

Modeling microbial diversity

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How can we explain species' diversity?

An astonishing characteristic of life is its great variety:

- In tropical rainforests more than 300 tree species may be found on a single hectare.
- In one gram of soil the number of distinct microbial genomes has been estimated at ~ 2000 -- 18,000.

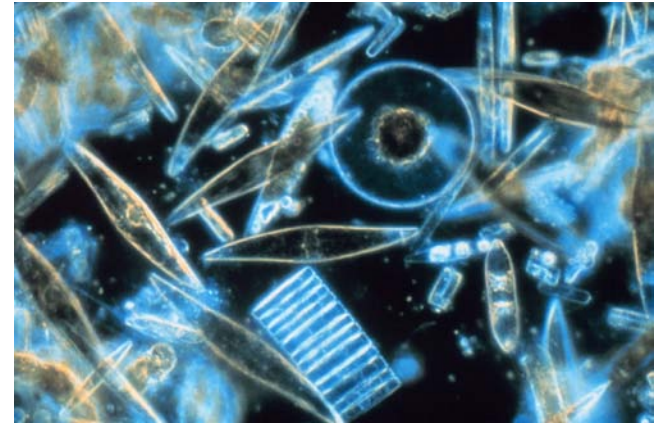
But...the competitive exclusion principle says:

- Two species that compete for the same limiting resource cannot stably coexist.
- In resource competition models, the number of species coexisting in equilibrium cannot exceed the number of resources.

Paradox of the plankton

Originally described by G. E. Hutchinson in 1961:

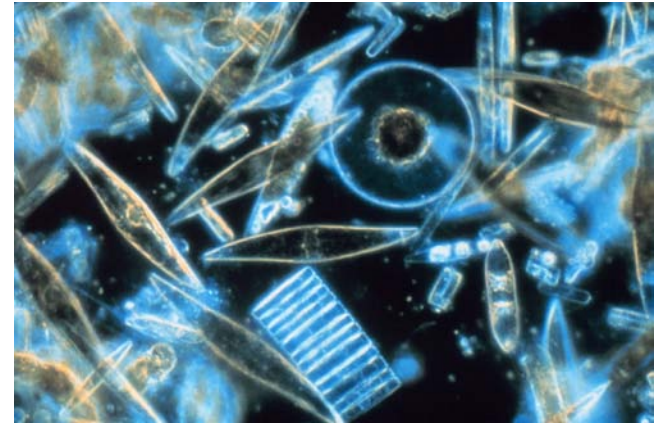
“...a limited range of resources supports an unexpectedly wide range of plankton species, apparently flouting the competitive exclusion principle...”



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Possible solutions:

- Oscillatory or chaotic population dynamics
- Temporal variation of environment, e.g. weather changes, seasonal cycles
- Spatial variation of environment, e.g. gradients such as temperature, salinity, exposure to light
- Other limiting factors, e.g. predation

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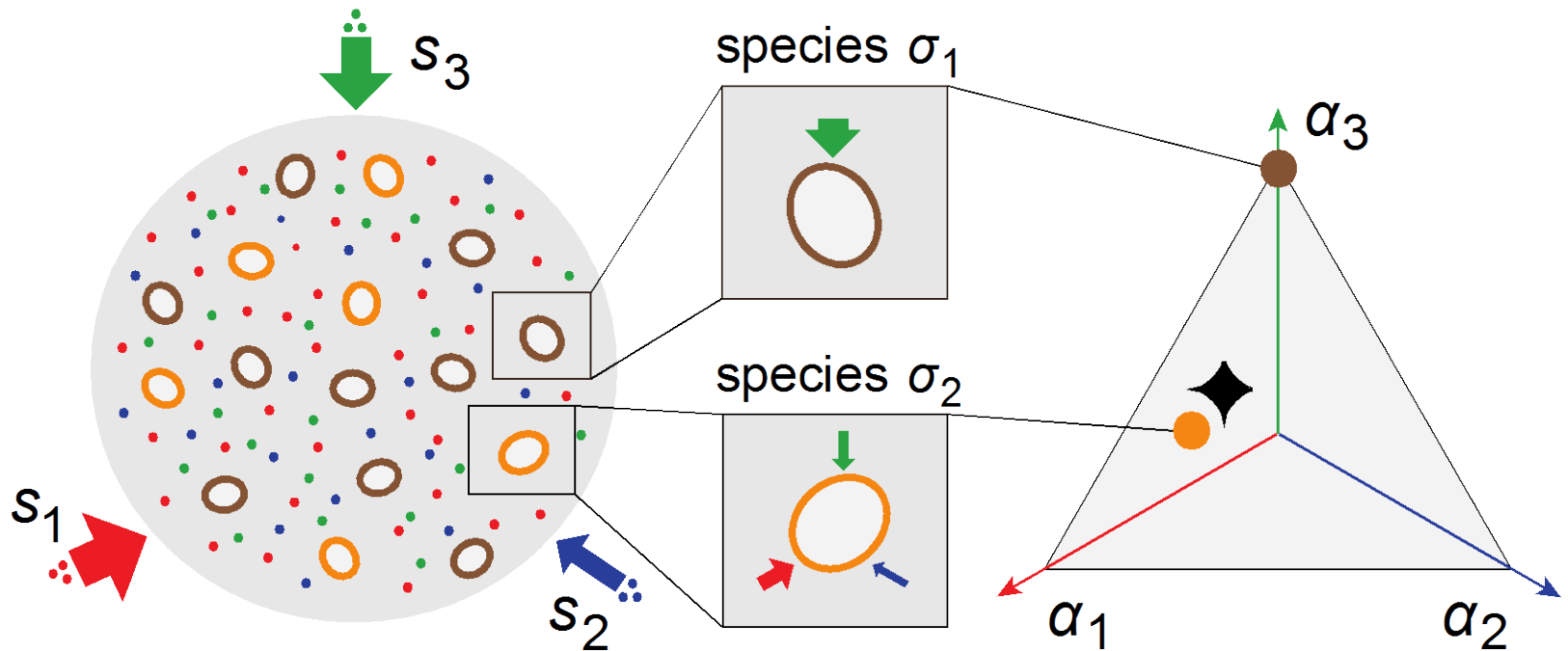
Resource-competition model with trade-offs

p resources

Species: $\vec{\alpha}_\sigma = (\alpha_{\sigma 1}, \dots, \alpha_{\sigma p})$

Trade-offs in ability to utilize different resources:

$$\sum_{i=1}^p w_i \alpha_{\sigma i} = E$$



Resource-competition model with trade-offs

Nutrient concentrations dynamics:

$$\dot{c}_i = s_i - \sum_{\sigma} n_{\sigma} \alpha_{\sigma i} \frac{c_i}{K_i + c_i} - \mu_i c_i$$

= supply – consumption – loss

Growth rate of species σ :

$$g_{\sigma}(\vec{c}) = \sum_{i=1}^p v_i \alpha_{\sigma i} \frac{c_i}{K_i + c_i}$$

Population dynamics:

$$\dot{n}_{\sigma} = (g_{\sigma}(\vec{c}) - \delta) n_{\sigma}$$

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Simplified parameters:

- no nutrient loss: $\mu_i = 0$
- separation of time scales: $\dot{c}_i = 0$
- “symmetric” nutrients: $w_i = K_i = v_i = 1$

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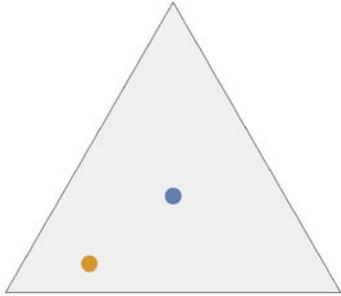
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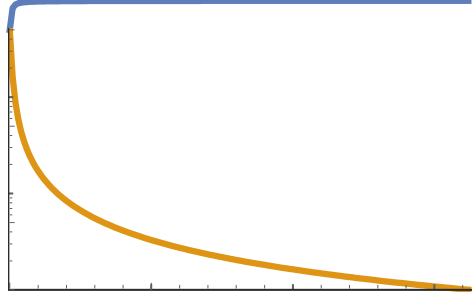
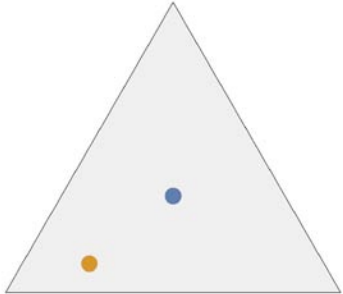
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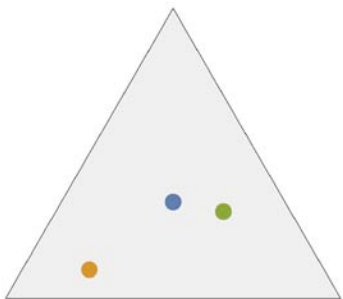
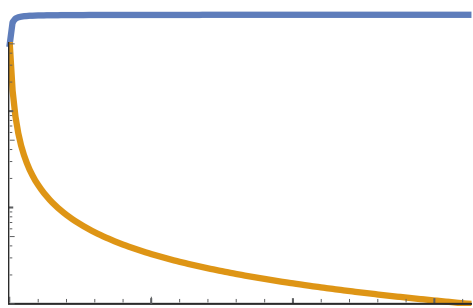
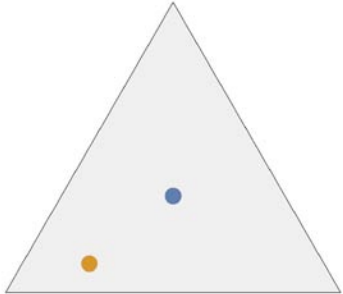
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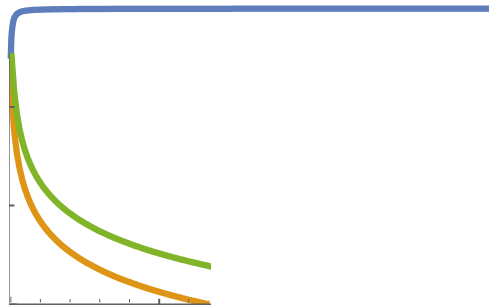
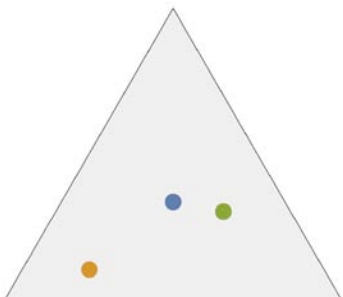
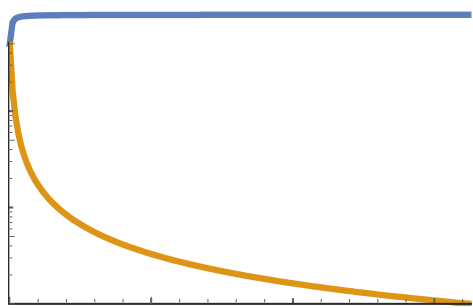
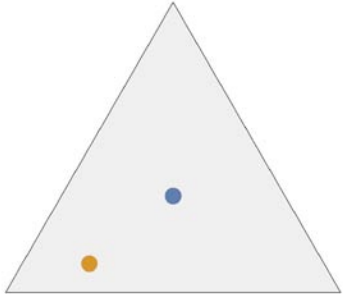
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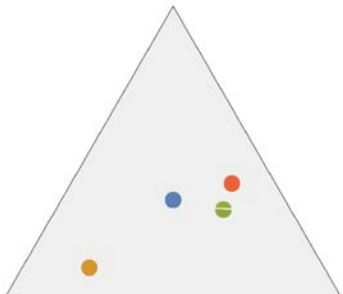
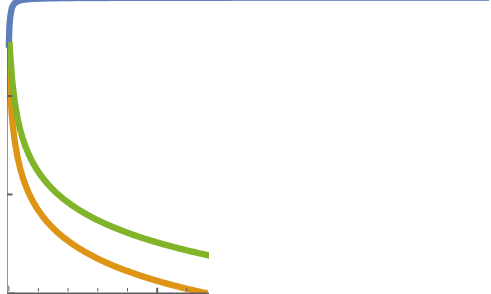
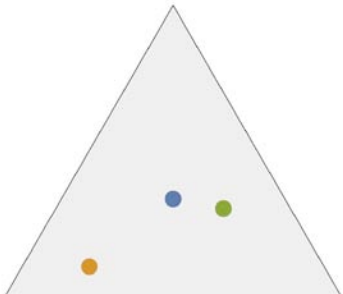
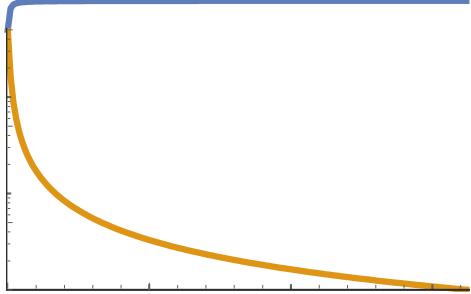
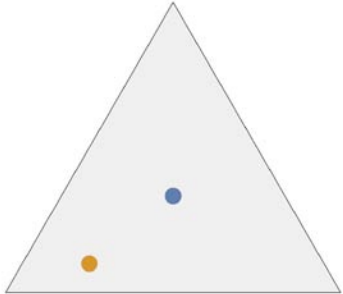
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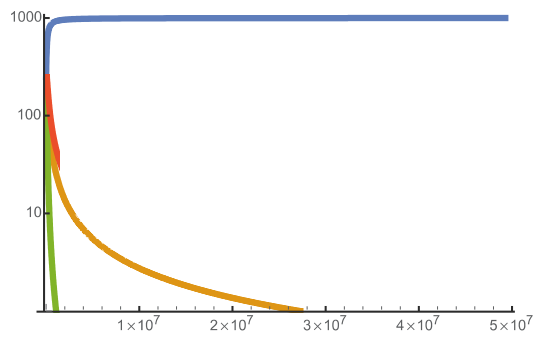
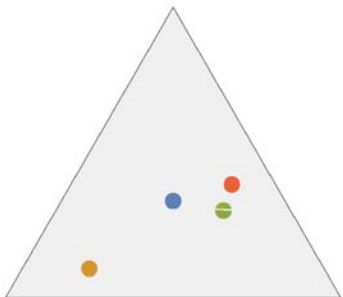
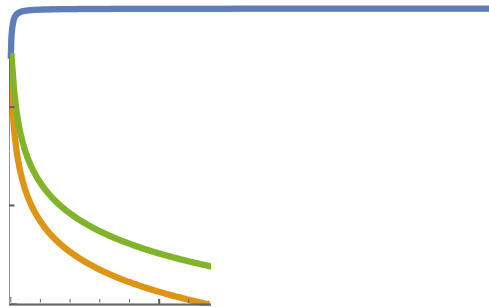
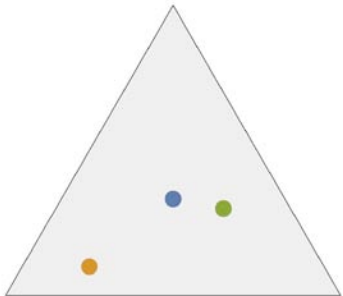
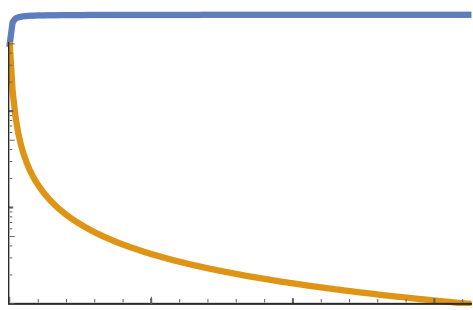
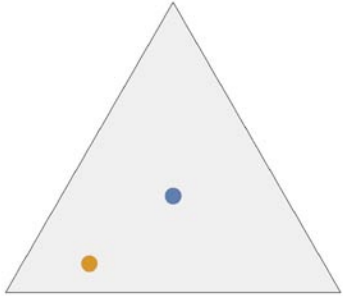


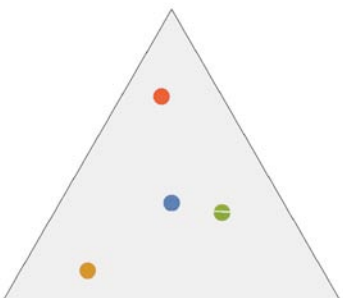
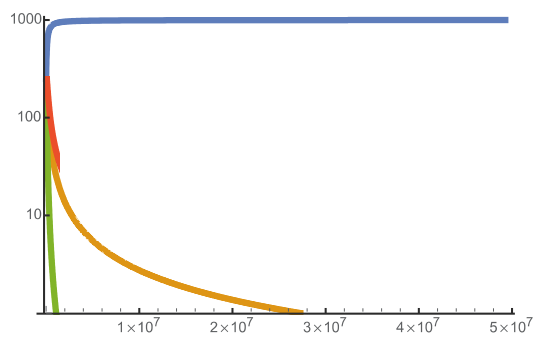
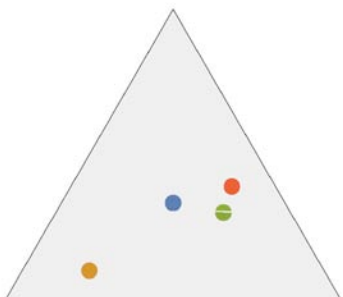
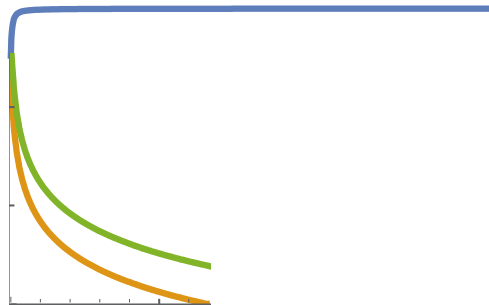
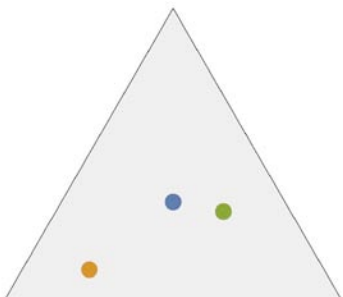
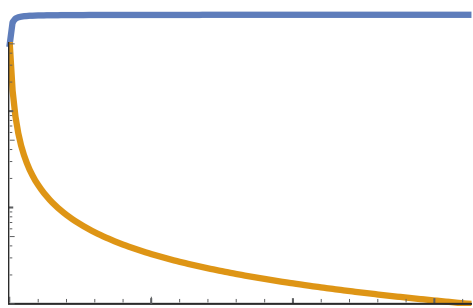
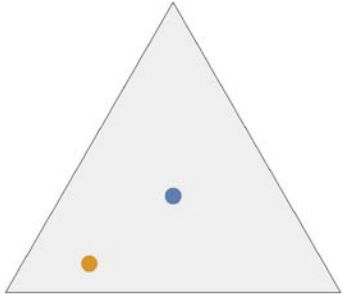


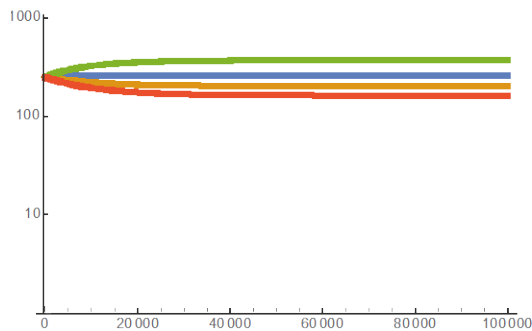
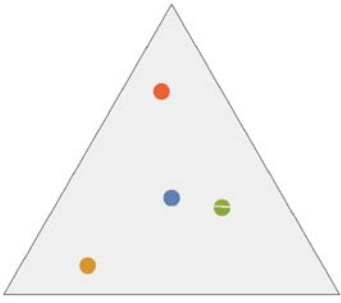
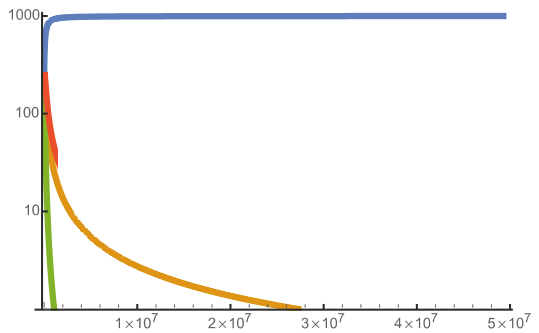
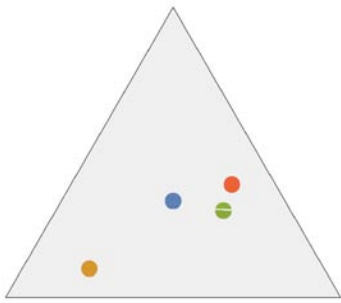
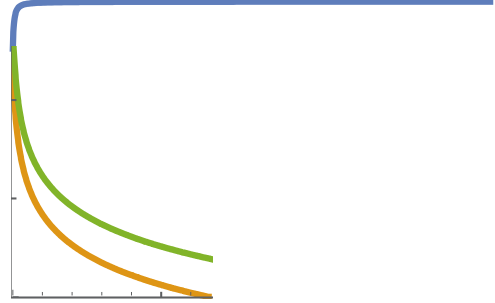
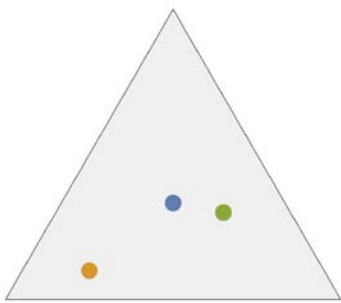
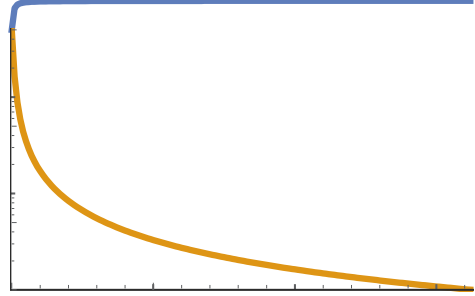
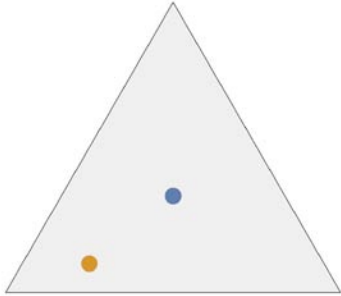


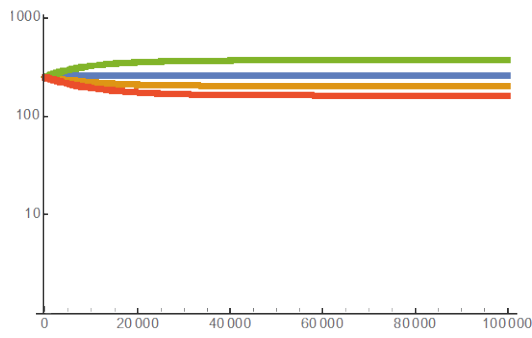
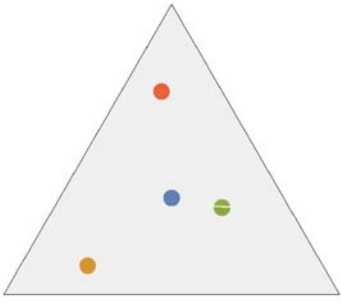
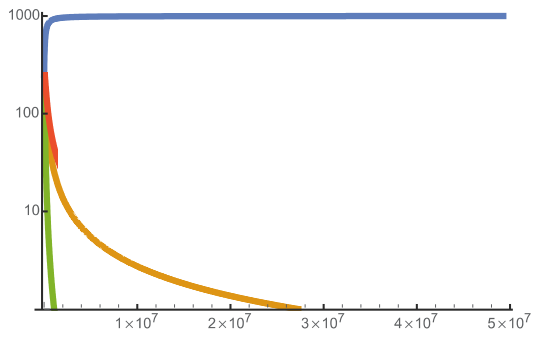
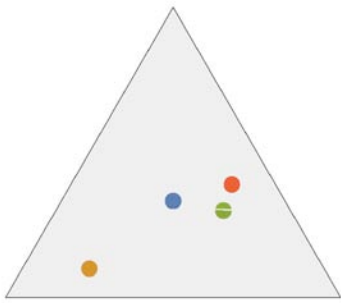
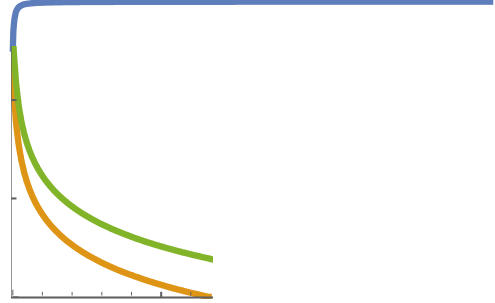
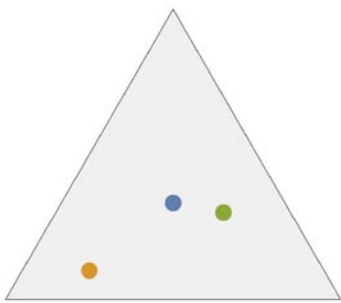
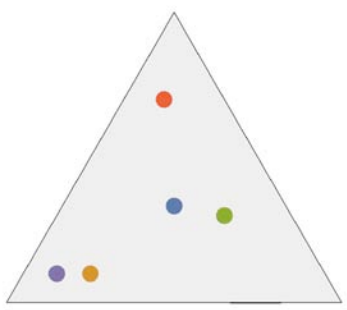
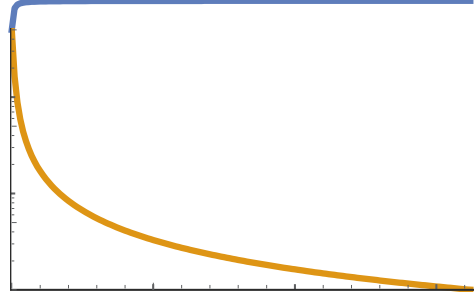
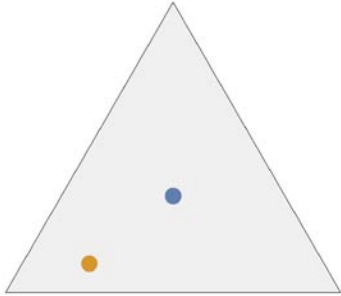


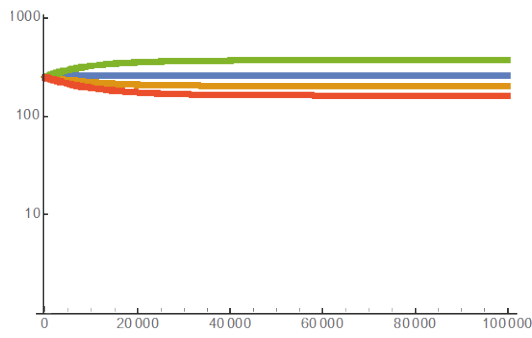
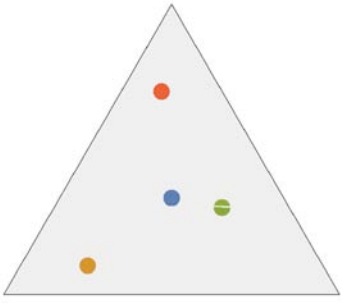
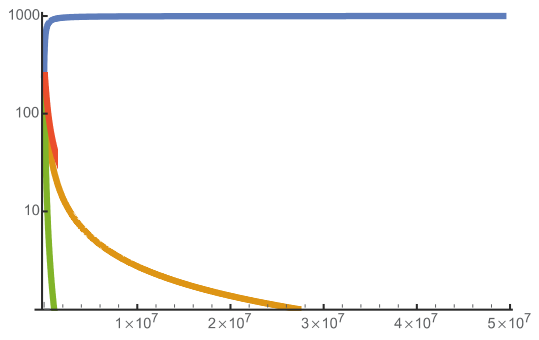
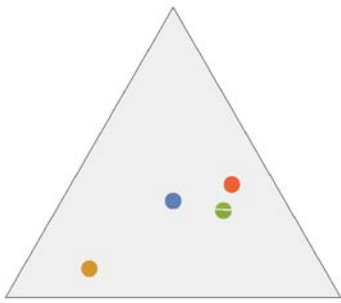
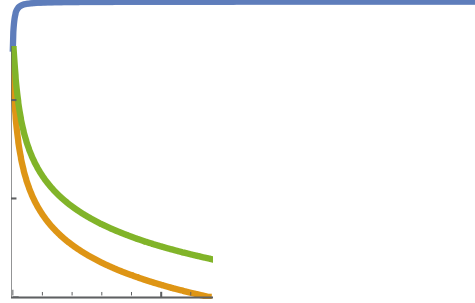
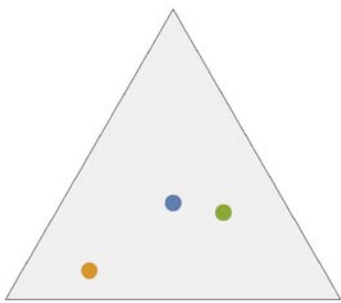
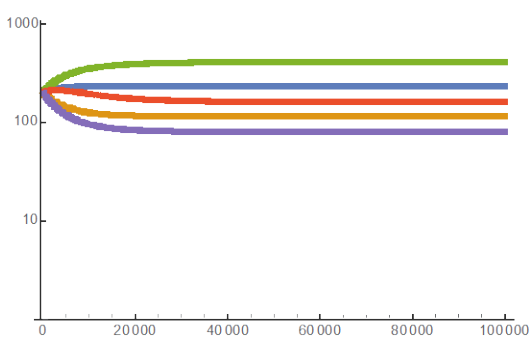
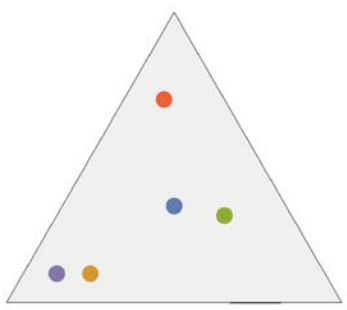
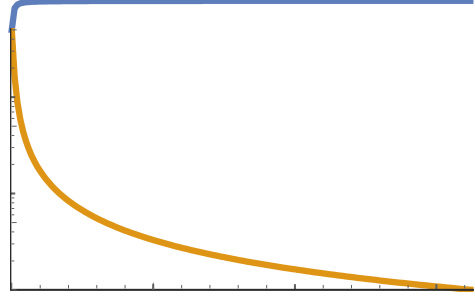
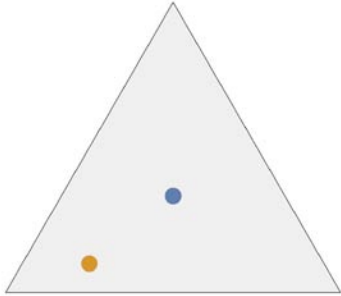


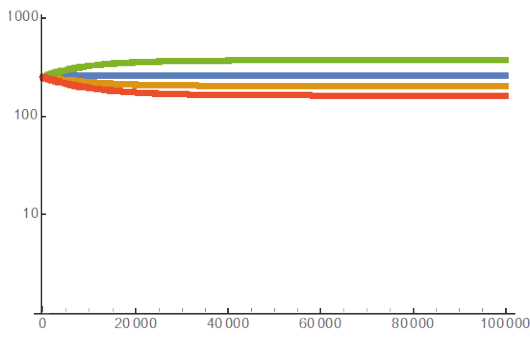
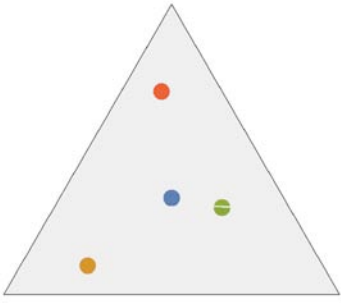
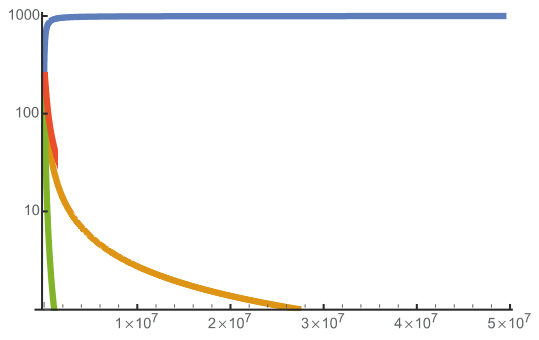
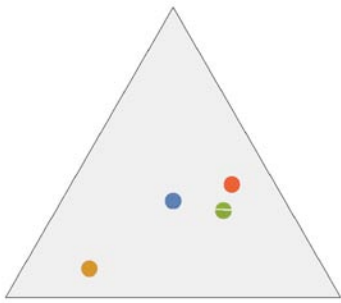
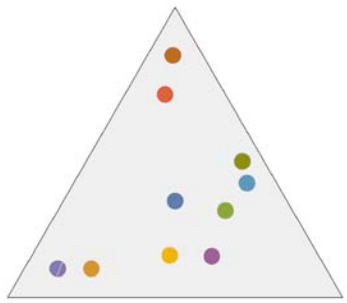
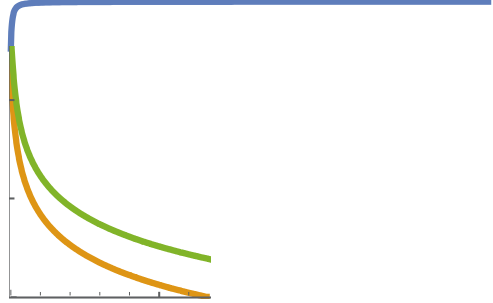
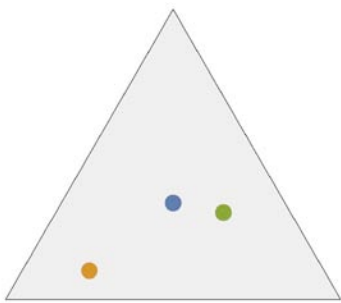
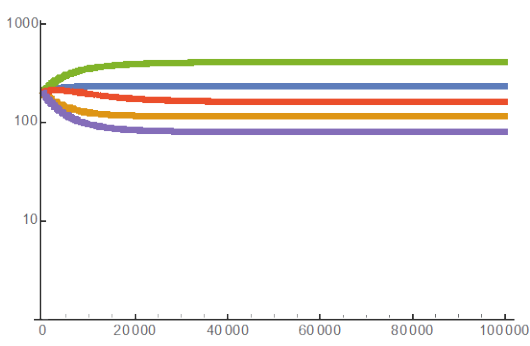
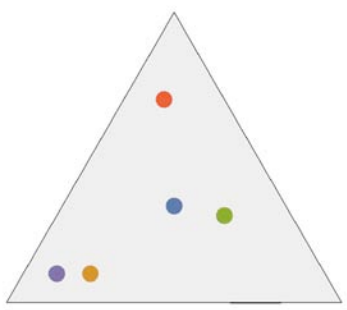
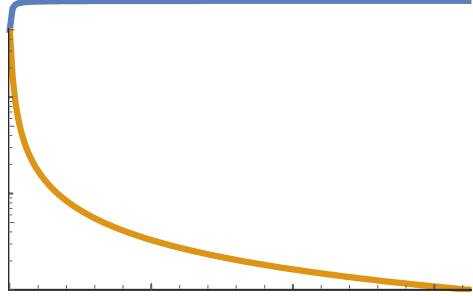
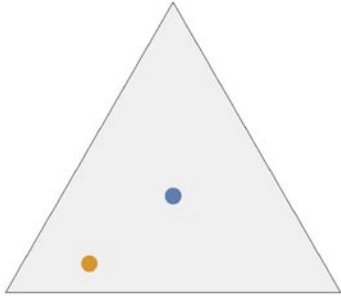


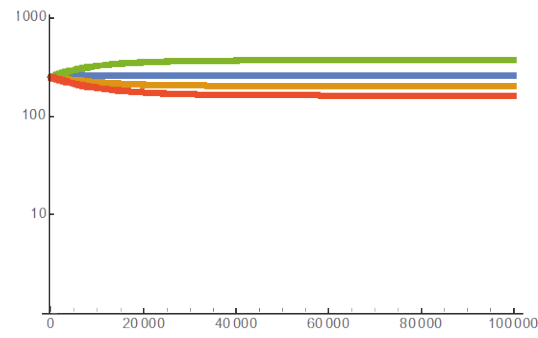
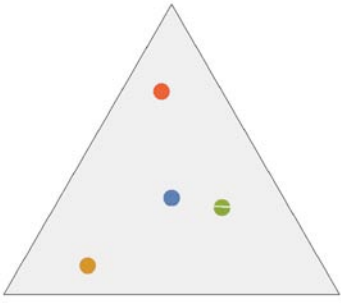
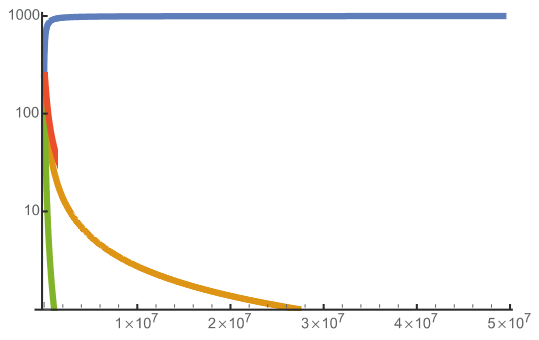
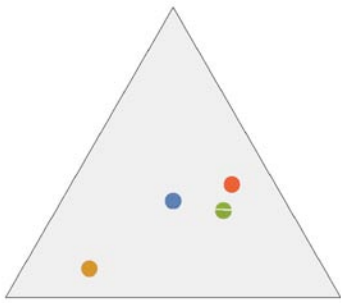
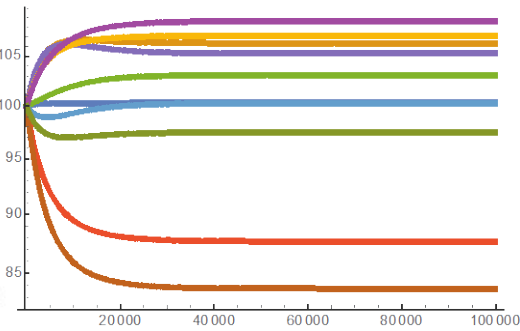
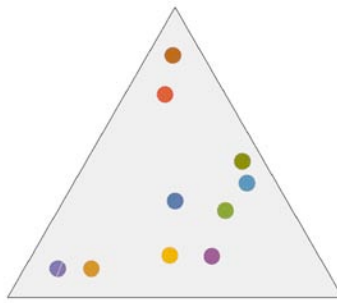
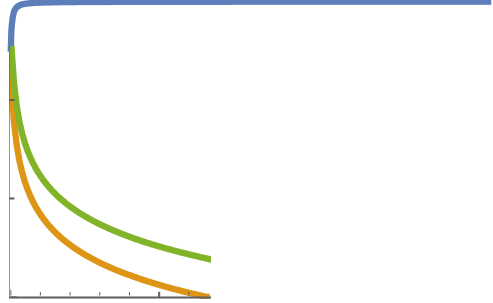
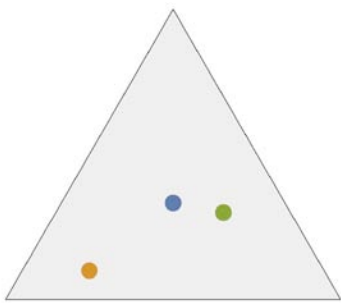
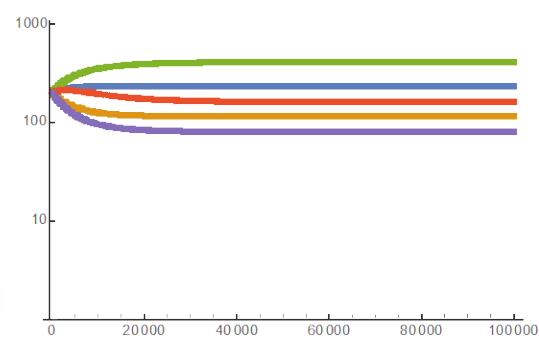
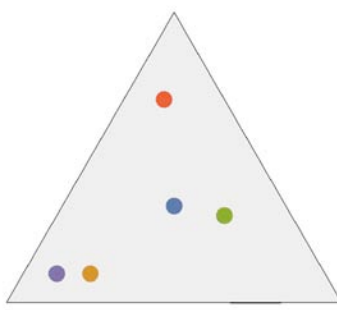
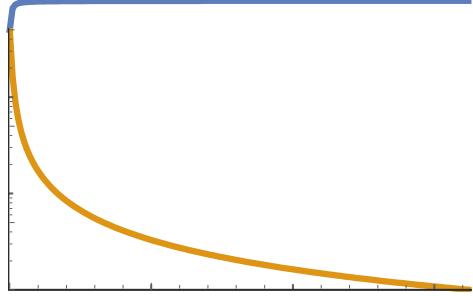
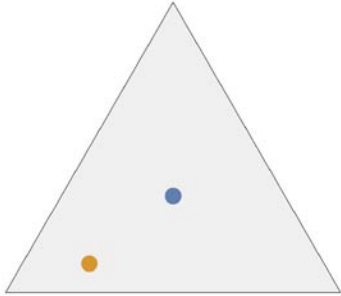


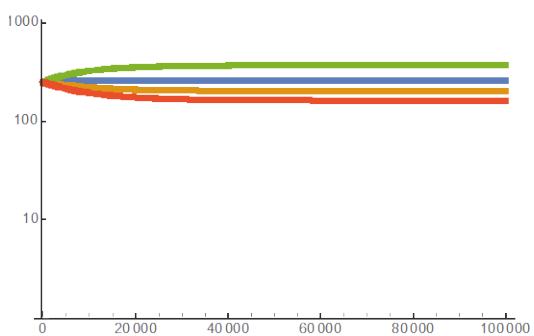
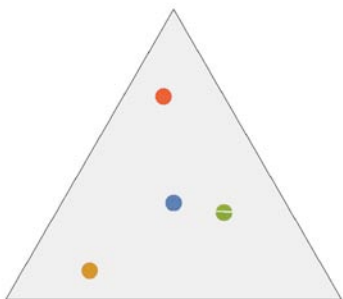
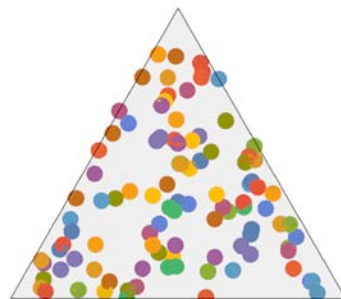
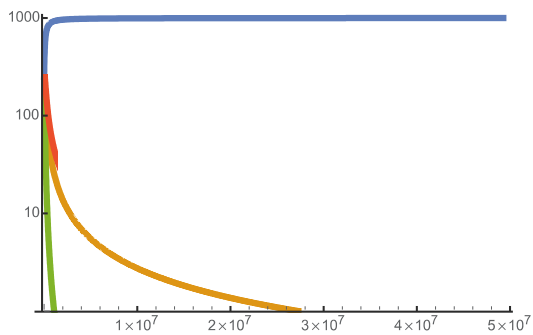
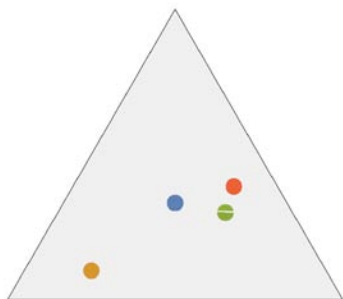
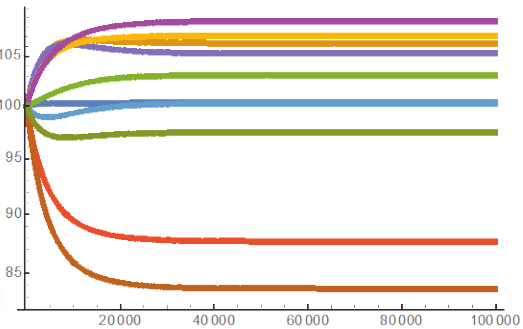
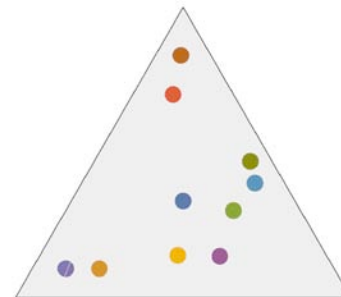
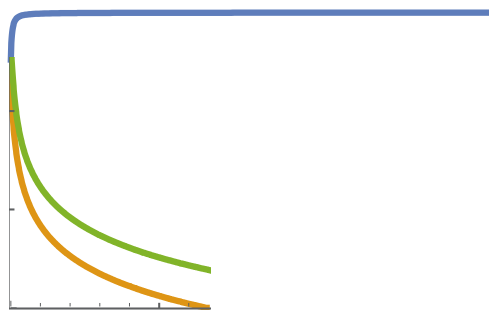
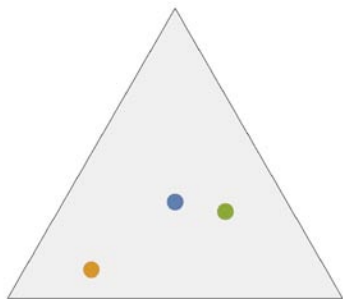
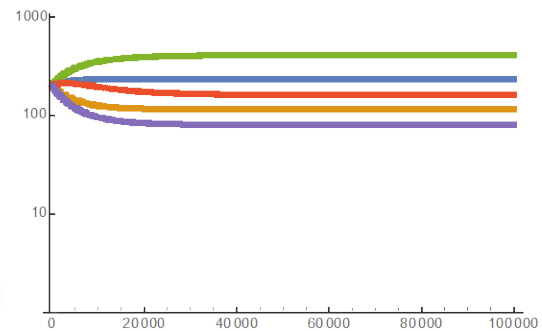
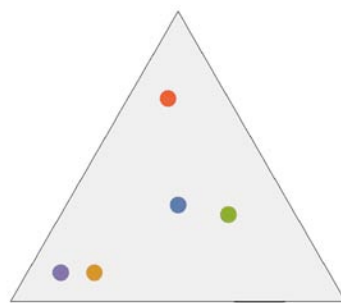
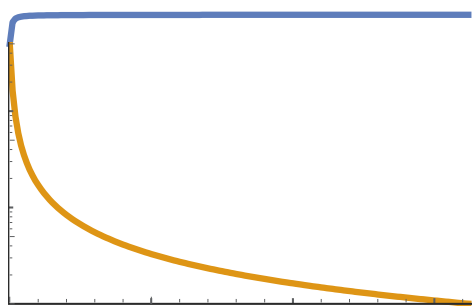
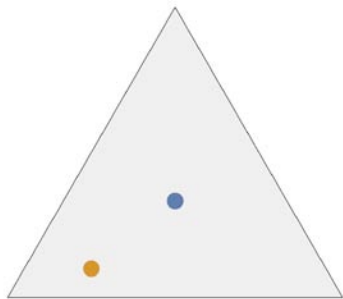


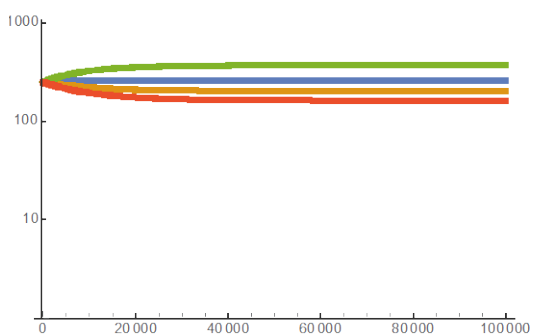
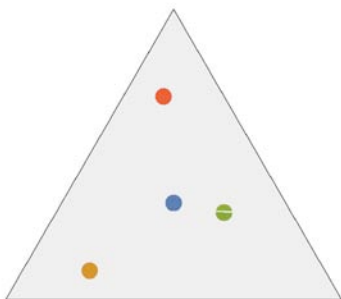
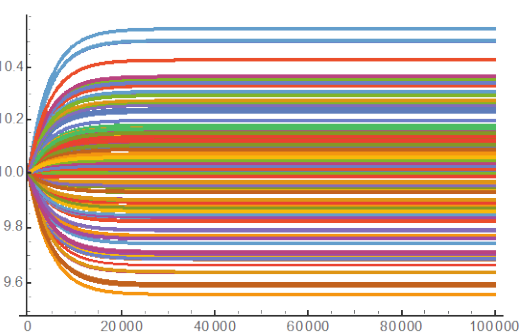
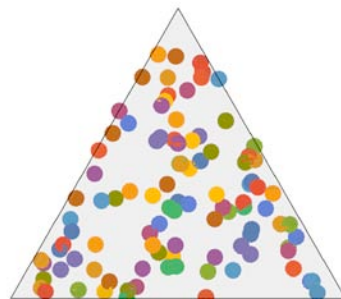
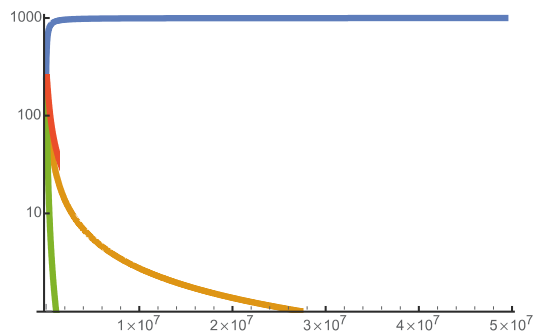
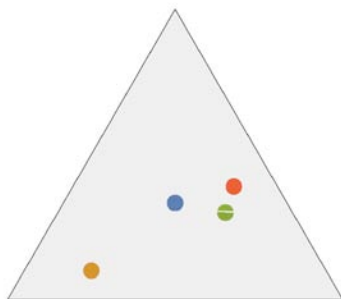
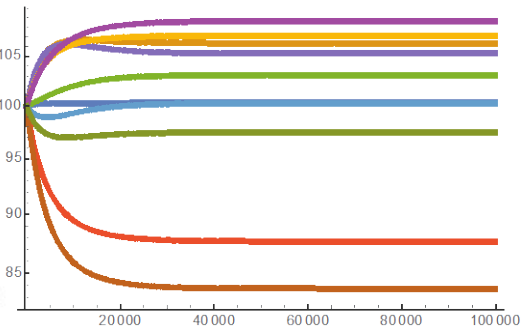
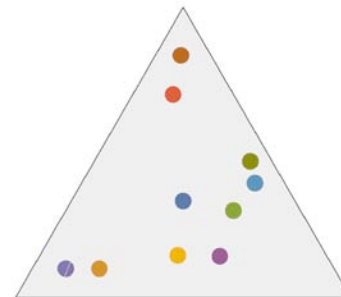
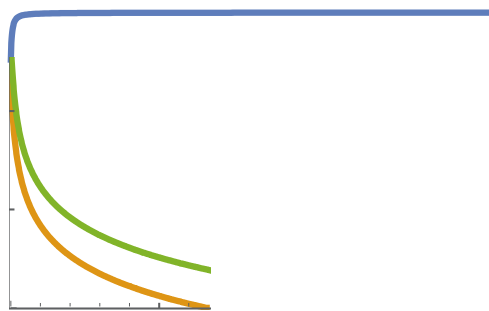
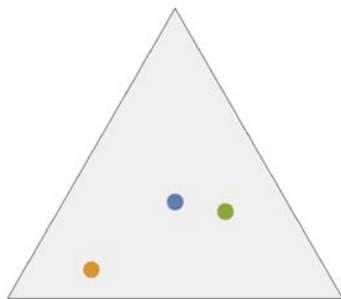
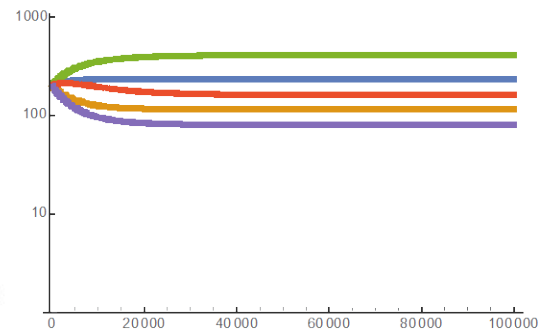
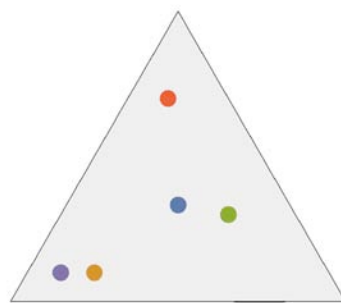
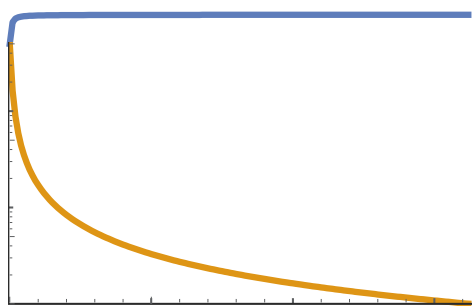
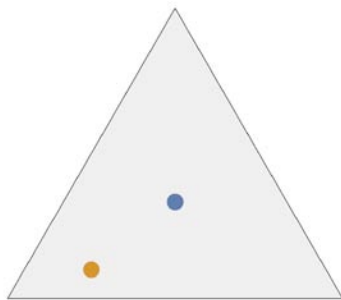




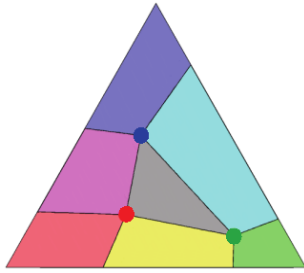




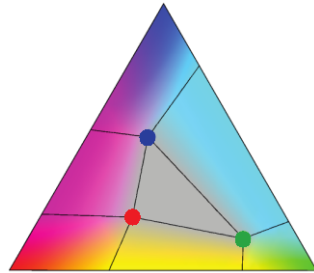




species = # resources:



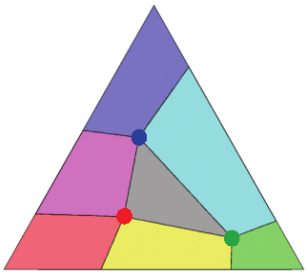
Surviving species



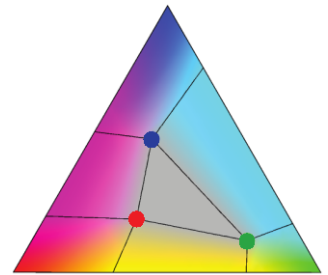
Nutrient concentrations

$$g_{\sigma}(\vec{c}) = \sum_i \alpha_{\sigma i} \frac{c_i}{1 + c_i}$$

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Surviving species

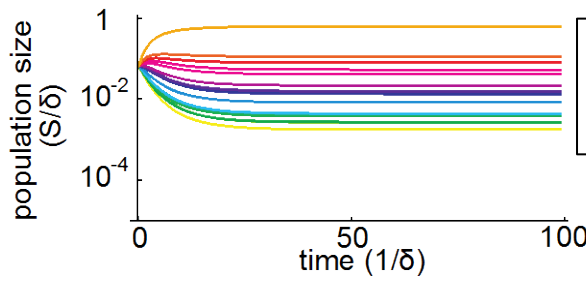
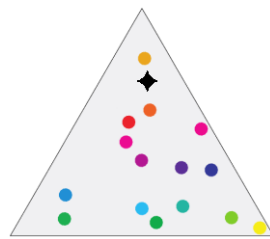
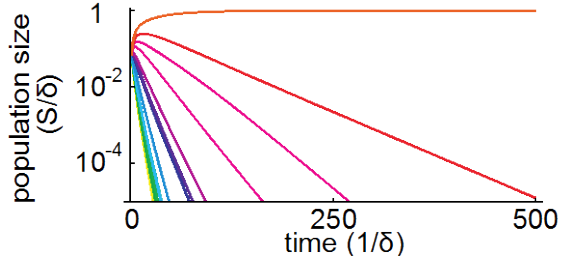
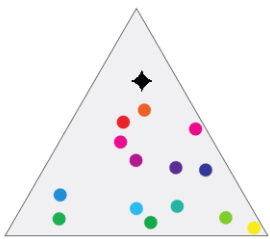


Nutrient concentrations

$$g_{\sigma}(\vec{c}) = \sum_i \alpha_{\sigma i} \frac{c_i}{1 + c_i}$$

species ≥ # resources:

A collection of $\{\vec{\alpha}_{\sigma}\}$ species coexist in steady state \Leftrightarrow the supply \vec{s} lies within the convex hull of the species $\{\vec{\alpha}_{\sigma}\}$.



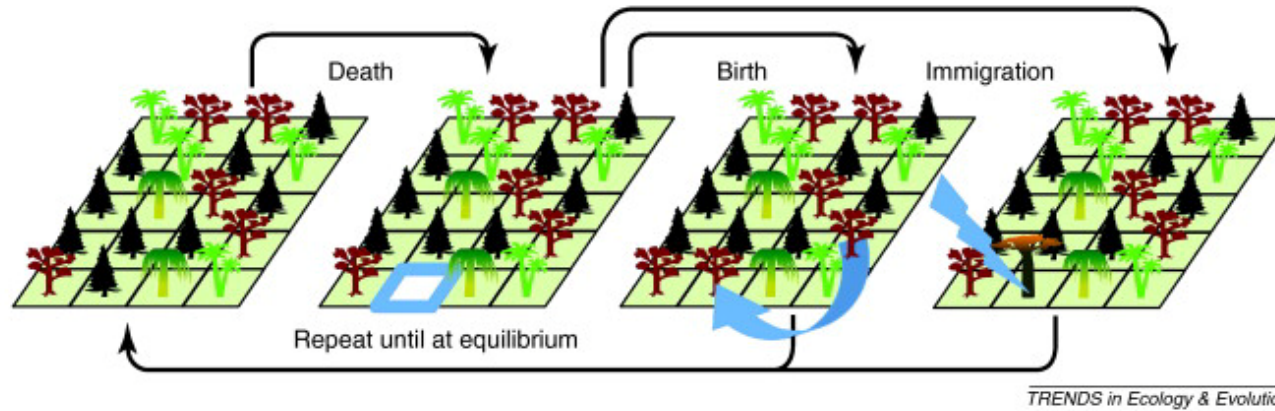
Gold species acts as "keystone species".

Competition \rightarrow nutrient concentrations become too low for certain species to survive \rightarrow at most #resources - 1 species survive.

Cooperation \rightarrow balanced nutrient concentrations \rightarrow all species are equally fit \rightarrow all species survive!

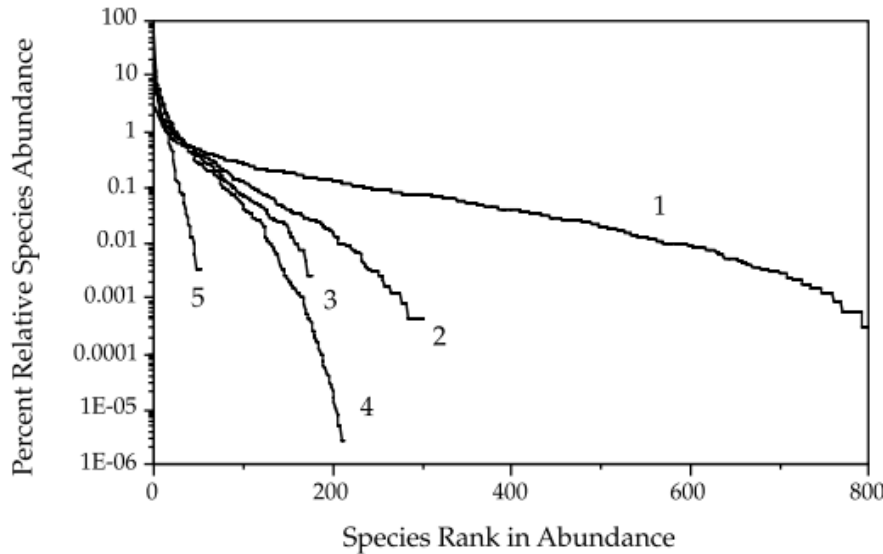
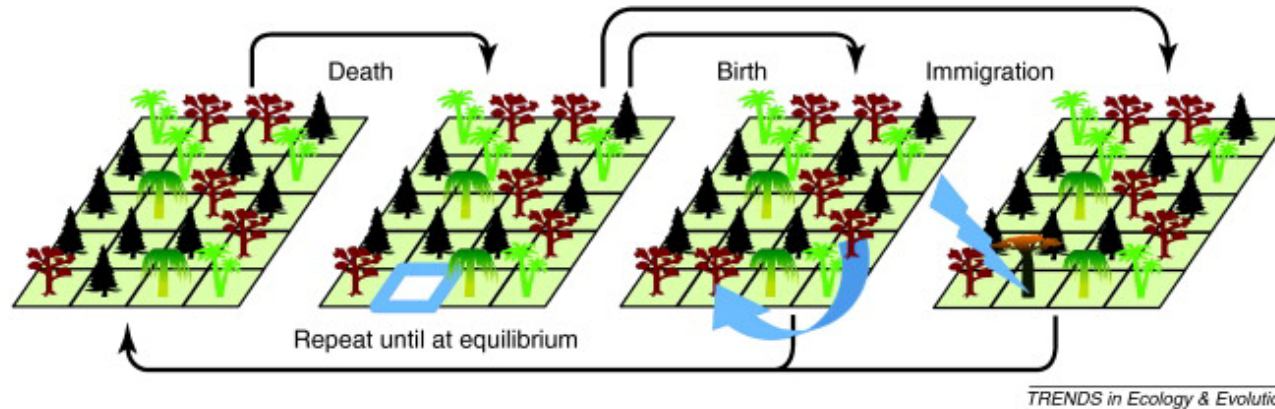
The neutral theory of biodiversity

Neutral theory: species are ecologically equivalent, and diversity emerges from ecological drift.



The neutral theory of biodiversity

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1. Tropical wet forest in Amazonia
2. Tropical dry deciduous forest in Costa Rica
3. Marine planktonic copepod community from the North Pacific gyre
4. Terrestrial breeding birds of Britain
5. Tropical bat community from Panama

Discrete, stochastic version of the model

Birth-death-immigration process $\vec{N}(T)$:

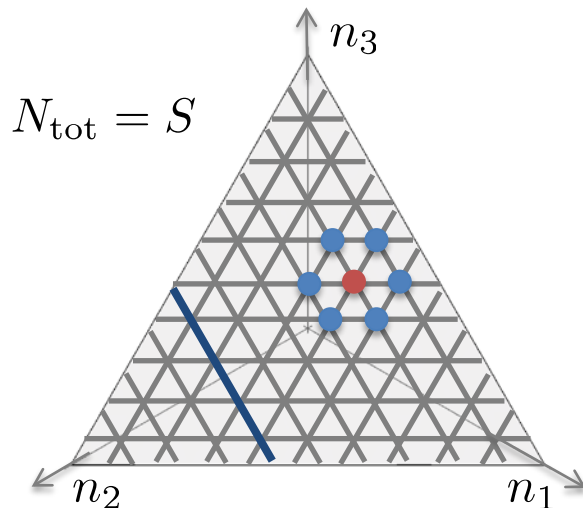
1. At $T = 0$, $\vec{N}(T)$ starts from a fixed initial population $\vec{N}(0)$.
2. At each time step

(i) a random individual dies:

$$P(\text{individual from species } \sigma \text{ dies}) = \frac{N_\sigma}{N_{\text{tot}}}$$

(ii) it is replaced by a random new individual:

$$P(\text{new individual is an immigrant}) = \nu$$
$$P(\text{new individual is from species } \sigma) = (1 - \nu) \frac{g_\sigma(\vec{N}, \vec{S}) N_\sigma}{N_{\text{tot}}}$$



In deterministic model:

$$\dot{n}_\sigma = (g_\sigma(\vec{n}, \vec{s}) - \delta) n_\sigma, \quad n_{\text{tot}} = s/\delta$$

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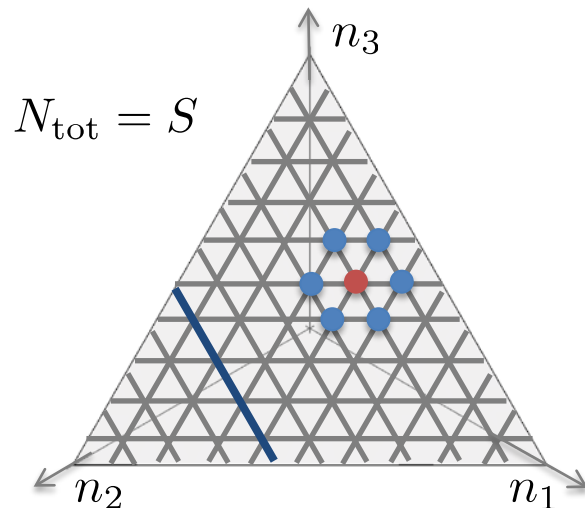
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Connection between deterministic and stochastic models:

If there are many organisms per each non-extinct species, i.e. S and N_σ are large, and $T = O(S)$ then

$$N_\sigma(T) \approx \frac{\delta S}{s} n_\sigma \left(\frac{s}{\delta^2 S} T \right).$$

Connection to the neutral theory of biodiversity

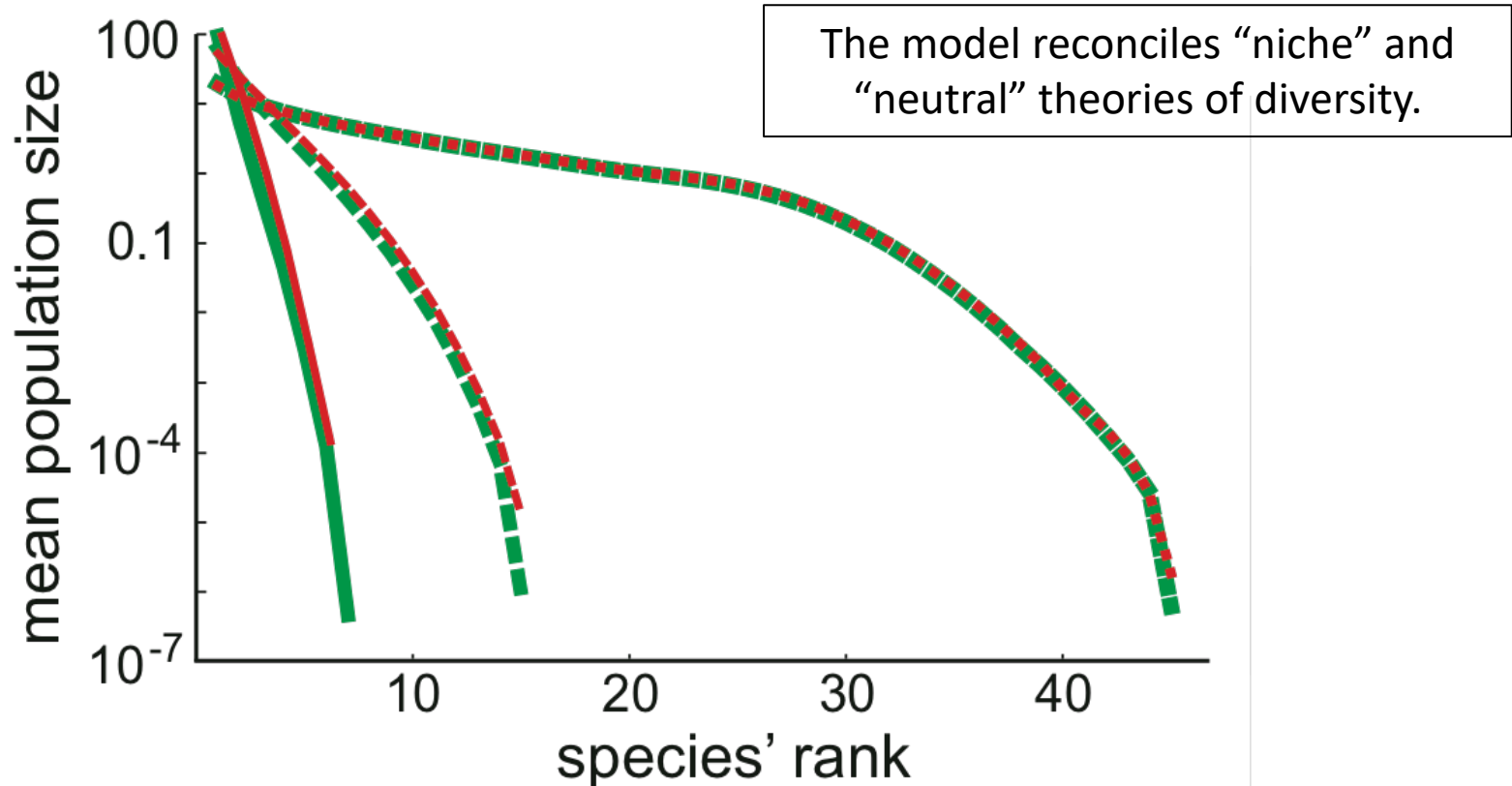
Rank-abundance curves

red – resource-competition model

green – neutral model

total population: $N_{\text{tot}} = 100$

immigration probabilities: $\nu = 0.001, 0.01, 0.1$



Robustness of coexistence

- I. Against population disturbances
- II. Against fluctuations in nutrient availability
- III. Against variability in species' budgets and death rates

Robustness of coexistence

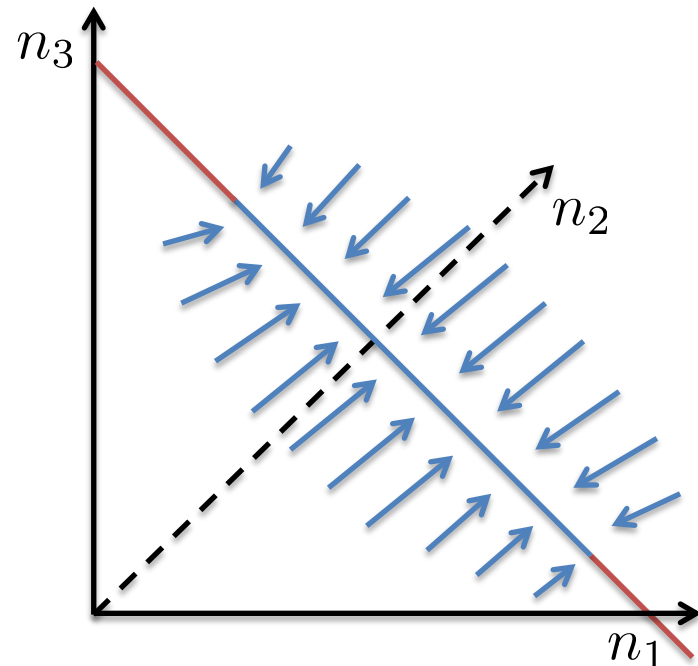
- I. Against population disturbances
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- III. Against variability in species' budgets and death rates

I. Against population disturbances:

The population fixed points are the solutions to the system of linear equations with #resources equations and #species variables:

$$n_1 \vec{\alpha}_1 + \dots + n_m \vec{\alpha}_m = \frac{E}{\delta} \vec{s}$$

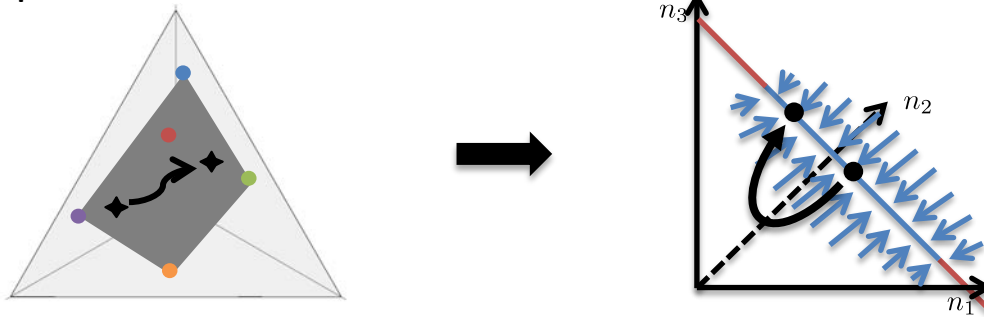
The population fixed points form the non-negative subset of a #species-#resources dimensional plane, and this set is an attractor of the dynamical system.



Robustness of coexistence

II. Against fluctuations in nutrient availability:

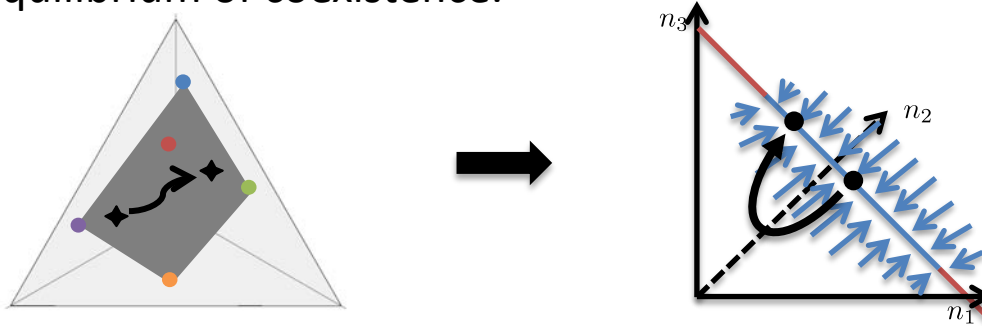
If nutrient supply changes, but remains in the convex hull of the species, the populations find a new equilibrium of coexistence.



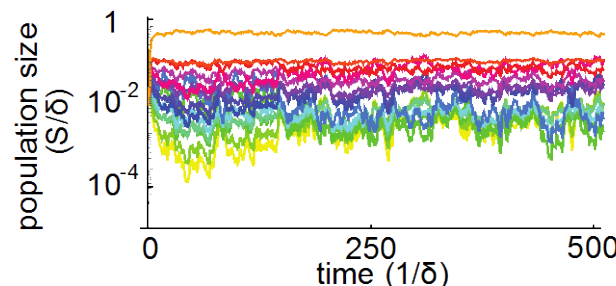
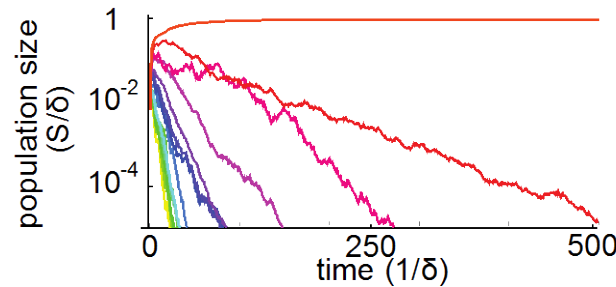
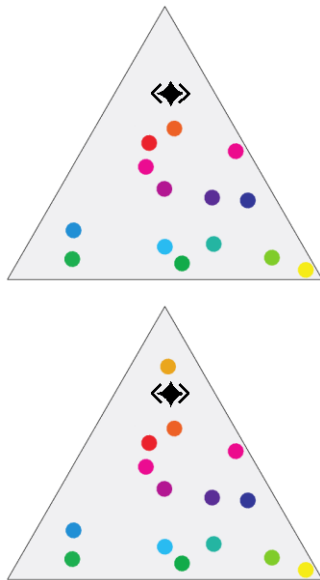
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Time-dependent nutrient supply: The supply regularly changes, at a fixed time interval T , to a new randomly selected supply, while the total supply is fixed:



Mean supply hypothesis:

A collection of species $\{\vec{\alpha}_\sigma\}$ coexist.



The mean supply $\langle \vec{s} \rangle$ lies within the convex hull of the species $\{\vec{\alpha}_\sigma\}$.

Robustness of coexistence

III. Against variability in species' budgets and death rates:

Definition of species σ :

enzyme distribution: $(\max(\alpha_{\sigma 1} + \xi_{\sigma 1}), \dots, \max(\alpha_{\sigma p} + \xi_{\sigma p}))$ with $\sum_{i=1}^p \alpha_{\sigma i} = E$

death rate: $\delta_{\sigma} = \delta + \xi_{\sigma}$,

where $\xi_{\sigma i}$ and ξ_{σ} are iid random variables with Gaussian distribution $\mathcal{N}(0, \Sigma^2)$

Robustness of coexistence

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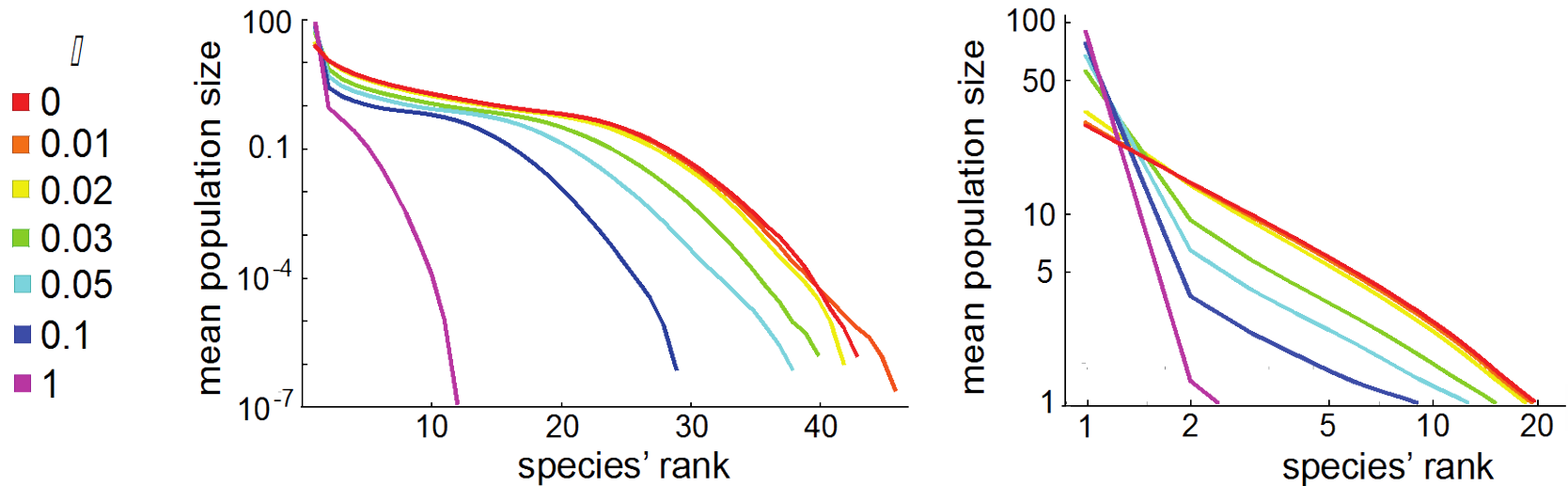
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In the deterministic version of the model, diversity is lost. However, in the stochastic version, diversity can be maintained:



$\Sigma \leq 0.02$ \Rightarrow High diversity

$\Sigma \geq 0.03$ \Rightarrow Low diversity

Robustness of coexistence

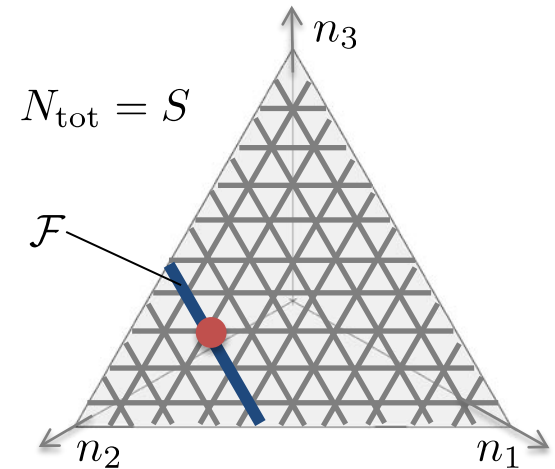
There is a critical standard deviation Σ^* that separates the high and low diversity regimes:

Estimate a species' lifetime T_0 when $\Sigma = 0$:

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$\vec{N}(T)$ converges to \mathcal{F} , and on \mathcal{F} the process is governed by neutral birth and death probabilities.

$$T_0 \propto S^2$$



Robustness of coexistence

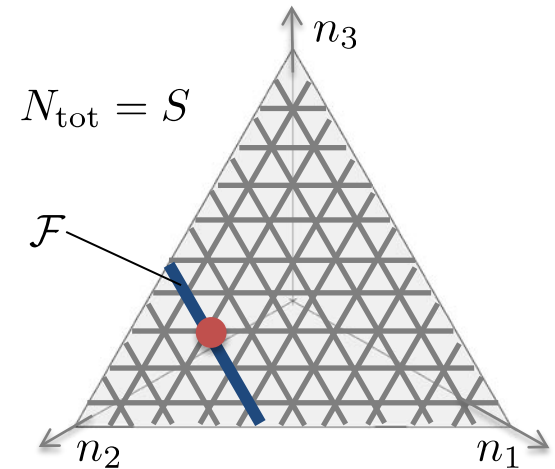
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$$\frac{dn_\sigma}{dt} = \left(\left(\sum_{i=1}^p (\alpha_{\sigma i} + \xi_{\sigma i}) c_i \right) - \delta \right) n_\sigma \propto -\Sigma n_\sigma$$

$$T_\Sigma \propto \frac{S \log(S)}{\Sigma}$$

Robustness of coexistence

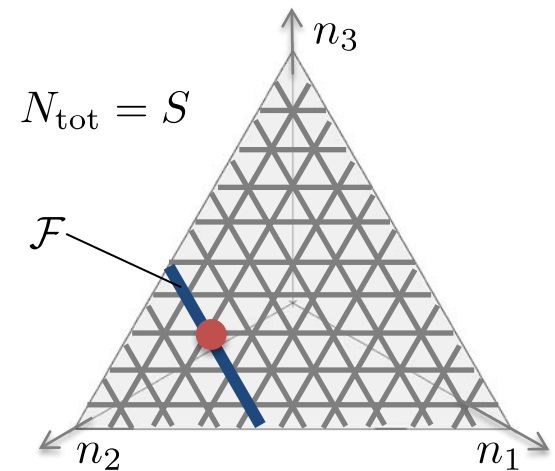
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$$T_0 = T_\Sigma \Rightarrow \Sigma^* \propto \log(S)/S$$

Conclusions

- Features of model that allow for coexistence:
 - Organisms take part in shaping their environment.
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Thank you!

Proof of convex hull condition

What are the steady-state concentrations of nutrients required for coexistence?

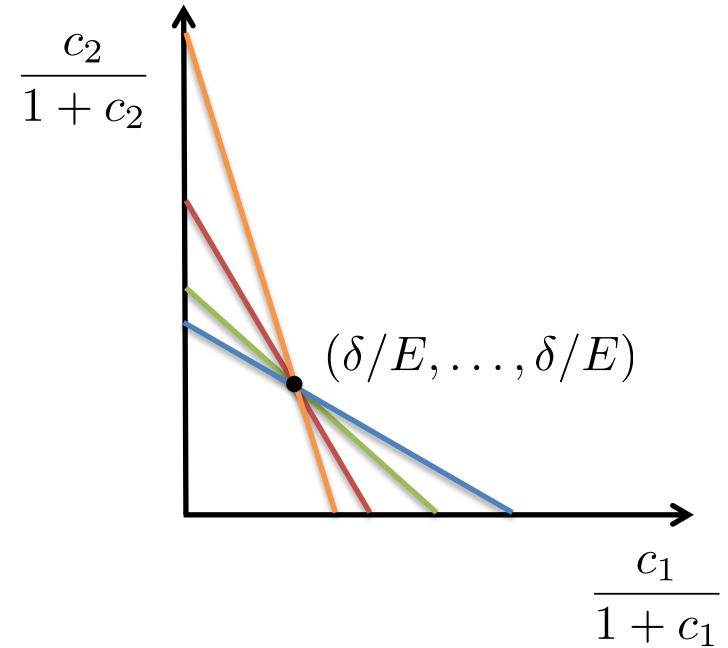
growth rate = death rate

$$\sum_{i=1}^p \alpha_{\sigma i} \frac{c_i}{1 + c_i} = \delta$$

⇓ at least #resources equations

$$\frac{c_1^*}{1 + c_1^*} = \dots = \frac{c_p^*}{1 + c_p^*} = \frac{\delta}{E}$$

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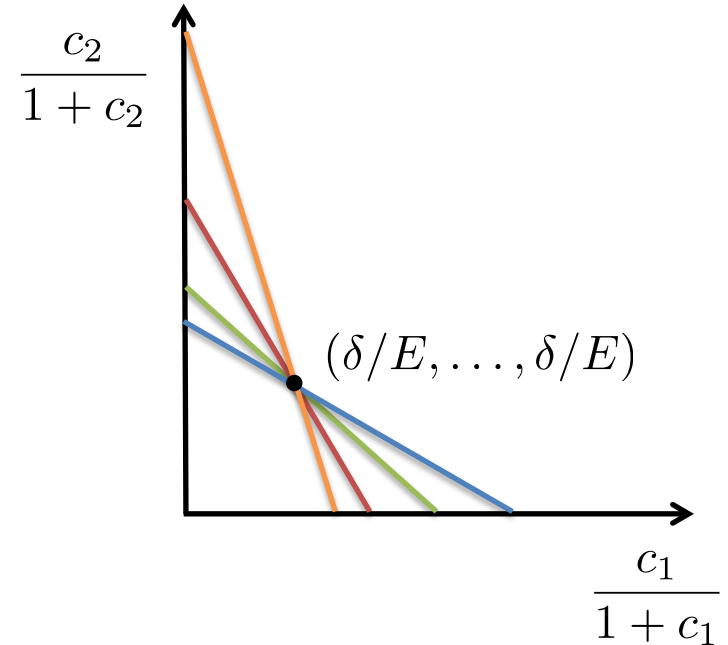
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What strategies can adjust their populations to achieve these steady-state nutrient concentrations?

$$\frac{\delta}{E} = \frac{c_i^*}{1 + c_i^*} = \frac{s_i}{\sum_{\sigma} n_{\sigma} \alpha_{\sigma i}}$$

$$n_1 \vec{\alpha}_1 + \dots + n_m \vec{\alpha}_m = \frac{E}{\delta} \vec{s}$$

The solutions \vec{n} form a #species – #resources dimensional plane in the space of population sizes.

\exists solution $\vec{n} > 0$ if the supply \vec{s} is in the convex hull of the strategies $\vec{\alpha}_{\sigma}$.