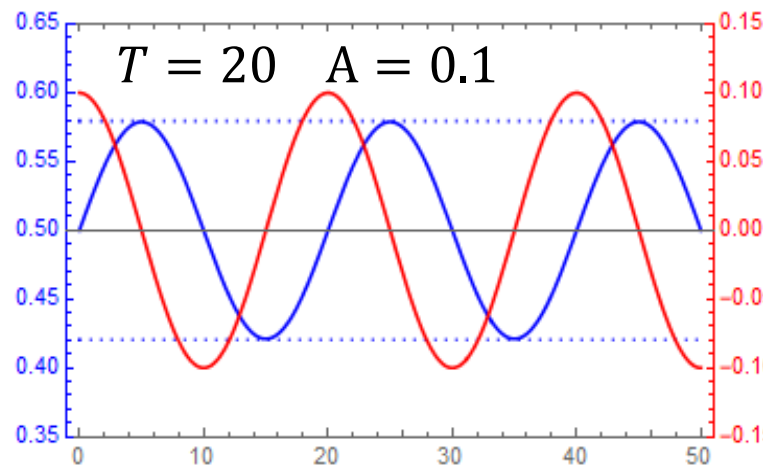
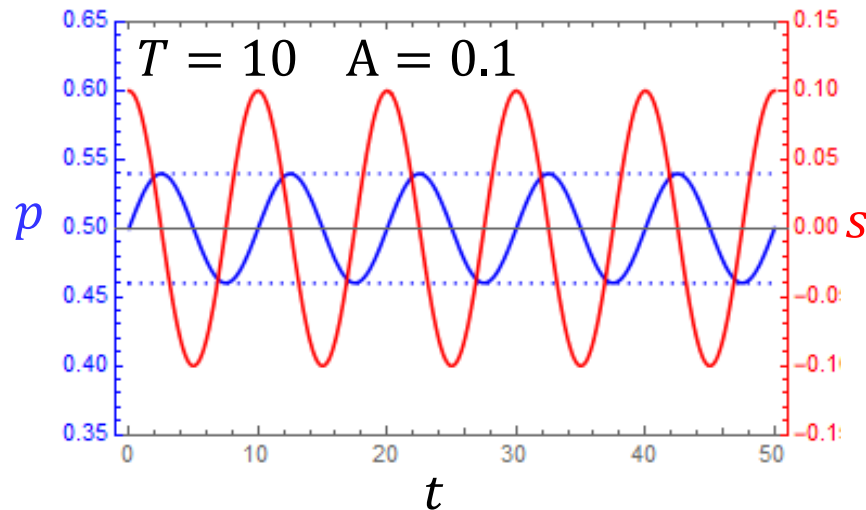
Cycling environment

- **Model**¹: $\theta = A \sin\left(\frac{2\pi t}{T}\right)$, amplitude A and period T
- **Polygenic trait** with constant G
- After $\sim \frac{1}{SG}$ generations, mean phenotype settles into sine wave with same period as optimum, but:
 - amplitude multiplied by $\zeta = \frac{SGT}{\sqrt{(SGT)^2 + (2\pi)^2}} \leq 1$
 - phase shifted (delayed) by $\varphi = -\text{arcTan}\left(\frac{2\pi}{SGT}\right)$
- **Higher adaptive potential SG** and **slower oscillations** (larger T) lead to closer adaptive tracking of optimum ($\zeta \rightarrow 1$ and $\varphi \rightarrow 0$)



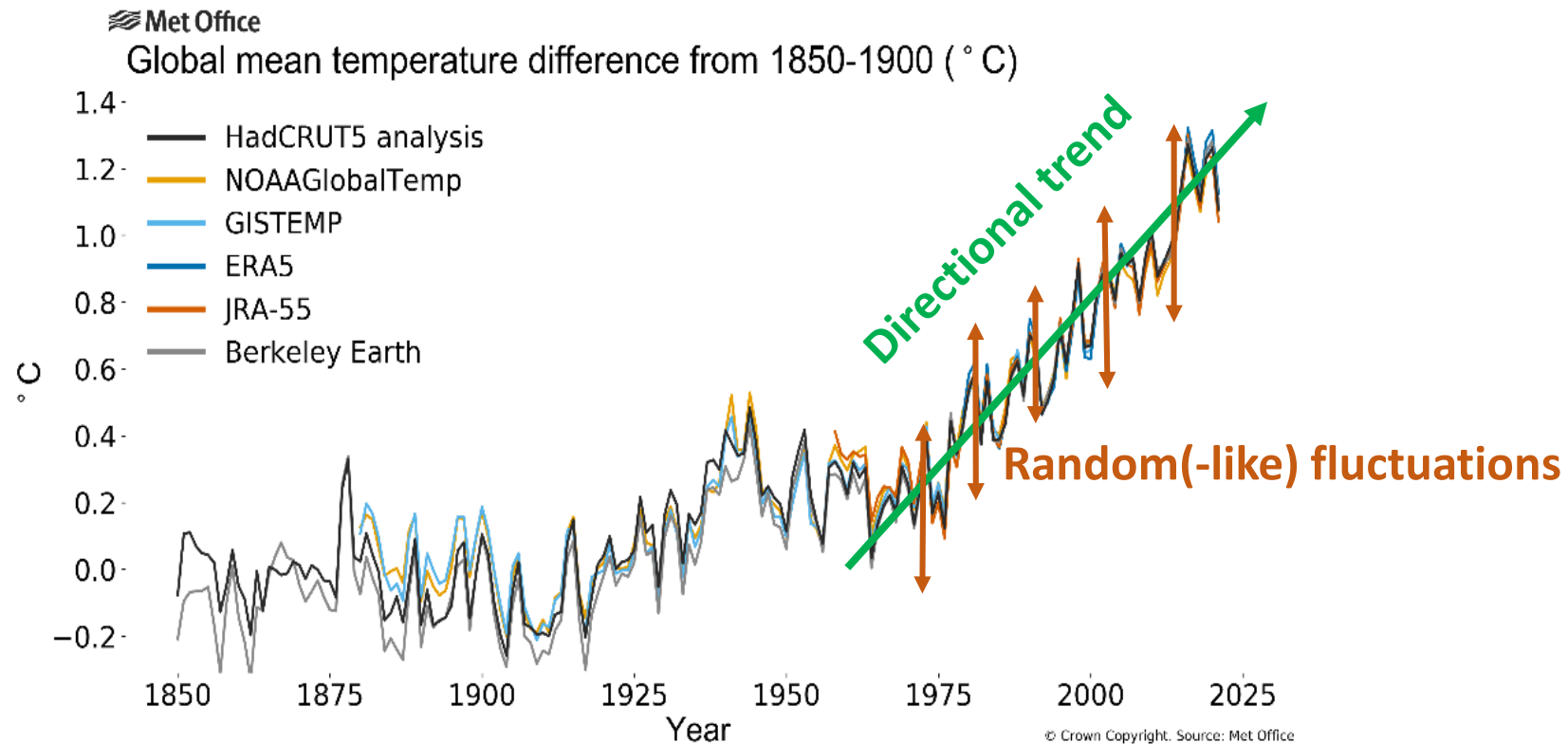
Cycling environment

- **Single locus** with selection coefficient $s = A \sin\left(\frac{2\pi t}{T}\right)$
 - Quarter-period lag between frequency p and s ($\max(p)$ when $s = 0$)
 - Amplitude of p is $A_p \approx A \frac{T}{8\pi} \rightarrow$ larger under larger period and maximum s



Randomly fluctuating environment

- Most environments exhibit residual noise, after removing any trend



Randomly fluctuating environment

- Most environments exhibit residual noise, after removing any trend
- These fluctuations may well have deterministic causes, but if the latter are
 - (i) unknown
 - (ii) external to the system (and potentially complex)
 - (iii) beyond reach of measurement precision,

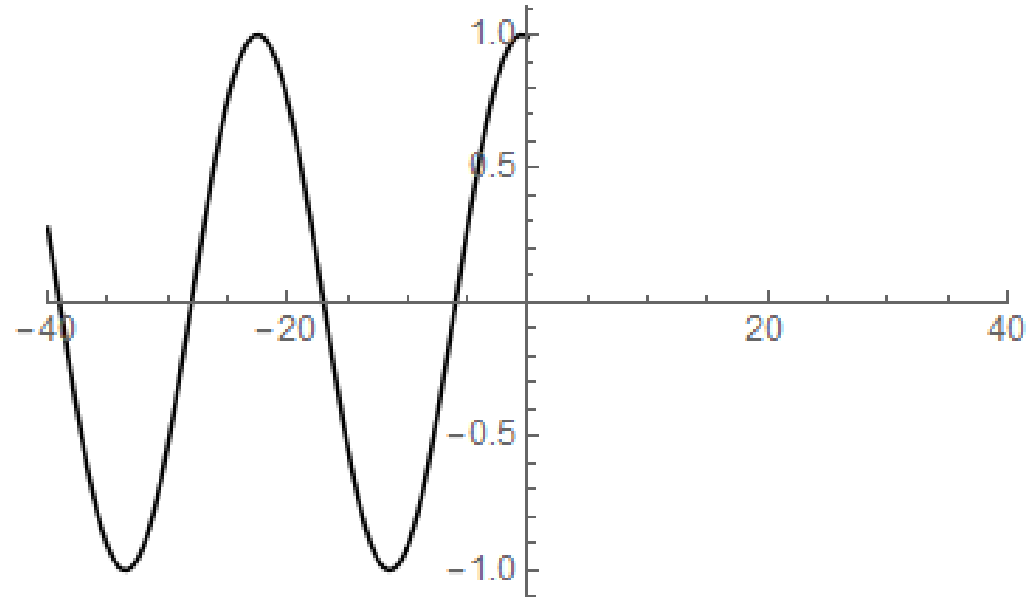
then fluctuations are **effectively random**, both to scientists analyzing them and to organisms experiencing them.

→ Treated as **stochastic processes** = random variable with time dependence

Prediction in stochastic environment

- Randomness matters when making predictions

Deterministic time series:

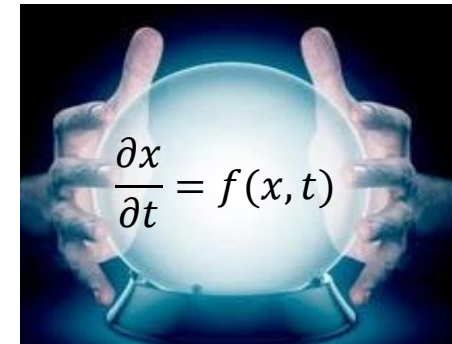
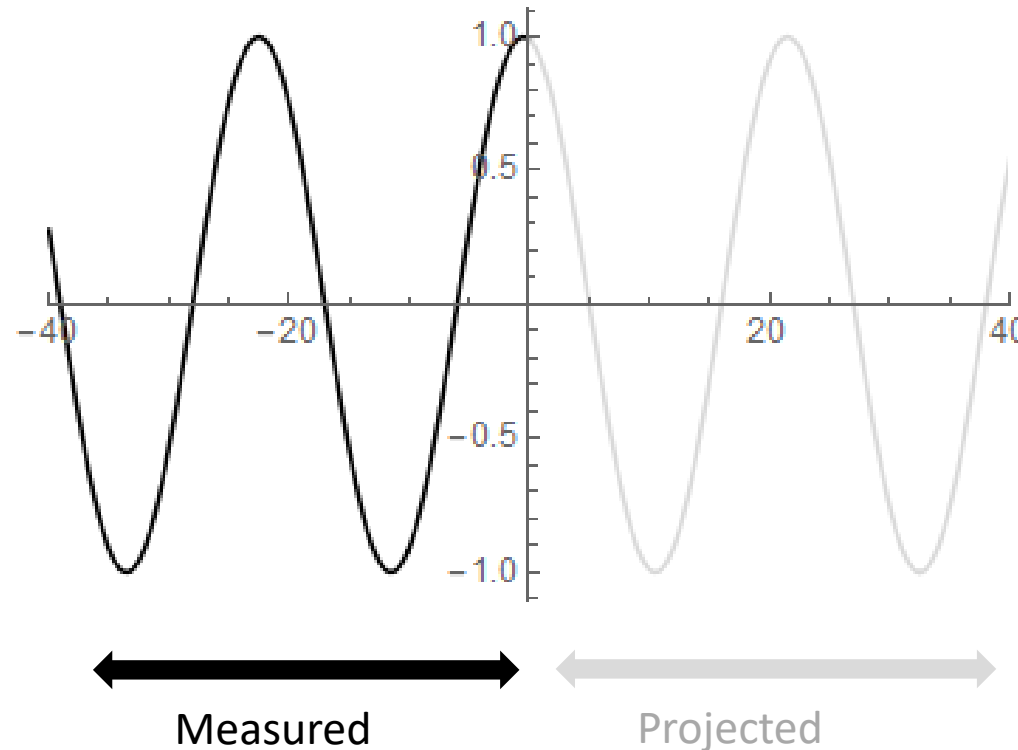


Measured

Prediction in stochastic environment

- Randomness matters when making predictions

Deterministic time series:

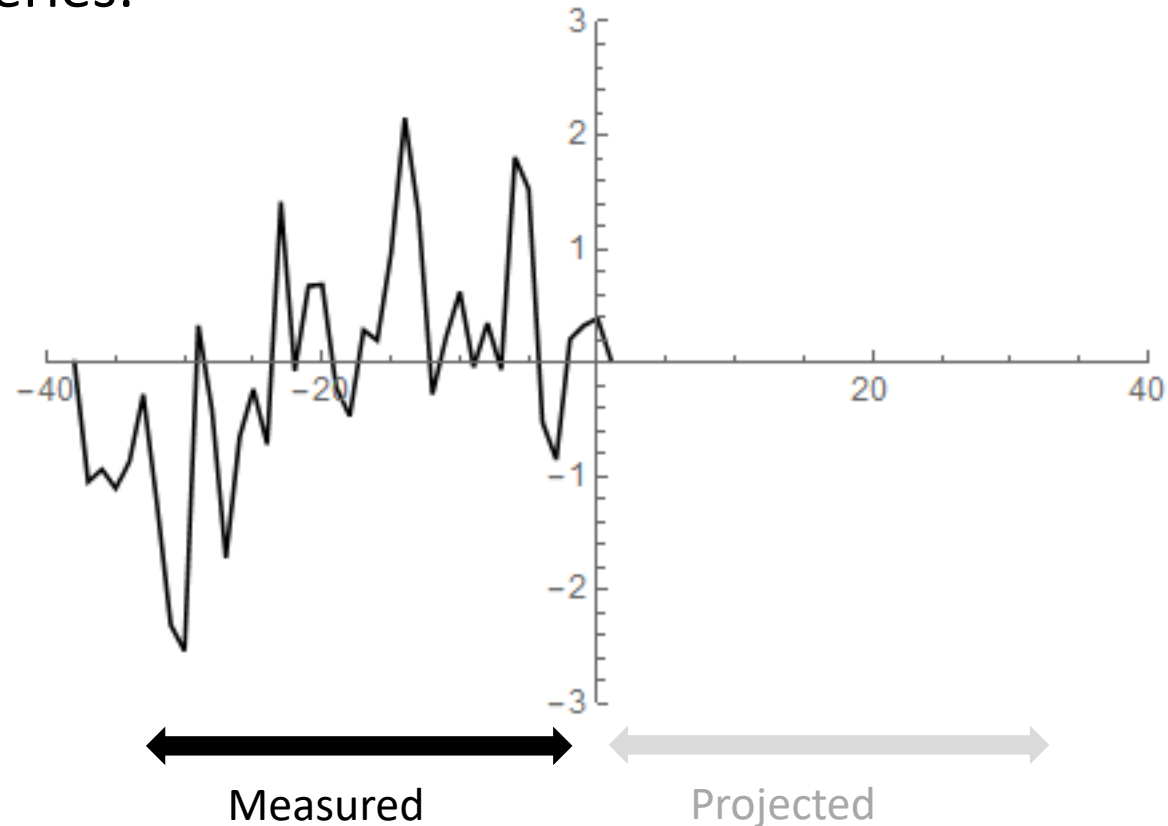


The **future is certain** provided accurate measurement of the past, and perfect knowledge of causal factors.

Prediction in stochastic environment

- Randomness matters when making predictions

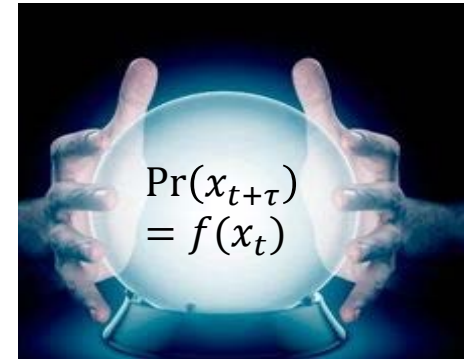
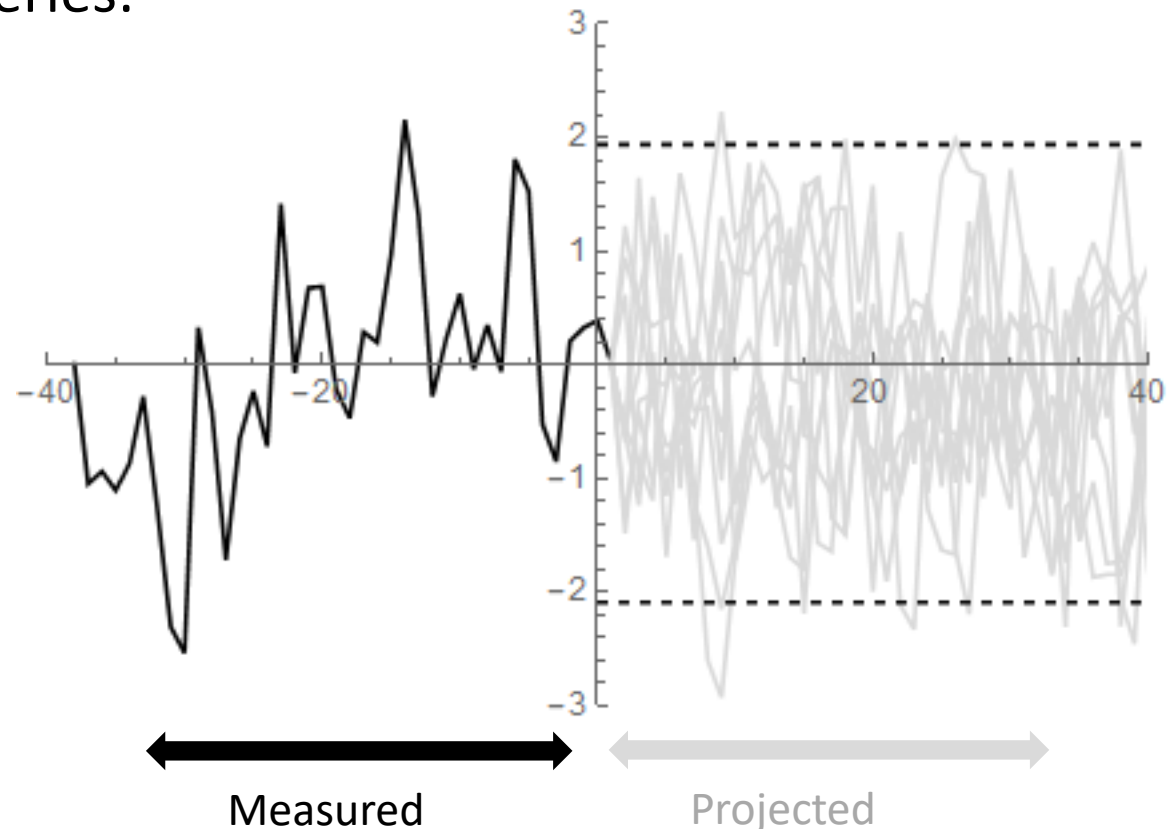
Stochastic time series:



Prediction in stochastic environment

- Randomness matters when making predictions

Stochastic time series:

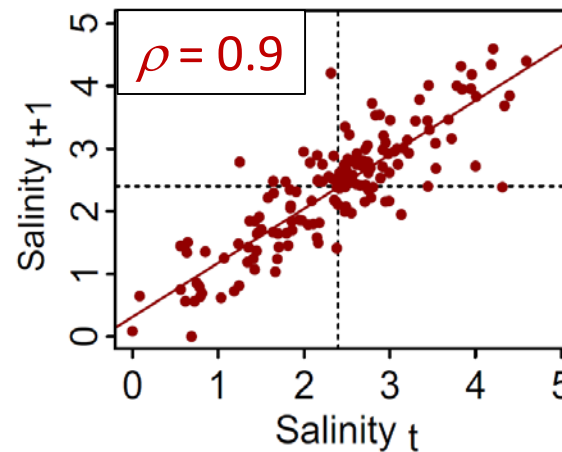
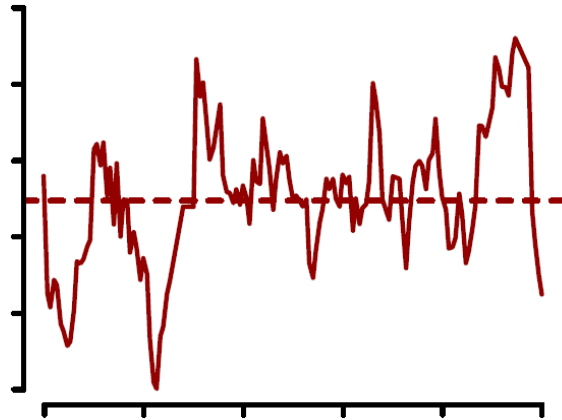
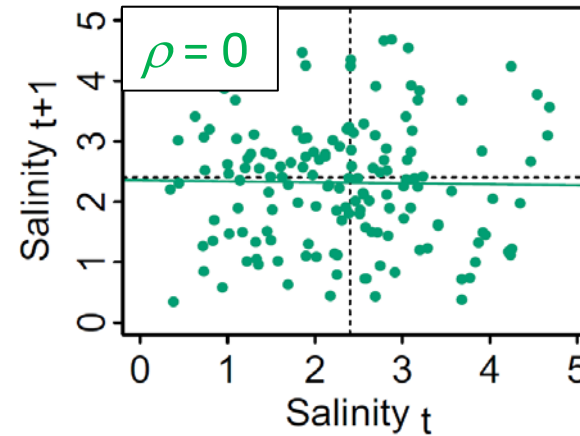
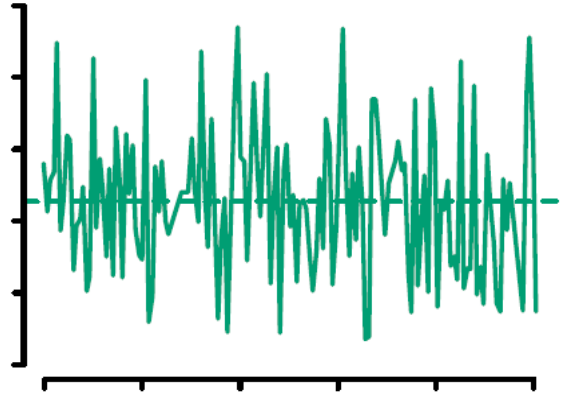


The future is probabilistic even with perfect measurement

Variance of the process matters, not just expectation

Prediction in stochastic environment

- Temporal autocorrelation ρ determines timescale of predictability



- Related to “colour » of environmental noise¹

Fig. from Leung et al (2020 Ecol Lett)
1: Vasseur & Yodzis (2004 Ecology)

Evolutionary responses to fluctuating optimum

- Selection on **mutation with phenotypic effect α** in background phenotype m , in haploid population¹.

Denoting $\psi = \ln\left(\frac{p}{q}\right)$, ie the logit frequency,

$$\Delta\psi = \ln W_{m+\alpha} - \ln W_m = -\frac{s\alpha}{2} [\alpha + 2(m - \theta)]$$
$$\Rightarrow \psi_t = \psi_0 - \frac{s\alpha}{2} \left[\alpha t + 2 \sum_{i=0}^{t-1} (m_i - \theta_i) \right]$$

Additive in mismatch \rightarrow **If optimum θ follows a Gaussian process, so does ψ .**

If changes in background mean phenotype m can be neglected, then ψ simply integrates all past optimas, with equal weight on all times

1: Kimura (1954 Genetics), Gillespie (1991),
Chevin (2019 Genetics)

Evolutionary responses to fluctuating optimum

- Assume the optimum follows a stationary autocorrelated Gaussian process (AR1)

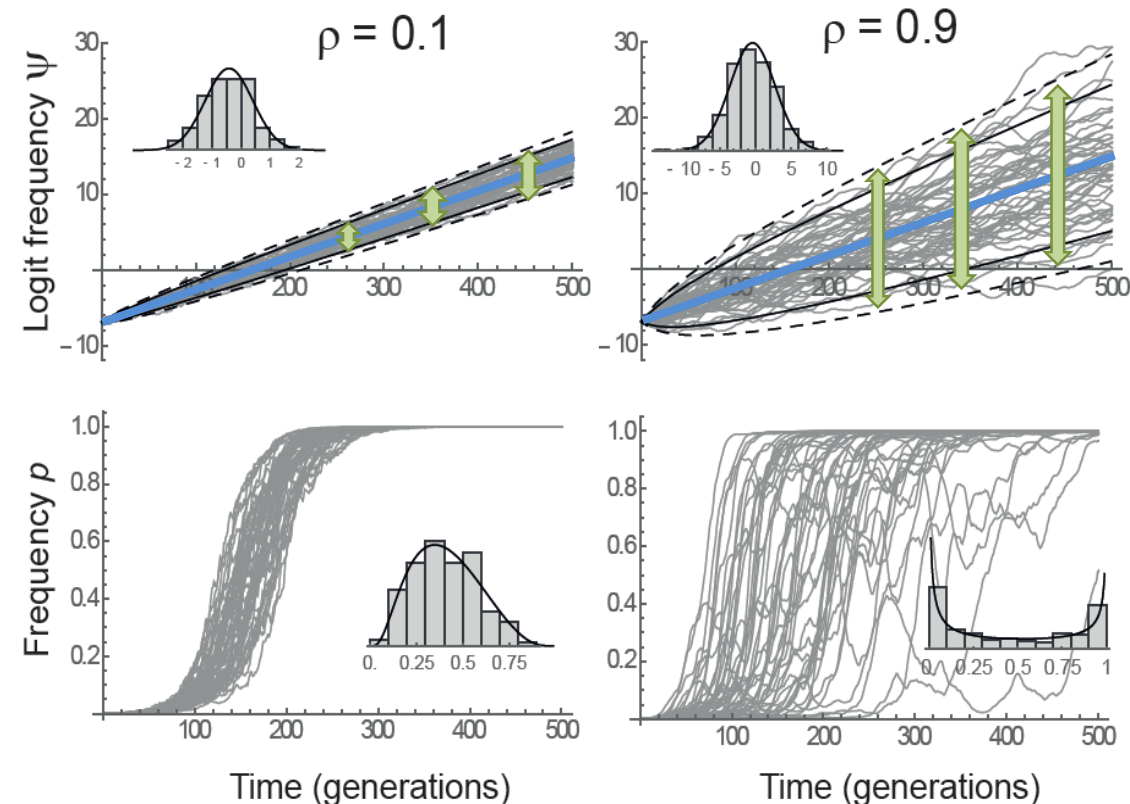
- Fluctuation pattern has **no influence on expected change** in (logit) frequency

- Stochastic **variance of ψ** is

$$\sigma_{\psi,t}^2 \approx \sigma_S^2 \frac{1+\rho}{1-\rho} t, \text{ with } \sigma_S^2 = (S\alpha\sigma_\theta)^2$$

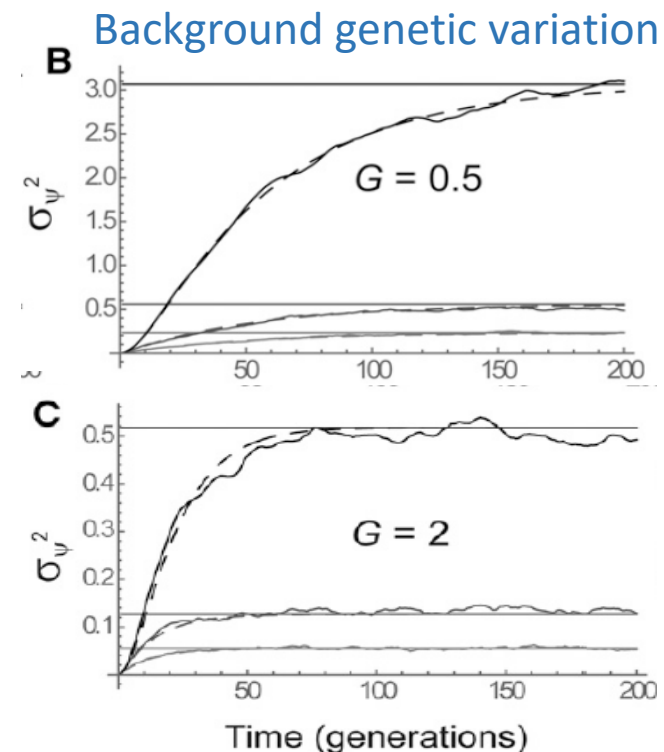
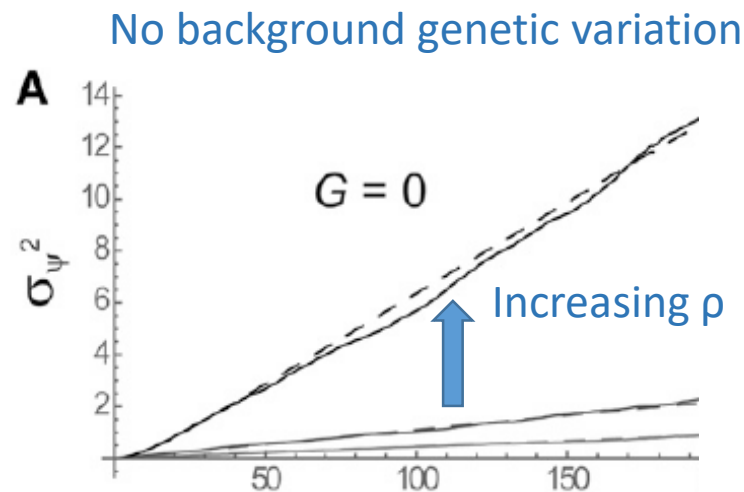
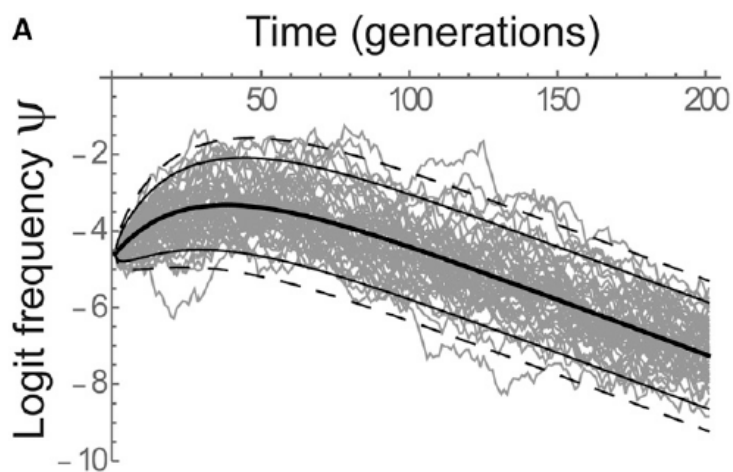
→ Increases linearly, faster under higher autocorrelation

- On p scale, variance of ψ translates into variance in the timing of selective sweeps



Evolutionary responses to fluctuating optimum

- If background genetic variance for the trait is normally distributed, then mean background also evolves in response to fluctuating optimum.
- The process for $\psi = \text{logit}(p)$ becomes **stationary, with variance that plateaus**
→ Other polymorphic loci buffer the stochasticity perceived by major gene



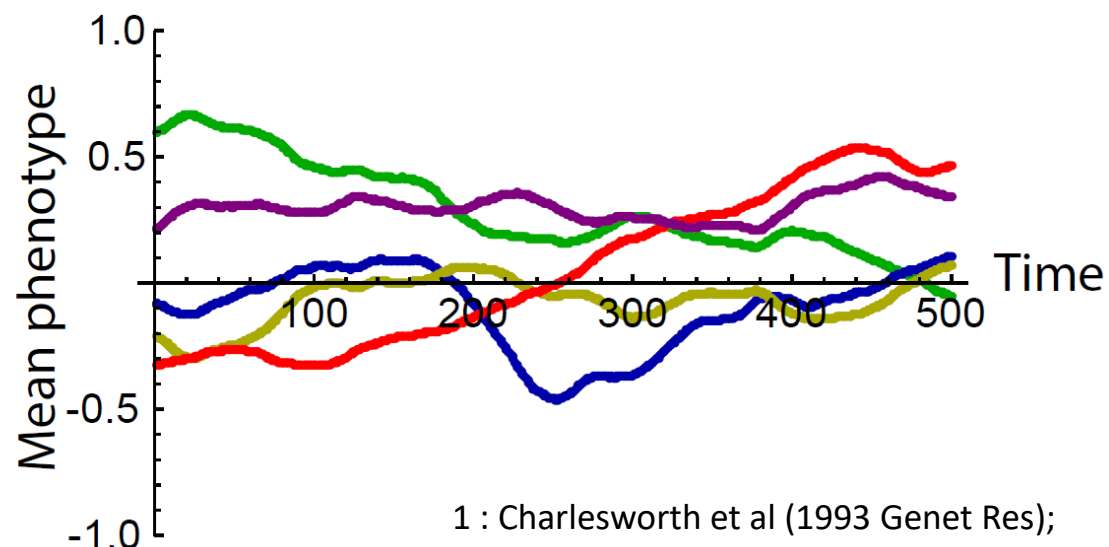
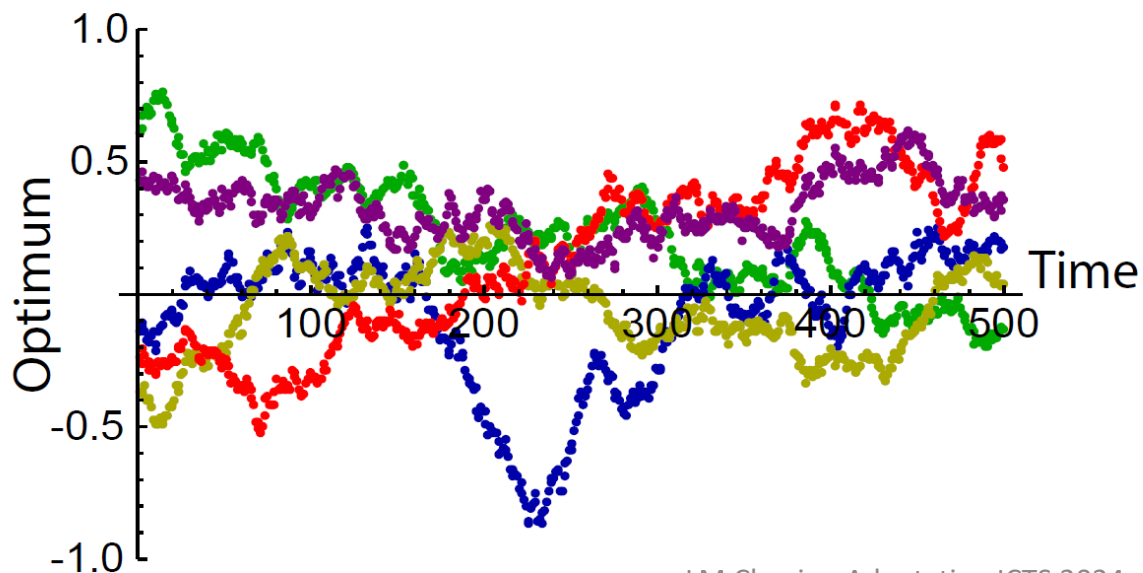
Evolutionary responses to fluctuating optimum

- For polygenic trait with constant variance, the mean phenotype¹ is

$$\bar{z}_t = \bar{z}_0(1 - GS)^t + GS \sum_{j=1}^t (1 - GS)^{j-1} \theta_{t-j} \xrightarrow{t \rightarrow \infty} GS \sum_{j=1}^{\infty} (1 - GS)^{j-1} \theta_{t-j}$$

→ **Weighted average of past optima**, with more weight on more recent ones.

Smooths environmental “signal”, all the more as adaptive potential SG is small



Evolutionary responses to fluctuating optimum

- For polygenic trait with constant variance, the mean phenotype¹ is

$$\bar{z}_t = \bar{z}_0(1 - GS)^t + GS \sum_{j=1}^t (1 - GS)^{j-1} \theta_{t-j} \xrightarrow{t \rightarrow \infty} GS \sum_{j=1}^{\infty} (1 - GS)^{j-1} \theta_{t-j}$$

- If optimum undergoes Gaussian process, so do:
 - the mean phenotype \bar{z} (linear combination of Gaussians)
 - the mismatch with optimum $x = \bar{z} - \theta$

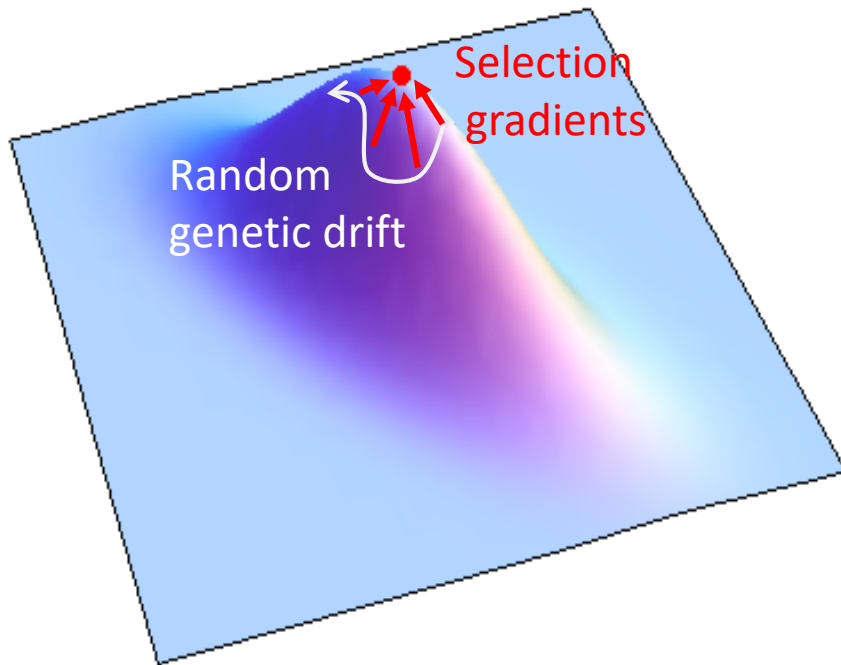
→ The distribution of maladaptation can be summarized by its mean and variance.

- At stationarity:
 - The expected mean phenotype matches the expected optimum
 - But the **variance and autocorrelation of mismatch** play important roles.

1 : Charlesworth et al (1993 Genet Res);

Fluctuations of selection gradient

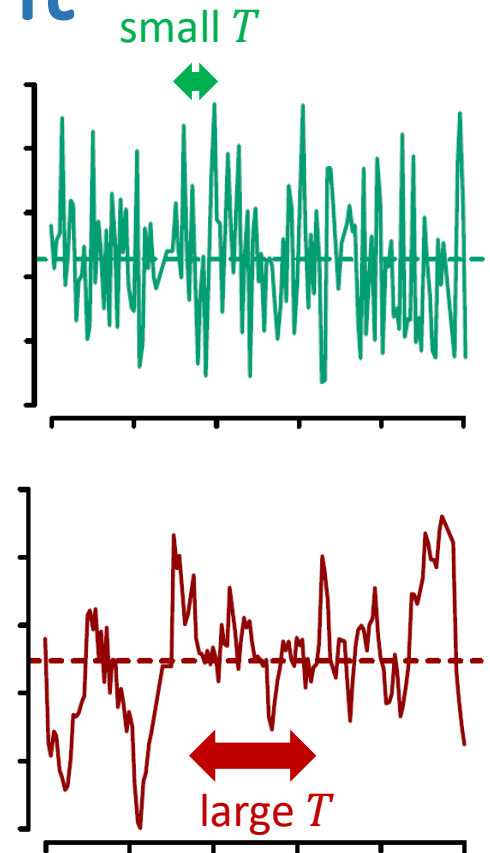
- Directional selection gradient is proportional to phenotypic mismatch, $\beta = -S(\bar{z} - \theta)$
- Even with a constant optimum, **drift causes temporal variation in mismatch** ($\bar{z} - \theta$)



- The variance of directional selection caused by drift around the constant optimum is $V(\beta) = \frac{S}{(2-SG)N_e}$
- **Lower bound for fluctuations** in directional selection, larger for lower N_e and larger S .
- The autocorrelation function of selection gradients is $ACF(\beta, \tau) = (1 - SG)^\tau$
- **Evolutionary inertia** over timescale $1/(SG)$ longer with lower evolutionary potential

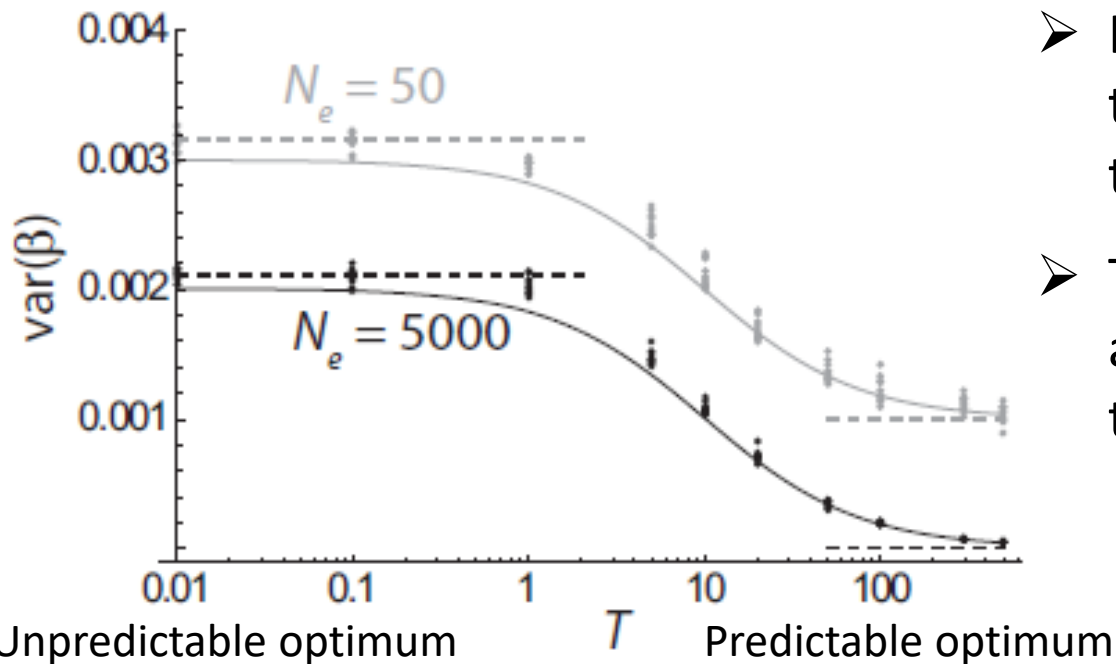
Fluctuations of selection gradient

- Autocorrelated fluctuating optimum (AR1), with T the characteristic time over which optimum is autocorrelated

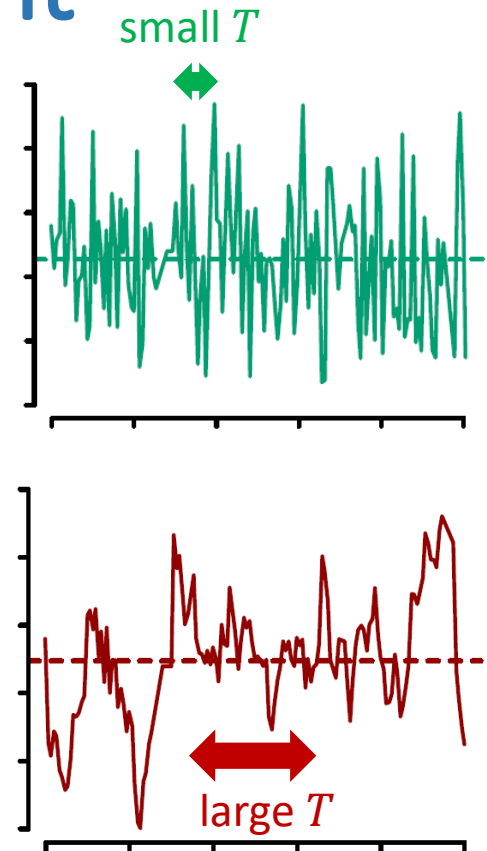


Fluctuations of selection gradient

- Autocorrelated fluctuating optimum (AR1), with T the characteristic time over which optimum is autocorrelated
- Without drift: $V(\beta) \approx \frac{S \sigma_{\theta}^2}{1+SGT}$



- Higher autocorrelation leads to **better adaptive tracking**, thus smaller fluctuations in β
- The variance due to drift around optimum adds up to that of optimum fluctuations



Fluctuations of selection gradient

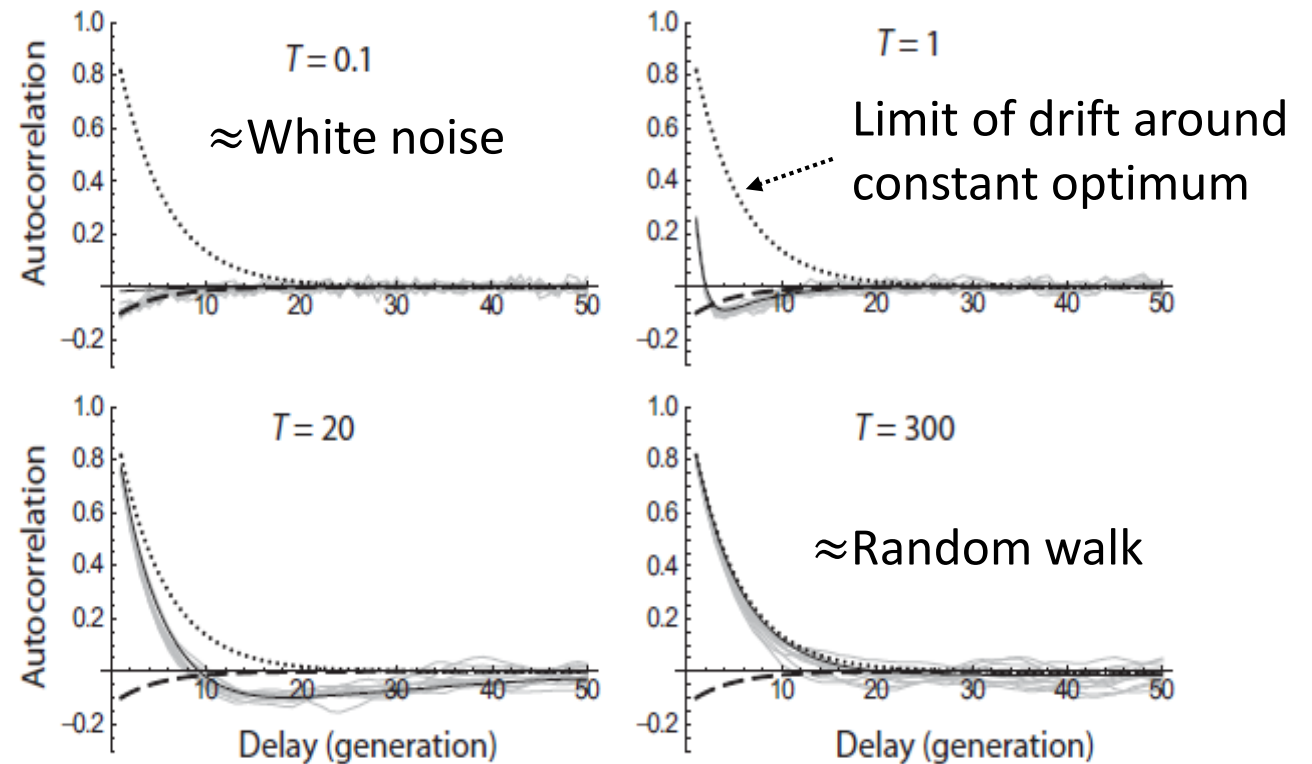
- Autocorrelated fluctuating optimum (AR1), with T the characteristic time over which optimum is autocorrelated

- Without drift: $V(\beta) \approx \frac{S \sigma_{\theta}^2}{1+SGT}$

$$\text{ACF}(\beta, \tau) = \frac{e^{-\frac{\tau}{T}} - SGT e^{-SG\tau}}{1 - SGT}$$

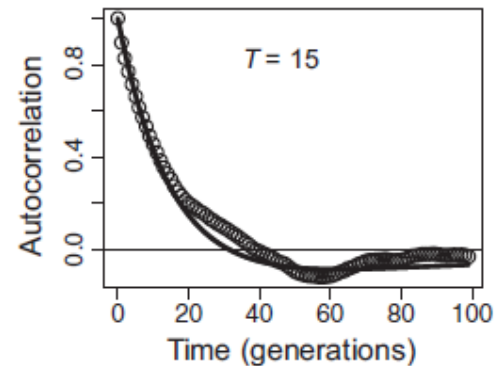
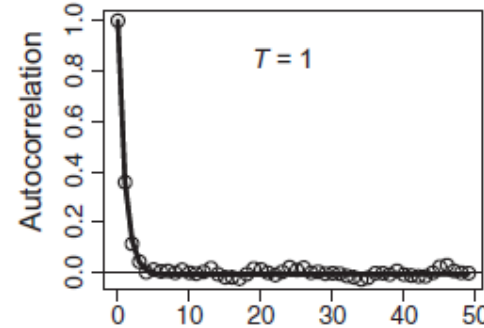
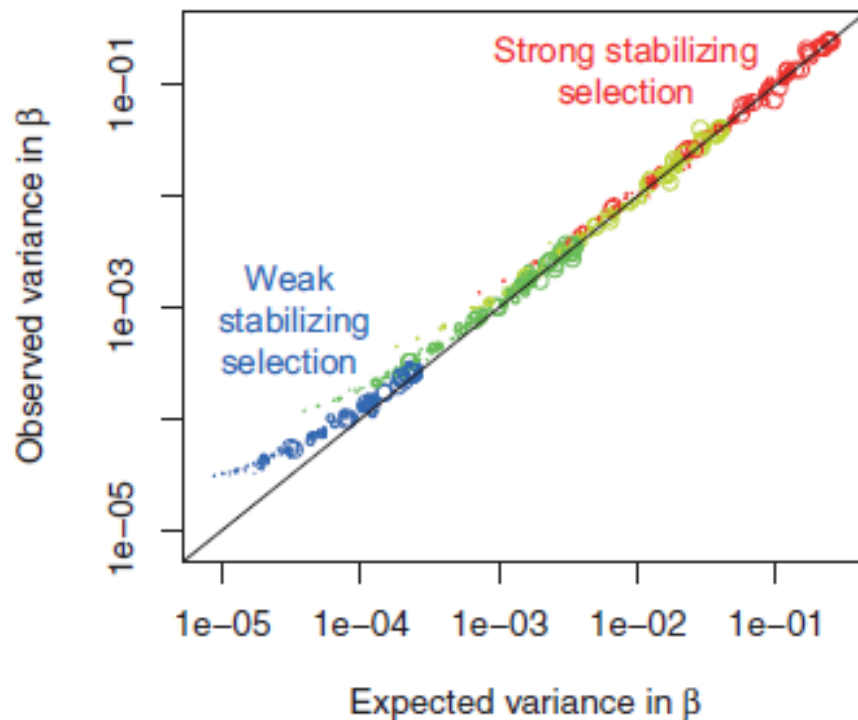
(Weighted) difference between autocorrelation of optimum and evolutionary inertia

→ Fluctuations in β do not tell the whole story about fluctuating selection!



Fluctuations of selection gradient

- Analytical predictions assuming constant genetic variance work well on individual-based simulations with high mutation rates



Population dynamics under moving optimum

- Evolution and demography are **connected through the fitness landscape**¹ relating population mean fitness \bar{W} to the mean phenotype \bar{z}
- Simple discrete-time model:

$$\text{Demography: } \ln N_{t+1} = \ln N_t + \ln \bar{W}_t$$

$$\text{Evolution: } \Delta \bar{z} = G \frac{\partial \ln \bar{W}}{\partial \bar{z}}$$

- With Gaussian fitness peak, **mean mismatch with optimum** drives eco-evo dynamics

$$\text{Demography: } \ln N_{t+1} = \ln N_t + r_{\max} - g(N_t) - \frac{S}{2} (\bar{z}_t - \theta_t)^2$$

$$\text{Evolution: } \Delta \bar{z} = -GS(\bar{z}_t - \theta_t)$$

1 : Wright (1937 PNAS)

Crow & Kimura (1970)

Lande (1976 Evolution, 1982 Ecology)

Population dynamics under moving optimum

- **Neglecting density dependence** (eg under severe stress):

$$n_t = \ln N_t = n_0 + r_{\max} t - \frac{s}{2} \sum_{k=0}^{t-1} (\bar{z}_k - \theta_k)^2$$

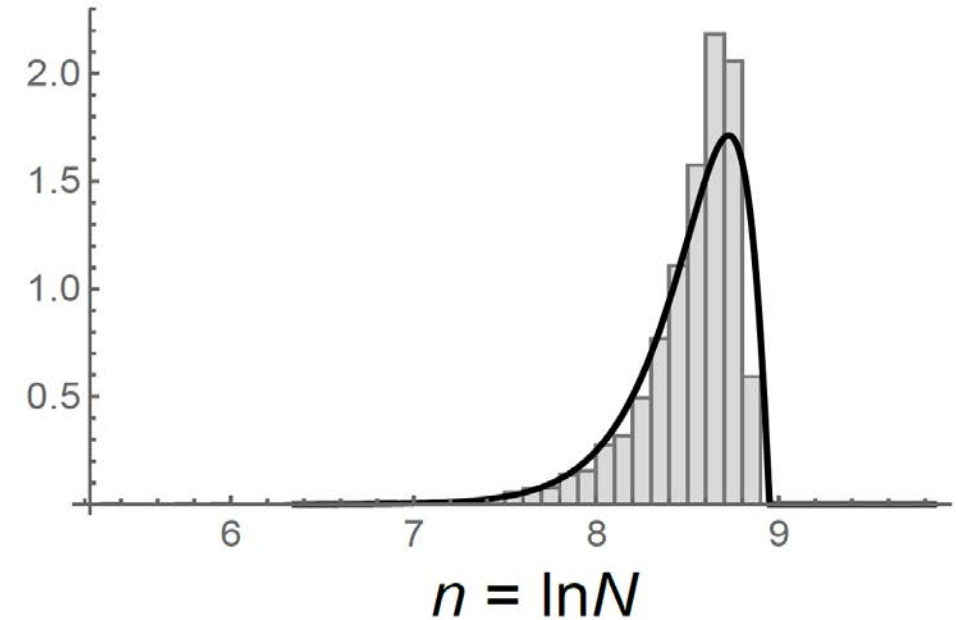
Unweighted sum of all **past maladaptation**

→ Past extreme events may have long-lasting consequences

- If θ is a Gaussian process, so are \bar{z} and $(\bar{z} - \theta)$
Then $n = \ln N$ is related to chi-square,
or gamma distribution with shape parameter increasing with time

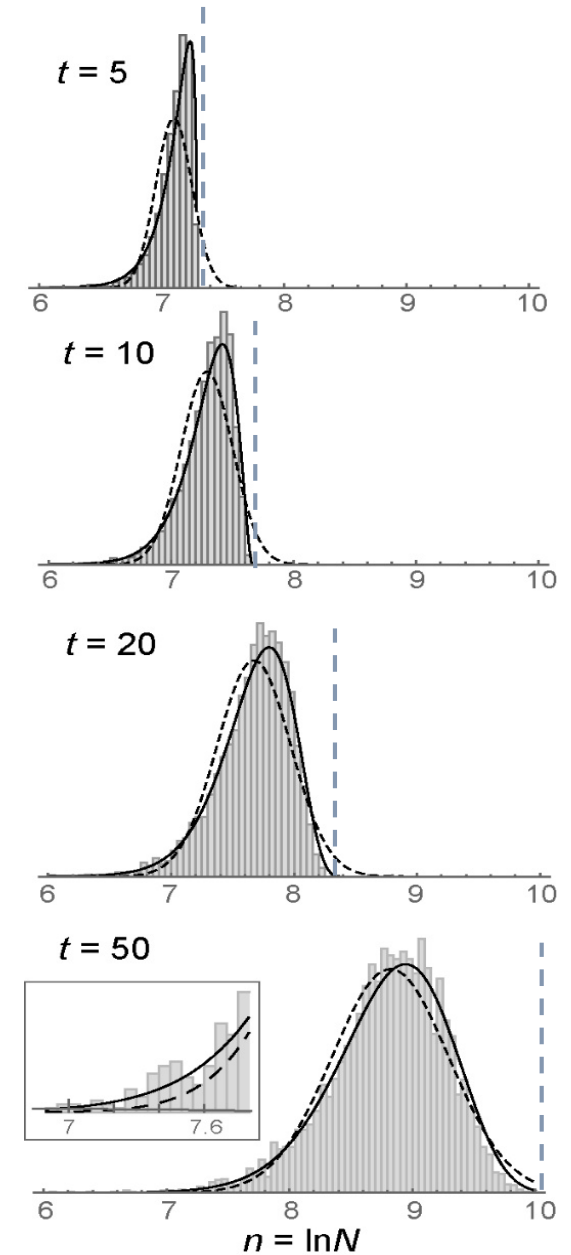
Distribution of population size

- The reverse gamma distribution is:
 - **Bounded above** by growth of optimum phenotype
 - **Left skewed**
 - **excess of small population sizes** at high extinction risk



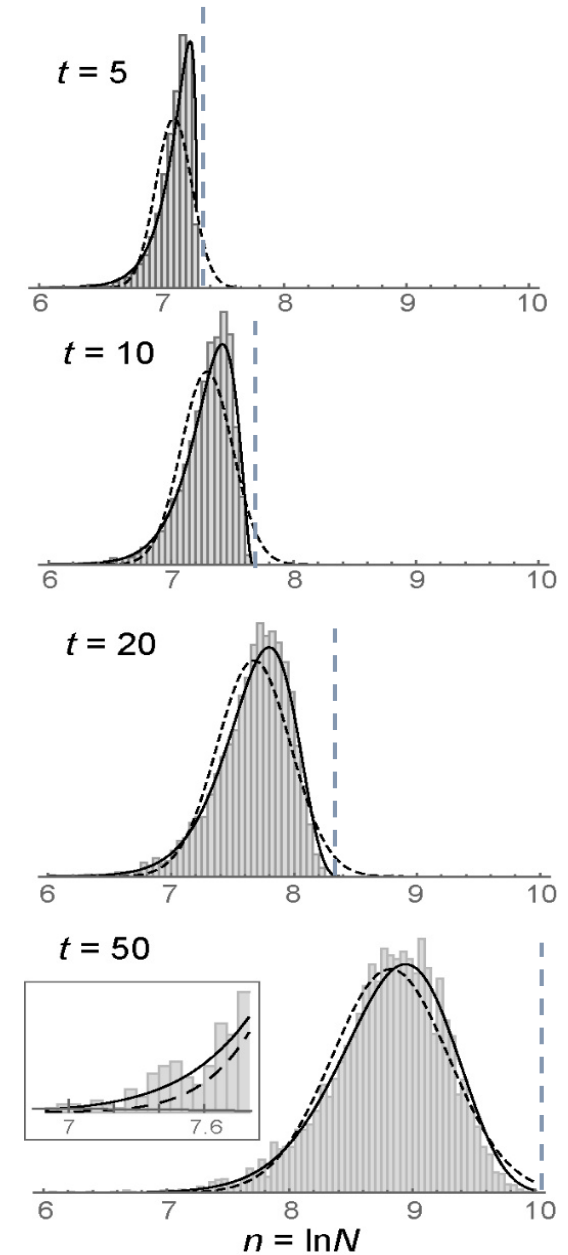
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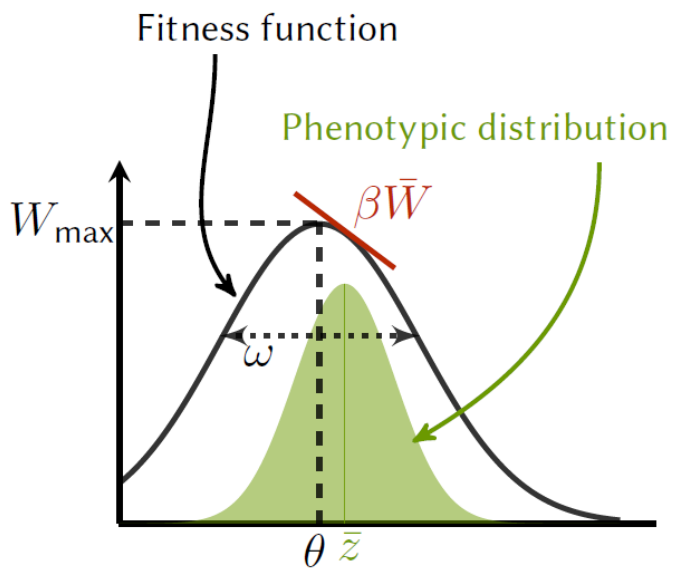
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 - **excess of small population sizes** at high extinction risk
 - Starting from fixed size, **tends to normal** over time, but slowly (excess of small N remains)
 - **Autocorrelation** of optimum :
 - **increases the expected** $\ln N$ (facilitates adaptive tracking^{1,2})
 - increases **variance** of population size (among independent lineages)¹.
 - possibly antagonistic for extinction risk



Conclusion

- Models of adaptation to an optimum phenotype rely on plausible biological assumptions
- They yield predictions about adaptation across a range of conditions (low/high mutation, fitness and traits).
→ Combine several lines of evidence.
- Can help understand adaptation, but only a starting point: reality is more complex! Multiple peaks, frequency dependence (flattening fitness peaks...), space, phenotypic plasticity...

Thanks!
Questions?



Gaussian fitness peak

Recursion for the mismatch with optimum:

$$x_t = \bar{z}_t - \theta_t = \bar{z}_{t-1} - GS(\bar{z}_{t-1} - \theta_{t-1}) - \theta_{t-1} + \theta_{t-1} - \theta_t$$

$$x_t = (1 - GS)x_{t-1} + \theta_{t-1} - \theta_t$$

$$\Delta x = -GSx_{t-1} - \Delta\theta$$

Gaussian fitness peak

- Recursion for mean phenotype \bar{z} :

$$\bar{z}_t = \bar{z}_{t-1} - GS(\bar{z}_{t-1} - \theta_{t-1}) = \bar{z}_{t-1}(1 - GS) + GS\theta_{t-1}$$

$$\bar{z}_1 = \bar{z}_0(1 - GS) + GS\theta_0, \quad \bar{z}_2 = \bar{z}_0(1 - GS)^2 + GS(1 - GS)\theta_0 + GS\theta_1,$$
$$\bar{z}_3 = \bar{z}_0(1 - GS)^2 + GS(1 - GS)^2\theta_0 + GS(1 - GS)\theta_1 + \theta_2 \dots$$

- Full solution for $t \geq 1$ is

$$\bar{z}_t = \bar{z}_0(1 - GS)^t + GS \sum_{k=0}^{t-1} (1 - GS)^{t-1-k} \theta_k$$

Replacing $j = t - k$, such that $k = t - j$

$$\bar{z}_t = \bar{z}_0(1 - GS)^t + GS \sum_{j=1}^t (1 - GS)^{j-1} \theta_{t-j}$$



Directional environmental change

- Highly polymorphic regime:

- Recursion for distance to optimum $x = \bar{z} - \theta$:

$$x_t = (1 - GS)x_{t-1} - v$$

$$x_0 = 0, x_1 = -v, x_2 = -v(1 + (1 - GS)), x_3 = -v[1 + (1 - GS) + (1 - GS)^2], \dots$$

- Full solution for $t \geq 1$ is

$$x_t = -v \sum_{k=0}^{t-1} (1 - GS)^k = -v \frac{1 - (1 - GS)^t}{1 - (1 - GS)} = -\frac{v}{GS} [1 - (1 - GS)^t]$$

$\Delta \bar{z} = G\beta = -GSx$

- At equilibrium

$$x_{\text{eq}} = (1 - GS)x_{\text{eq}} - v \iff x_{\text{eq}} = -\frac{v}{GS}$$



Cycling environment

- **Model**¹: Sine wave with amplitude A and period T

$$\theta = A \sin\left(\frac{2\pi t}{T}\right)$$

- Continuous-time evolutionary dynamics

$$\frac{d\bar{z}}{dt} = -GS \left[\bar{z} - A \sin\left(\frac{2\pi t}{T}\right) \right]$$

$$\frac{d\bar{z}}{dt} + GS\bar{z} = A \sin\left(\frac{2\pi t}{T}\right)$$

