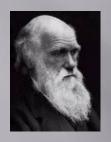
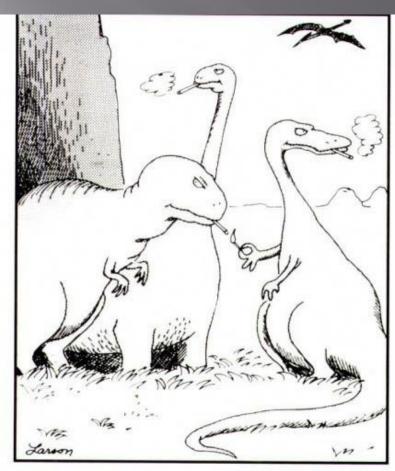
Neutral dynamics and family (type) statistics :











The real reason dinosaurs became extinct

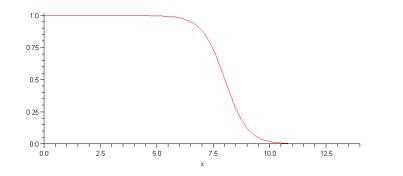
Nadav Shnerb (BIU) Yosi Maruvka (BIU) David Kessler (BIU)

Gur Yaari (Yale) Sorin Solomon (HUJI) Robert Ricklefs (St. Louis) <u>John Wakeley (</u>Harvard) PLoS computational biology 5(4): e1000359 (2009)

What is the difference between pants and glasses? Survival of the fittest Survival of the luckiest

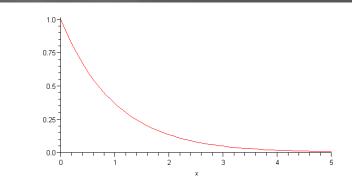


Pants "die" (ripped) due to accumulated wear.





Glasses "die" when they shattered, usually due to random and uncorrelated events



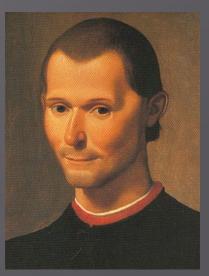
What is the signal to noise ratio in complex system? Social dynamics:

The prince, Capitolo XXV

[What fortune can effect in human affairs and how to withstand her]

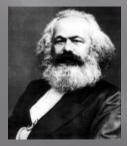
"Fortune is the mistress of one half our actions, and yet leaves the control of the other <u>half</u>, or a little less, to ourselves"





Stochastic (neutral)

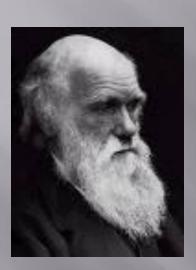
Deterministic (fitness)



it is the ultimate aim of this work, to lay bare <u>the</u> <u>economic law of motion of</u> <u>modern society</u> Karl Marx, Das Capital (introduction) Cleopatra's nose, had it been shorter, the whole face of the world would have been changed (Pascal)



Ecology and evolution: Neutral vs. adaptive (niche) theories



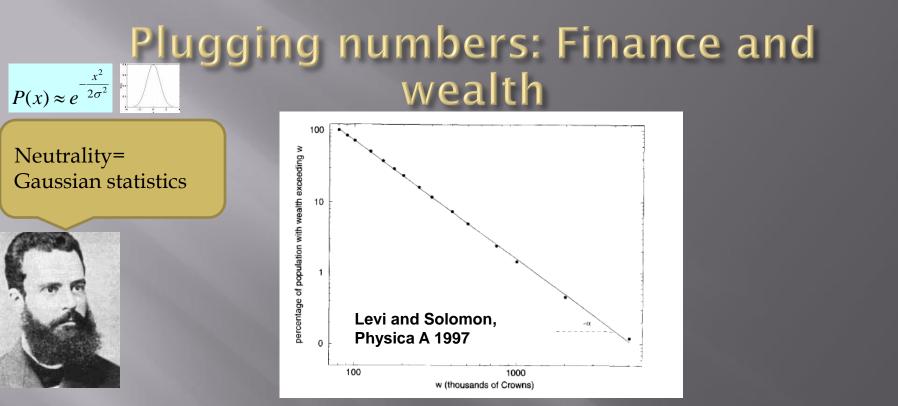




"When we look at the plants and bushes clothing an entangled bank, we are **tempted** to attribute their proportional numbers and kinds to what we call **chance.** But how **false** a view is this!" Darwin On the Origin of Species (1859) Hubbell started from a postulate that most consider preposterous: that one tree or one bird is just like any other. His patterns result **solely from random fluctuations** in births, deaths and the arrival of new species. Scientific American April 29 2002.

Deterministic (fitness)

Stochastic (neutral)



Pareto: Saying "everything is governed by chance" is a prediction. The central limit theorem guarantees that the resulting statistics will be Gaussian.

$$x_{t+1} = x_t + \eta_t$$

$$\eta \in [-1,1]$$

$$P(x) \approx e^{-\frac{x^2}{\sigma^2(t)}}$$

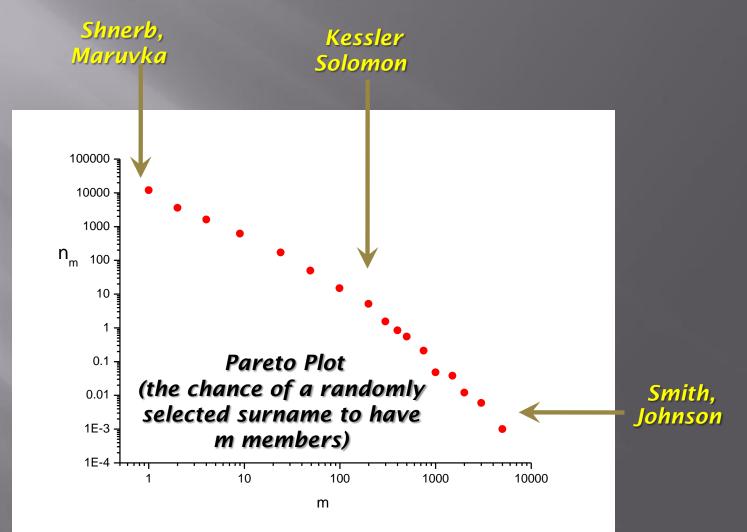
 $x_{t+1} = x_t \eta_t$ $\eta \in [1/2, 2]$ $\log(x_{t+1}) = \log(x_t) + \log(\eta_t)$

$$P(x) \approx e^{-\frac{\left[\log(x)\right]^2}{\sigma^2(t)}} \approx x^{-\frac{\log(x)}{\sigma^2(t)}}$$

Stochastic (neutral)

Deterministic (fitness)

example for neutral dynamics: surnames



The underlying stochastic process: 💑 (ule 1925) a special preference

for beetles ... here they are:



 $n_m \approx m^{-\alpha}$ $\log(n_m) \approx -\alpha \log m$



Genus = surname Species = first name Life

Domain

Kingdom

Phylum

Class

Order

Family

Genus

Species

0,00

3360

of species in genus varies tremendously

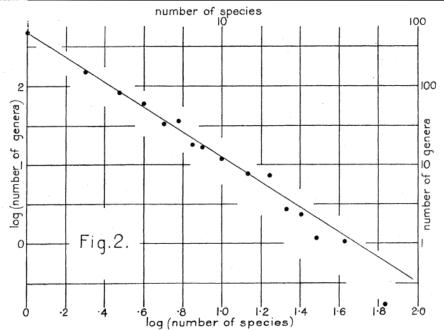
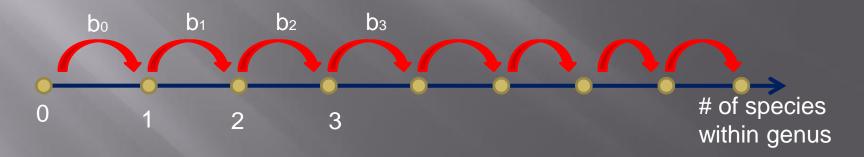


Fig. 2.-Double logarithmic chart for the frequency distribution of sizes of genera in the Cerambucina. Data in Appendix, Table B.

Modeling neutral processes : The Yule process (preferential attachment)

No fitness, no selection, everybody are equal

- **1.** Birth=speciation: A species is chosen at random to reproduce = to undergo speciation (no fitness, selection, whatever).
- **2.** Mutation = speciation that creates new genus: an offspring is identical to its father with probability $1-\mu$ and is a mutant (new species, surname, genus) with probability μ .

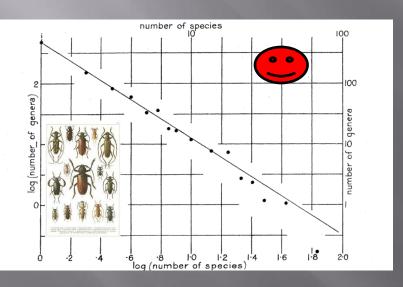


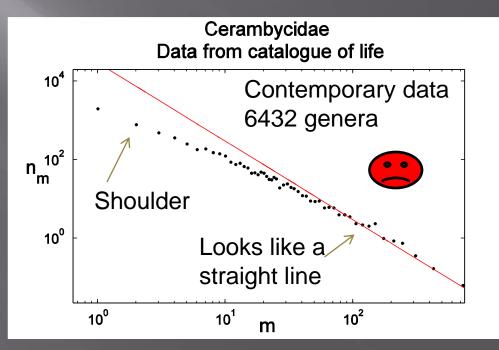
 $b_n = \gamma(1-\mu)n$ $(n \ge 1)$ $b_0 = \mu\gamma \cdot total population$ Where γ is the birth rate

Yule-Simon steady state

$$n_m = \left(1 + \frac{1}{\mu}\right) \frac{\Gamma(m)\Gamma(2 + 1/\mu)}{\Gamma(m + 2 + 1/\mu)} = \left(1 + \frac{1}{\mu}\right) B(m, 2 + 1/\mu) \approx m^{-\left(2 + \frac{1}{\mu}\right)}$$

Original Yule 1024 genera





JTB 262 245 (2010)

Beyond Yule: The Birth-Death-Mutation (BDM) process

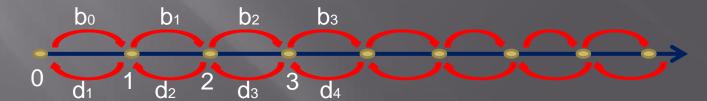
Moran Version ("overlapping generations")

1. Birth: choose a random agent to reproduce.

2. Mutation: an offspring is identical to its "mother" with probability $1-\mu$ and is a mutant (new genus, surname, allele) with probability μ .

3. Death: a random agent is chosen to die with probability δ .

Important parameters are μ and the growth rate $\gamma = 1-\delta$



JTB **262** 245 (2010)

Universality

The results are independent of the microscopic details of the process (overlapping or nonoverlapping generations, the order of events, IF

μ, **γ** <1.

Under these conditions the number of families of size m, n_m, varies only slightly with m so for both Moran and Wright-Fisher process, independent of their **details**, the family statistics is described by a universal differential equation, the Kummer equation:

$$\frac{\partial n_m}{\partial t} = (\mu - \gamma) \frac{\partial}{\partial m} (m n_m) + \frac{\sigma^2}{2} \frac{\partial^2}{\partial m^2} (m n_m)$$

$$\begin{split} \gamma > \mu & n_m = \frac{\nu R_c \Gamma(2+\nu)}{m} U \Big(1 + \nu, 0, \frac{R_c}{N} m \Big) \\ \gamma < \mu & n_m = \frac{R_c \Gamma(1+\nu)}{m} U \Big(\nu, 0, \frac{R_c}{N} m \Big) e^{-\frac{R_c}{N} m} \end{split}$$

Large m asymptotic

$$\gamma > \mu$$
$$n_m \approx m^{-1 - [\gamma/(\gamma - \mu)]}$$

$$\gamma < \mu$$

$$n_m \approx m^{-1 - \frac{\gamma}{\mu - \gamma}} e^{-(\mu - \gamma)m}$$

First presented by Manrubia and Zannette, JTB 216 461 (2002), in the context of Moran process

Effect of sampling

Let us assume that only R_0 individuals were sampled out of the total population of size N_0 .

Hypergeometric distribution

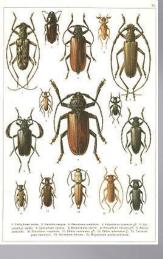
$$n_m^R = \sum_{p \ge m} n_p \frac{\binom{p}{m} \binom{N_o - p}{R_o - m}}{\binom{N_o}{R_o}}$$

Effective sampling strength

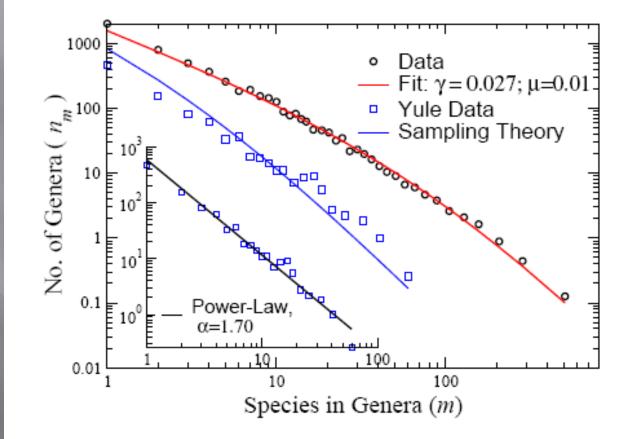
$$\nu \equiv \frac{\mu}{\gamma - \mu} \qquad s \equiv \frac{\sigma^2 (1 + \nu)}{2\gamma} \frac{R_0}{N_0}$$

 $n_m^R \approx \nu R_c B (2 + \nu, m) s^m {}_2F_1 (m, m + 1; m + 2 + \nu; 1 - s)$

Weak sampling \longrightarrow back to Yule-Simon



Back to the bugs

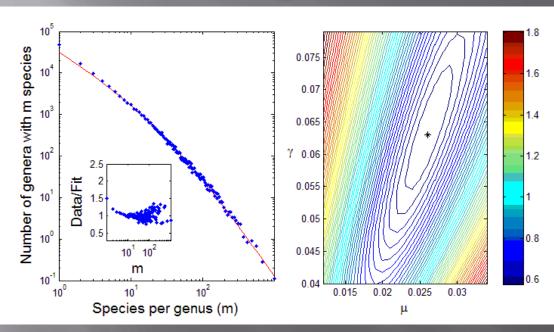


Applications Species within genera: animalia

•2 parameters fit.

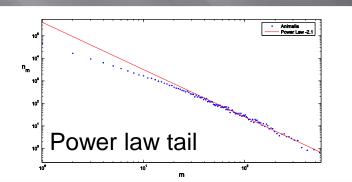
•Logarithmic binning.

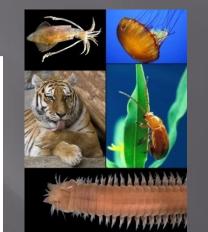
• PDF (NOT cumulative distribution)

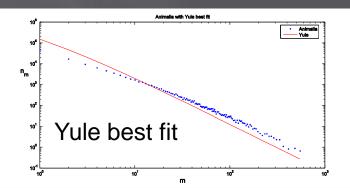


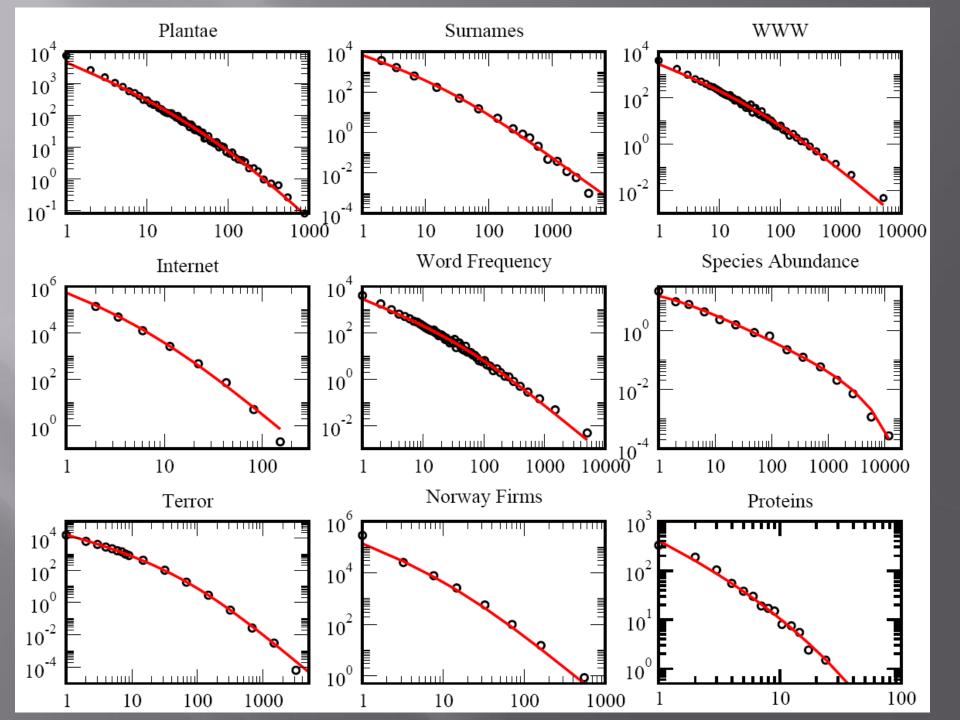
www.catalog ueoflife.org

Species 2000 & ITIS Catalogue of Life: 2009 Annual Checklist









Inferring historic demographic parameters from current polymorphism (SNP) data

J. Stat. Phys. **142** 1302 (2011) Molecular Biology and Evolution **28** 1617 (2011)

The data

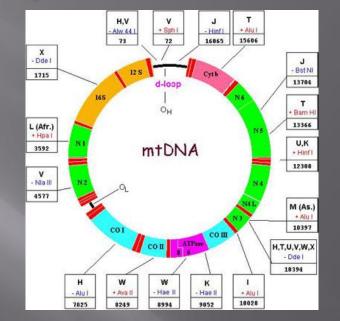
1212 sequences (China)

µ=0.00241/sequence/generation

The challenge:

Infer the growth rate γ

and the effective population size N_0





Haplotype (family) statistics:

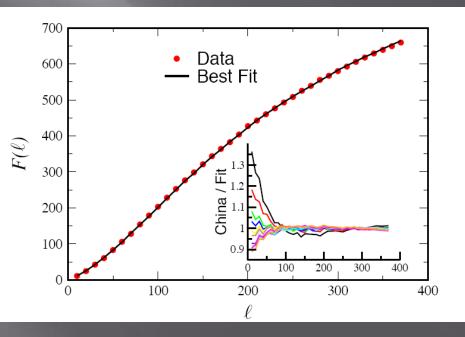
Any sequence inherited along maternal lineages is like a surname.

These surnames are subject to a birth-deathmutation process.

Ends up with haplotype (surname, family) statistics: the number of occurrences of singletons, doubletons and so one.

This statistics depends on the demographic parameters, thus in principle we can retrieve this information from it.

		Loci						
		1	2	3	4	5	6	7
Individual	α	Α	G	С	Т	Α	G	С
	eta	Α	Α	С	Т	Α	G	С
	γ	Α	G	С	А	Α	G	С
	δ	Α	G	Т	Т	Α	G	С
Π	ϵ	G	G	Т	Т	Α	G	Т



Conclusions

•The birth-death-mutation process yields the universal distribution function . It deviates strongly from Yule-Simon for small families since families cannot die in Yule.

• Yule is the weak sampling limit of BDM.

• One can use the BDM distribution in order to retrieve demographic parameters like growth rate in the prehistoric times.

Neutral theory, King Solomon Version

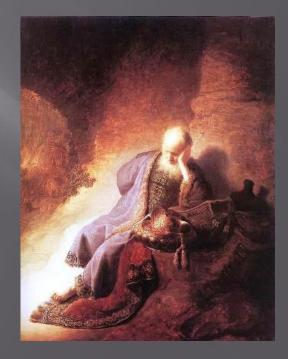
Ecclesiastes (Qoheleth), Chapter 2:

14: The wise man's eyes are in his head; but the fool walketh in darkness: and I myself perceived also that <u>one event happeneth to them all</u>.

15: Then said I in my heart, As it happeneth to the fool, so it happeneth even to me; and **why** was I then more wise?

יד הֶחָכָם אֵינִיו בְּרֹאשׁוֹ, וְהַפְּסִיל בַּחֹשֶׁך הוֹלֵך ; וְיָדַעְתִּי גַם-אָנִי, שֶׁמְקָרֶה אֶחָד יִקְרֶה אֶת-כֵּלָם.

טוּ וְאָמַרְתִּי אֲנִי בְּלִבִּי, כְּמִקְרֵה הַכְּסִיל גַּם-אֲנִי יִקְרֵנִי, וְלָמָה חָכַמְתִּי אֲנִי, אָז יֹתֵר; וְדִבַּרְתִי בְלִבִּי, שֶׁגַּם-זֶה הָבֶל.



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Благодаря Shukriya	Спасибо	ດລາ ດ3/ທ
тетсі беаисоир	вначения	شكرا