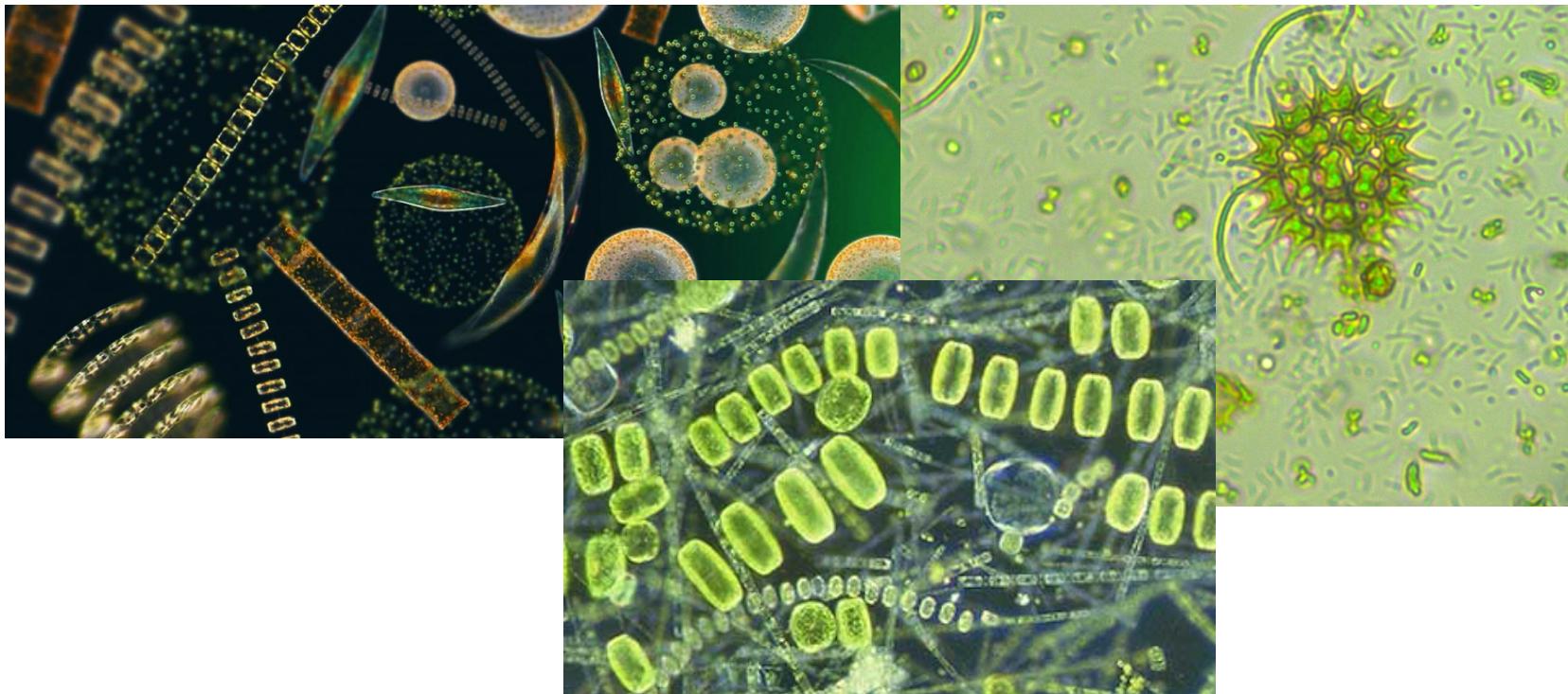
A microscopic image showing a dense population of phytoplankton. The cells are mostly small, oval-shaped, and green, distributed throughout the frame. There are also larger, more complex cells, including some with internal structures like chloroplasts and a few larger, elongated cells, possibly diatoms or dinoflagellates. The overall color palette is dominated by various shades of green and yellow.

Predicting phytoplankton metabolism from the individual size distribution

Daniel Padfield

Phytoplankton are key for the carbon cycle

- ~50% of annual carbon fixation
- Fuel entire ocean food webs



Current measurements of metabolism

Current measurements of metabolism

Bottle incubations – *in vivo*



- Snapshot in time
- Single community
- Control over environmental variables

Current measurements of metabolism

Bottle incubations – *in vivo*



- Snapshot in time
- Single community
- Control over environmental variables

Gas measurements – *in situ*



- Integrates over large areas
- Many communities and many different environmental conditions

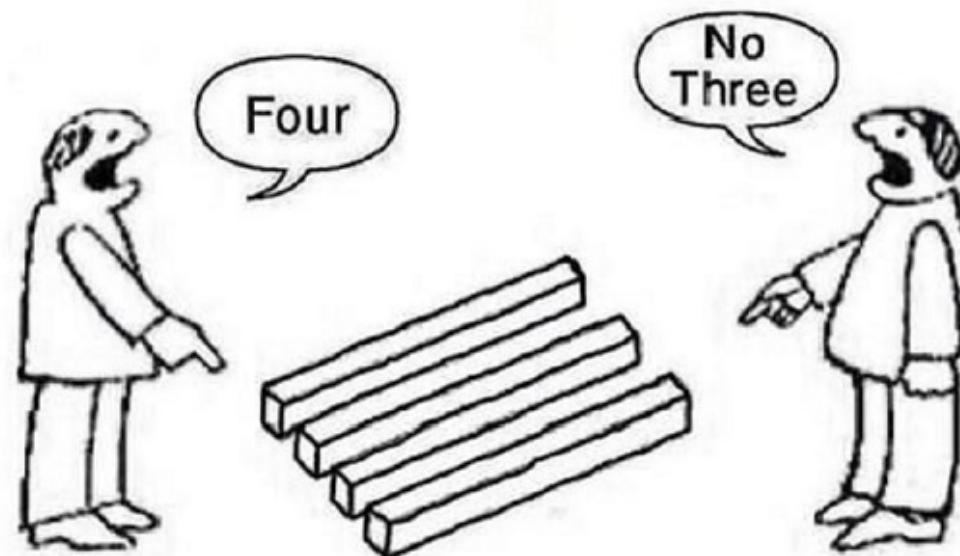
Current measurements of metabolism

Bottle incubations – *in vivo*

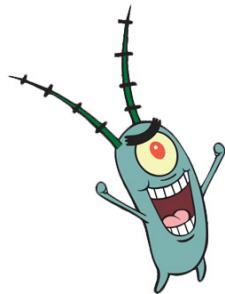
- Total photosynthesis < total respiration

Gas measurements – *in situ*

- Total photosynthesis > total respiration



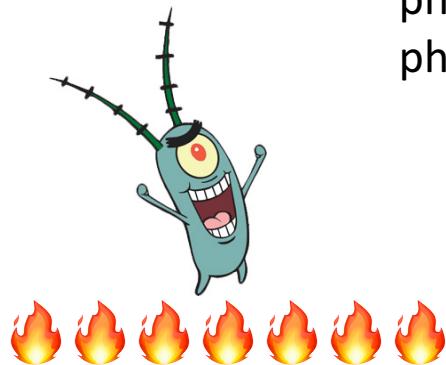
Crash course in metabolic theory



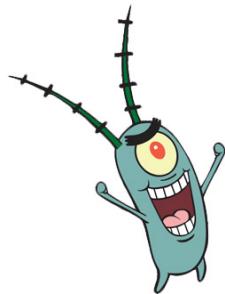
Crash course in metabolic theory

$$e^{E(\frac{1}{kT_c} - \frac{1}{kT})}$$

E previously found to be around 0.8 eV for photosynthesis and > 1 for respiration in phytoplankton



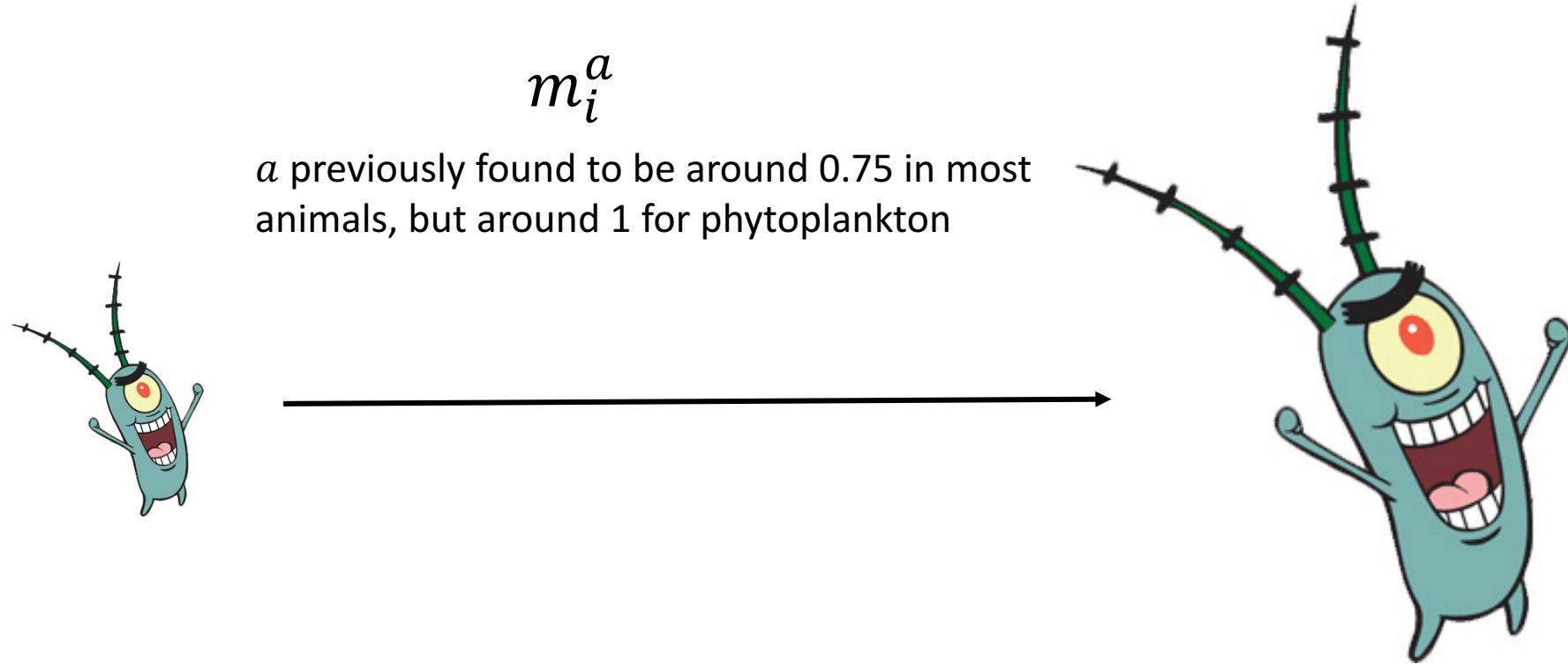
Crash course in metabolic theory



Crash course in metabolic theory

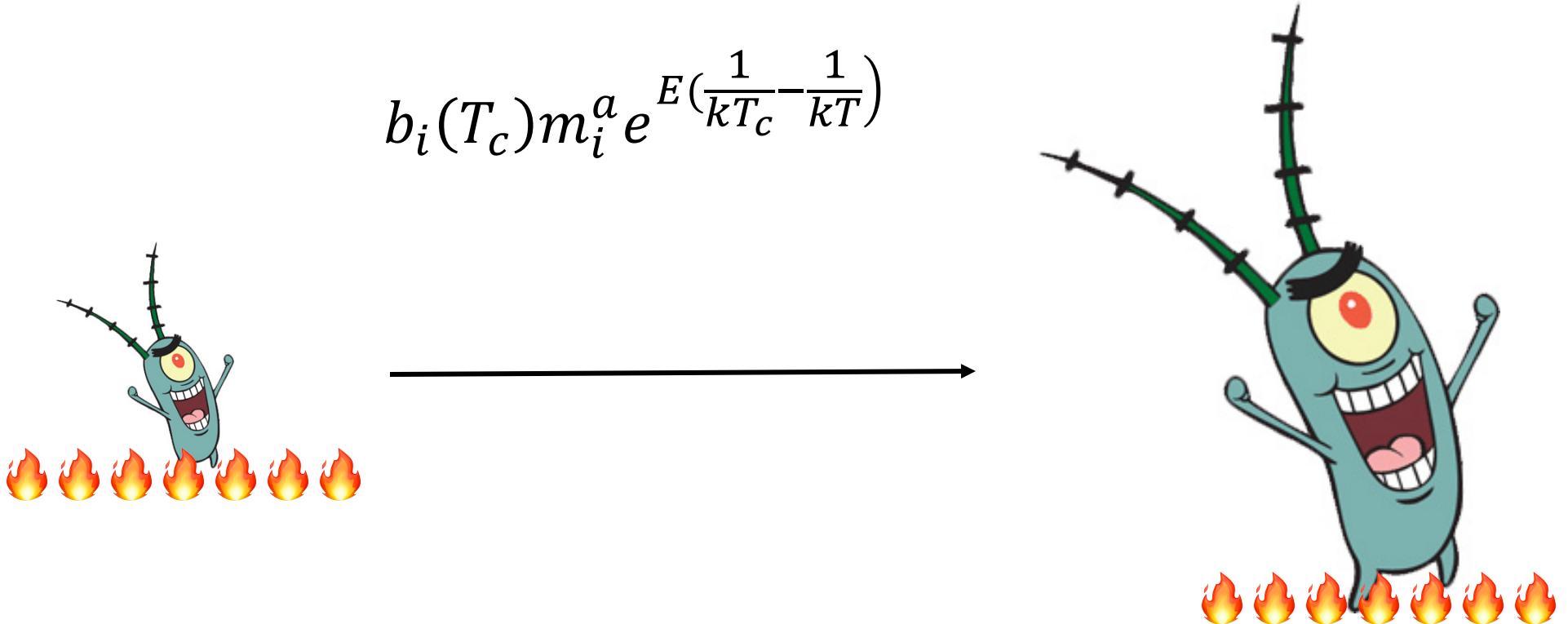
$$m_i^a$$

a previously found to be around 0.75 in most animals, but around 1 for phytoplankton



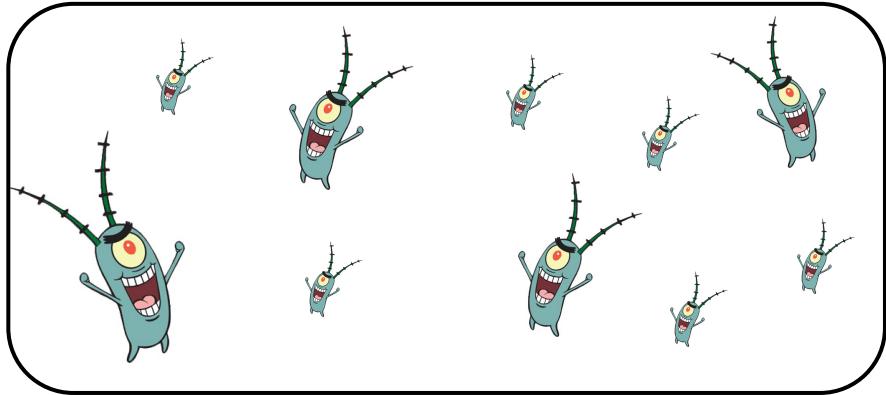
Crash course in metabolic theory

$$b_i(T_c)m_i^a e^{E\left(\frac{1}{kT_c} - \frac{1}{kT}\right)}$$



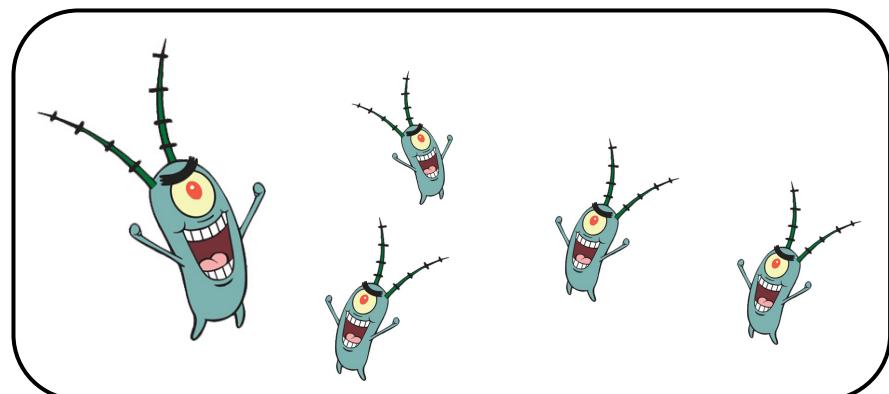
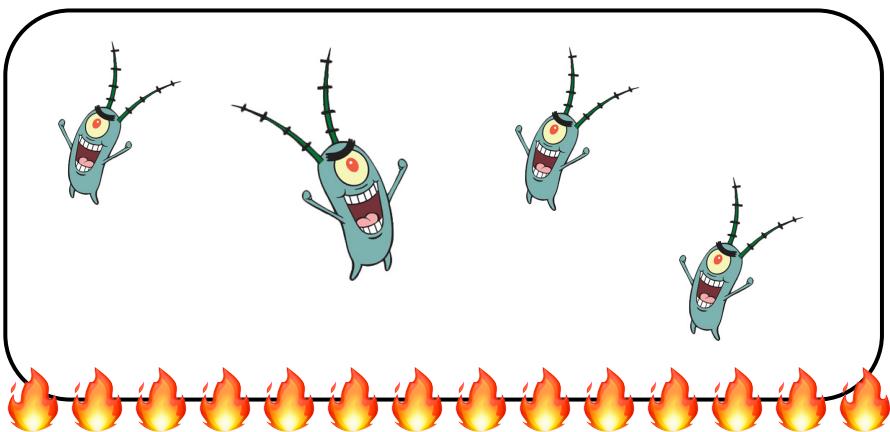
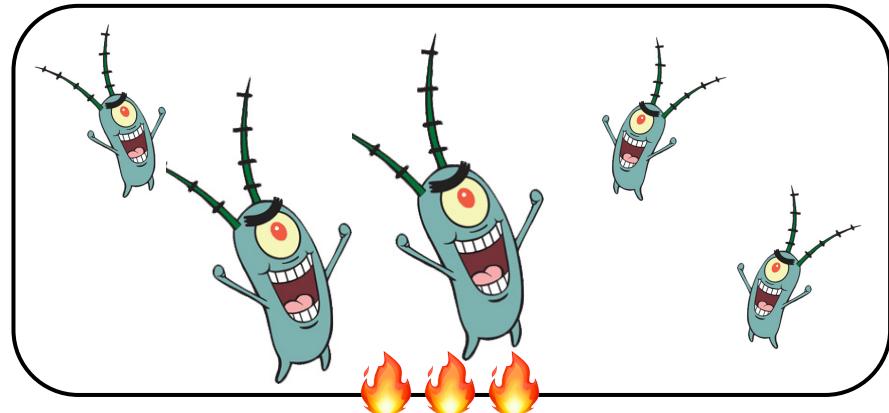
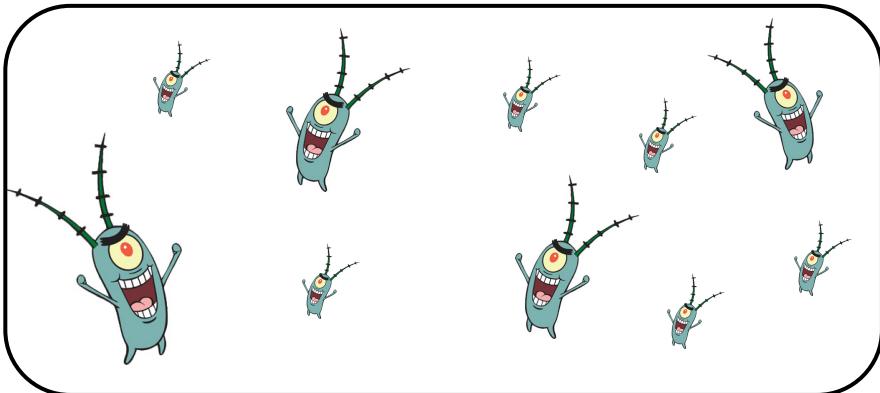
Crash course in metabolic theory

$$\sum_{i=1}^{n_{tot}} b_i(T_c) m_i^a e^{E(\frac{1}{kT_c} - \frac{1}{kT})}$$

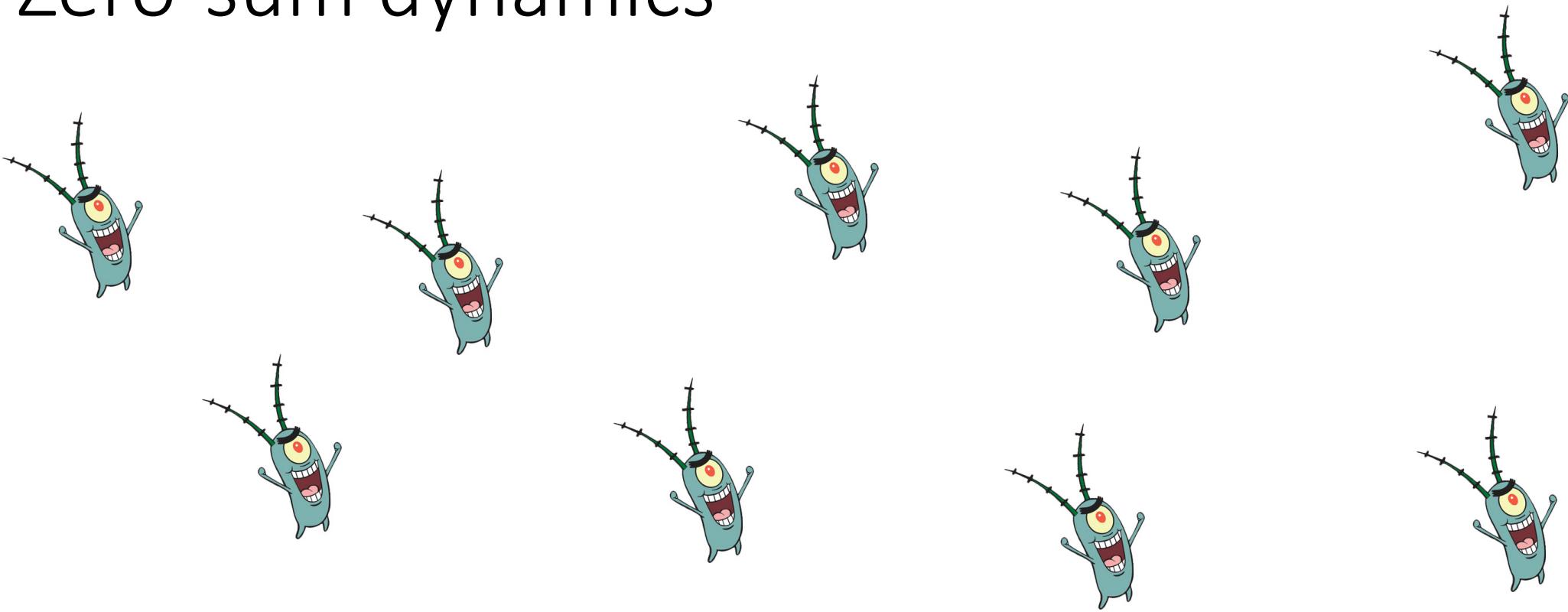


Crash course in metabolic theory

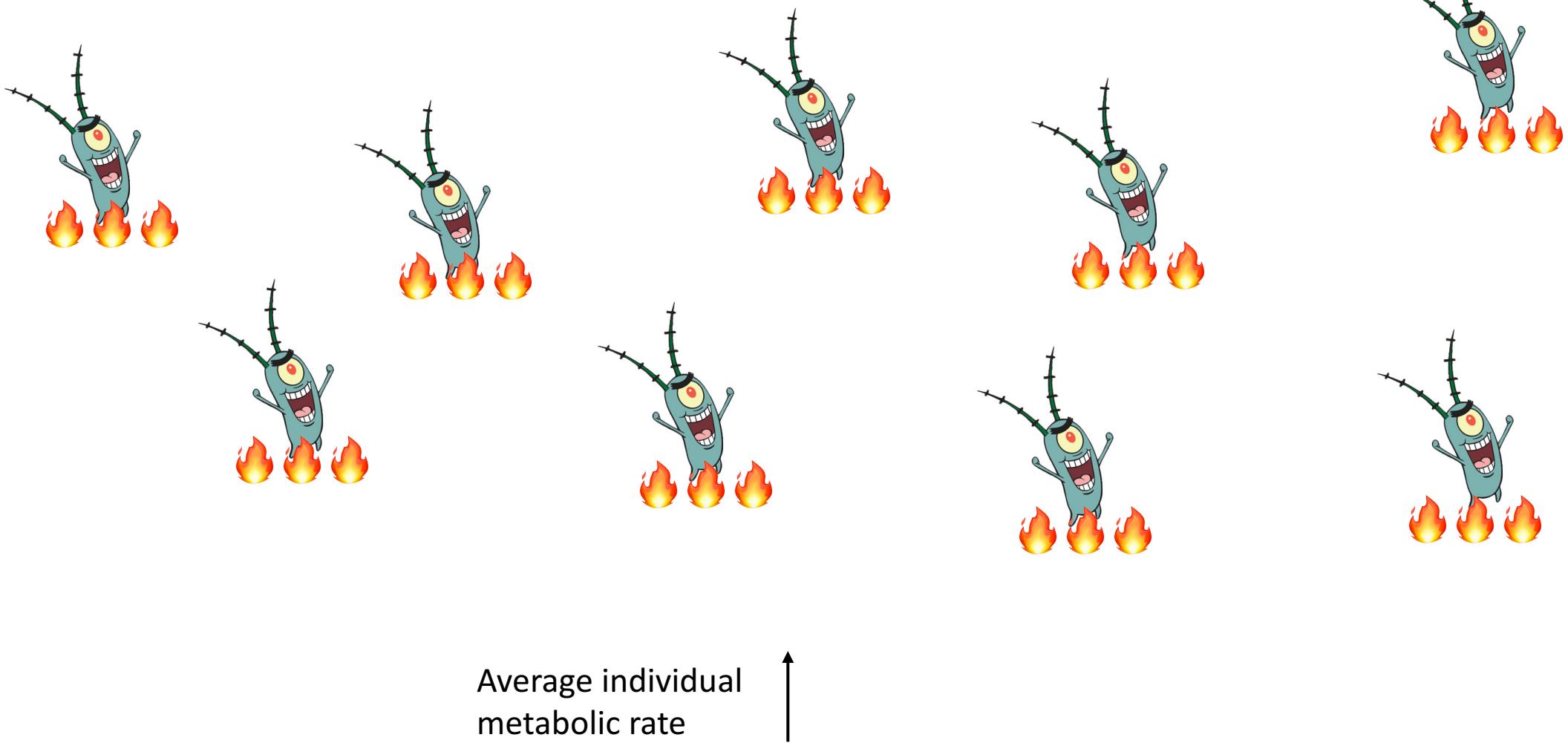
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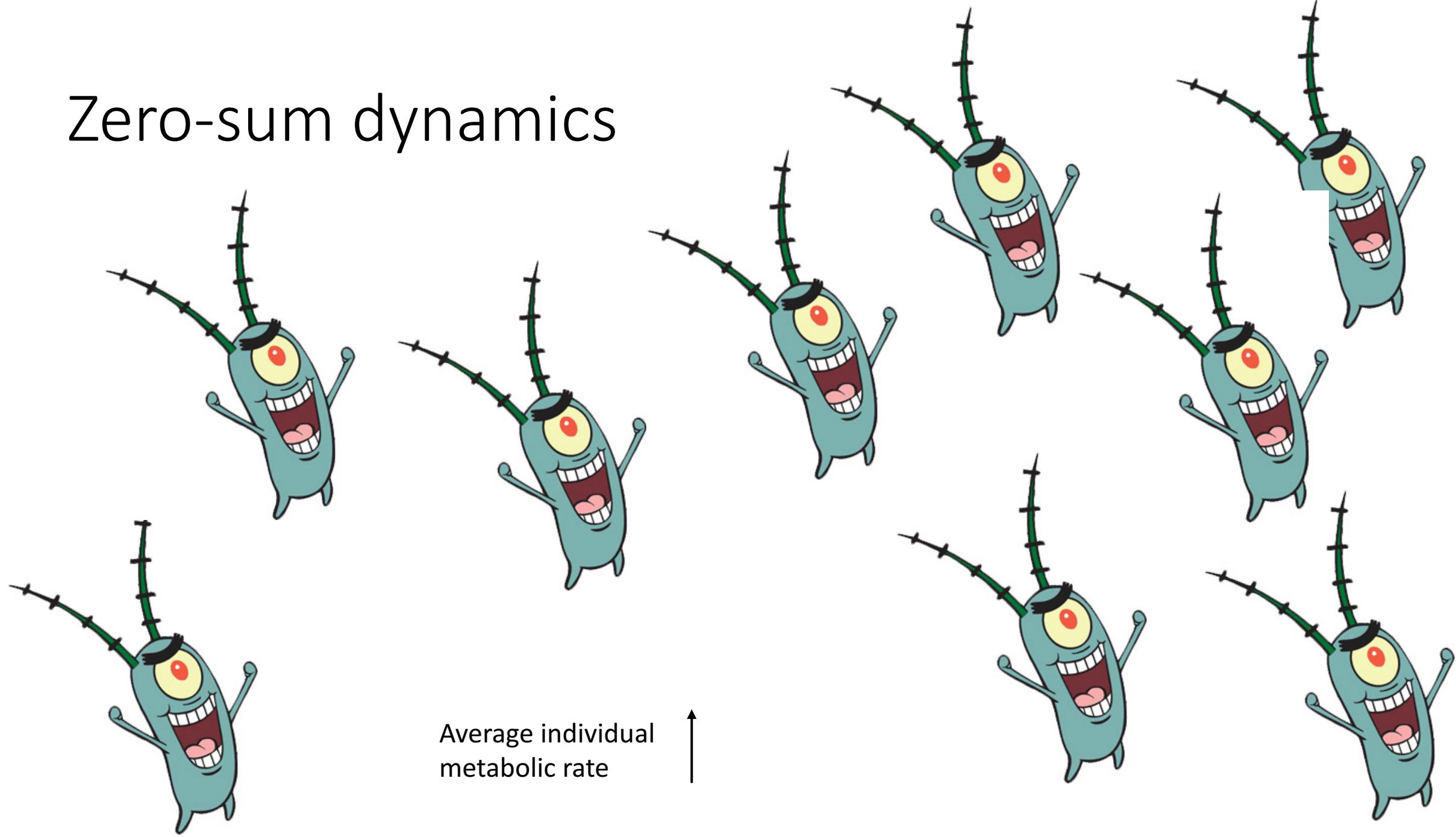
Zero-sum dynamics



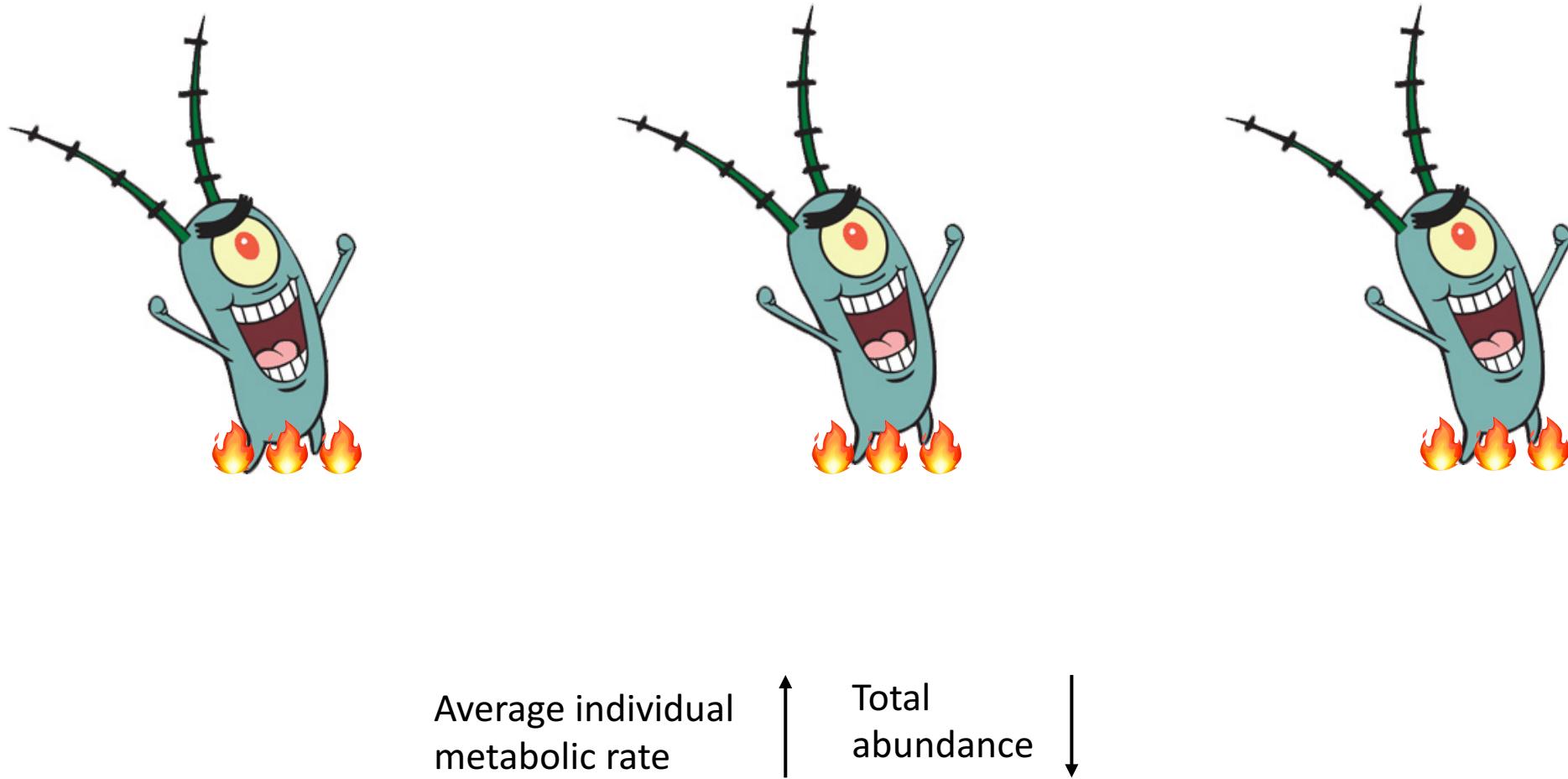
Zero-sum dynamics



Zero-sum dynamics



Zero-sum dynamics



Predictions

- Predict community metabolic rates from the size- and temperature-dependence of individuals in the size distribution
- Across-communities, there will be a trade off between total abundance and average individual metabolic rate
- Trade offs between community properties may impact community flux

Experimental setup

10 years warming : half 4°C above ambient



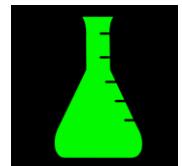
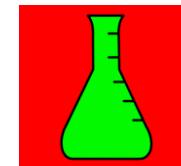
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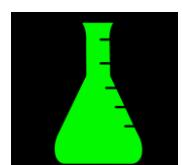
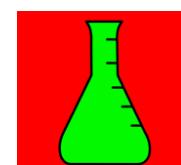


How do the communities and their functioning change in response to long-term and short-term warming

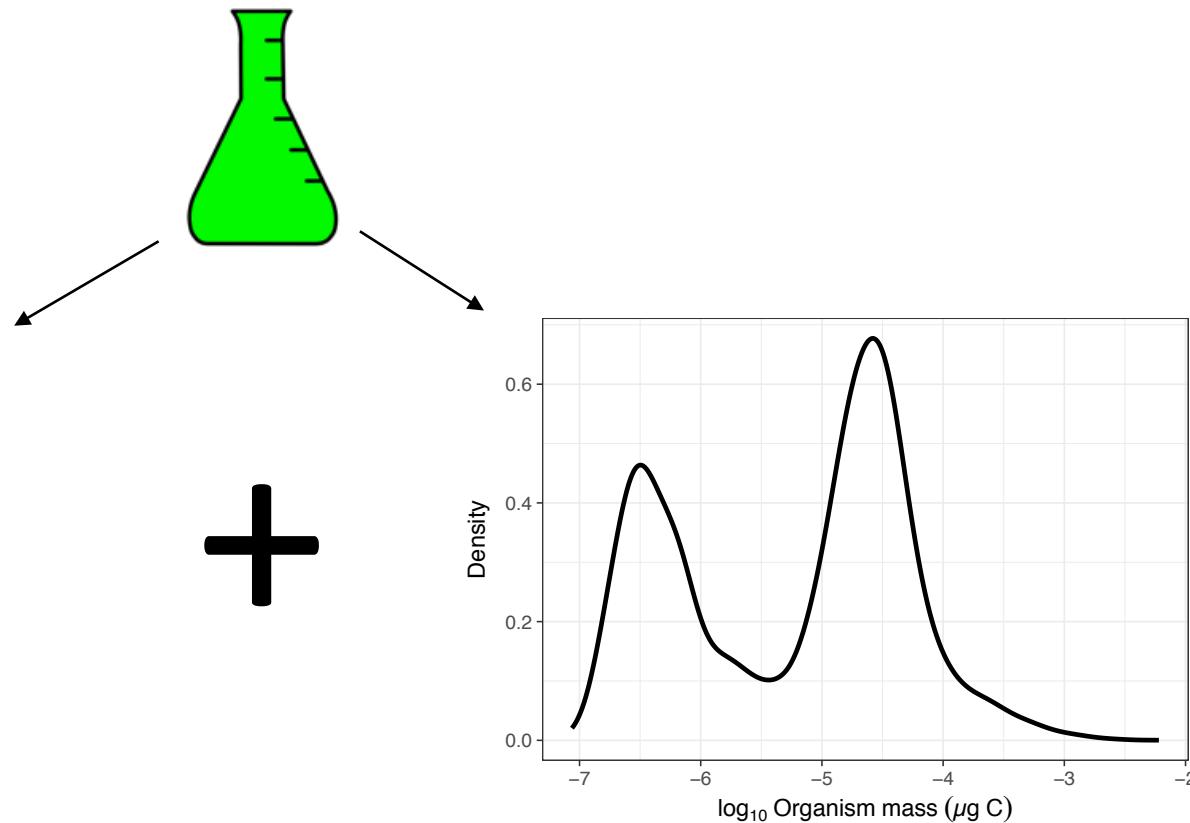
Cool incubator ~ 4 weeks



Warm incubator ~ 4 weeks



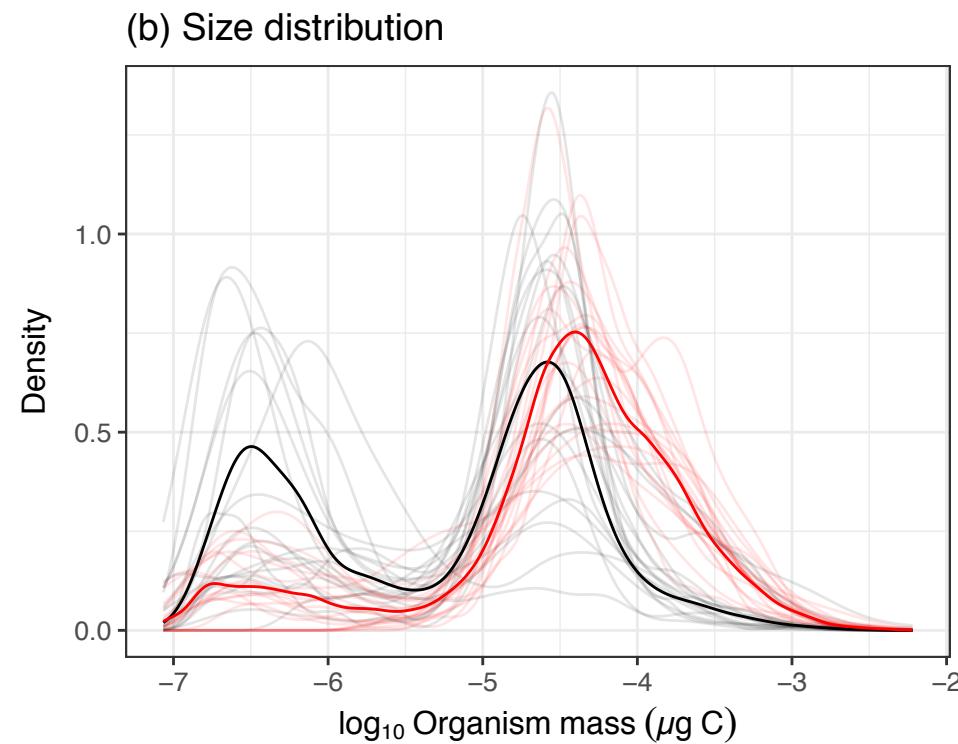
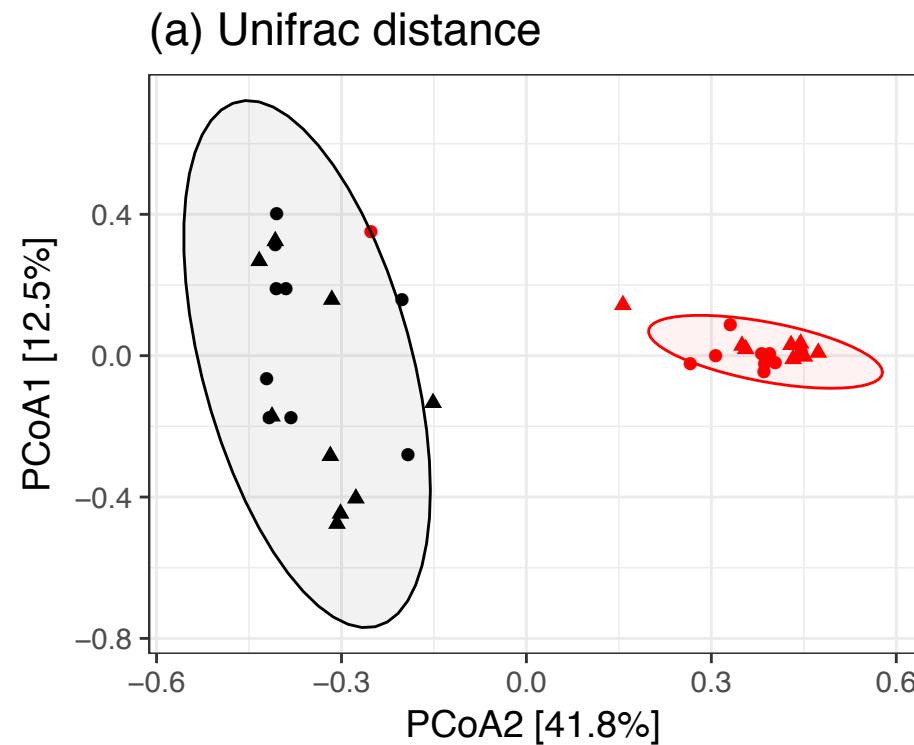
Measurements



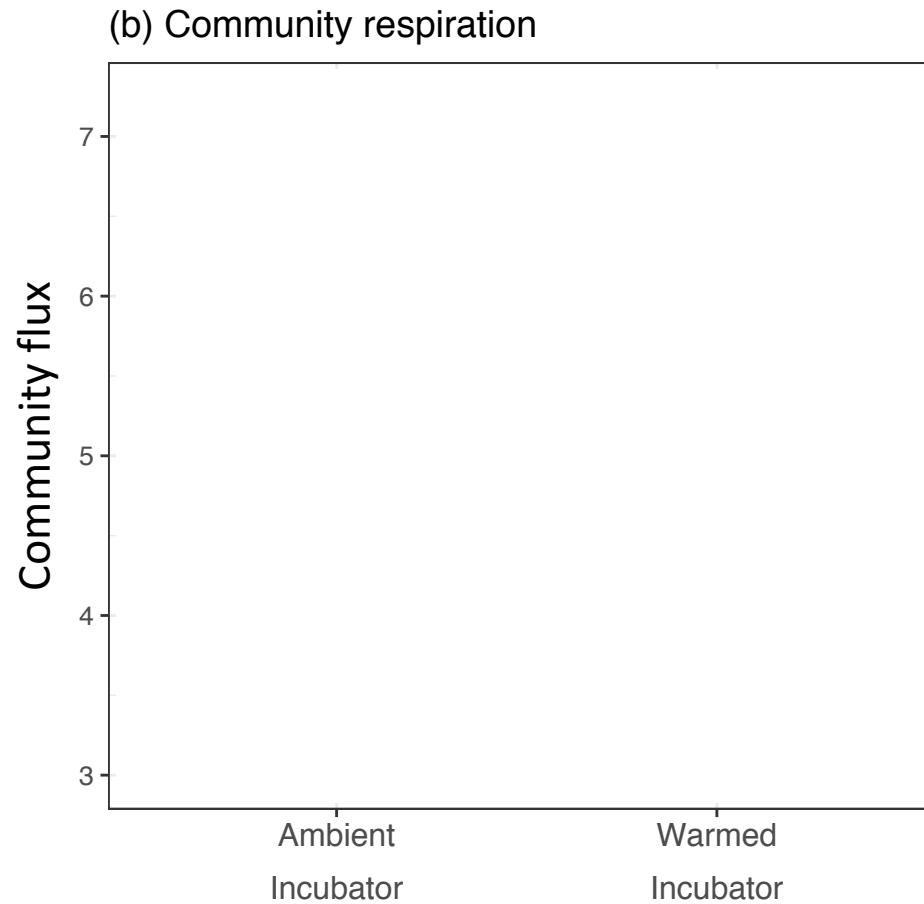
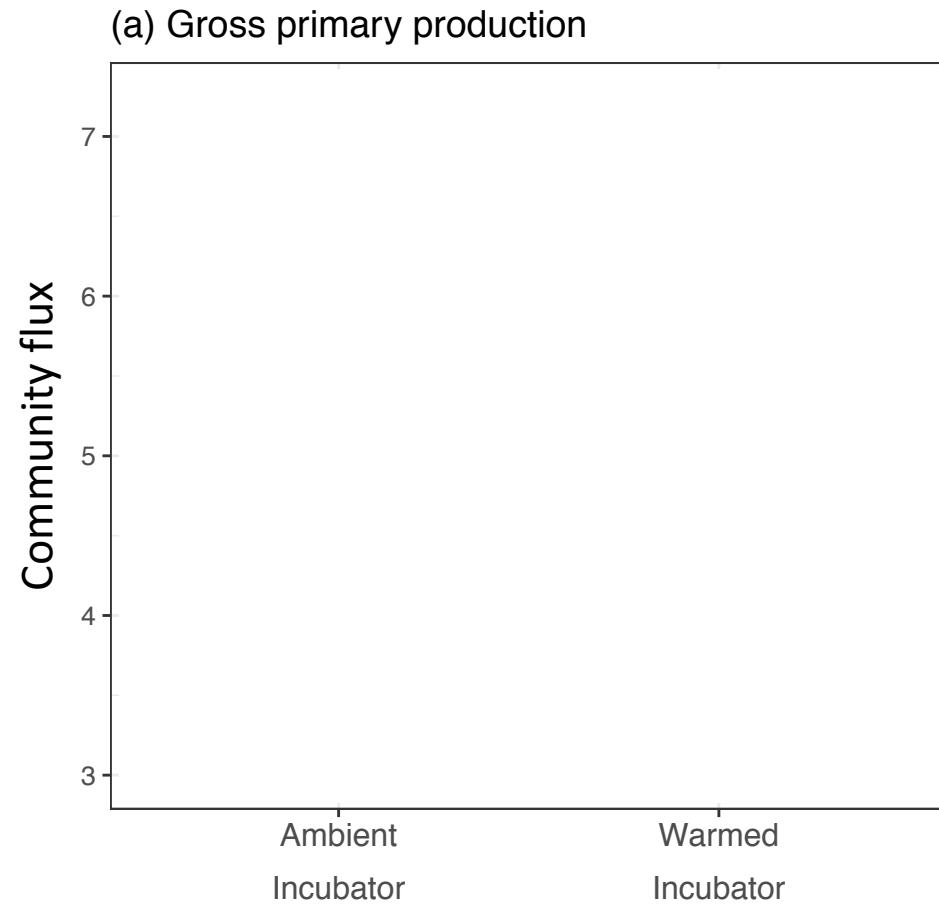
Metabolism @ incubator temperature
1 point for each microcosm

Size distribution
> 300 points per sample!

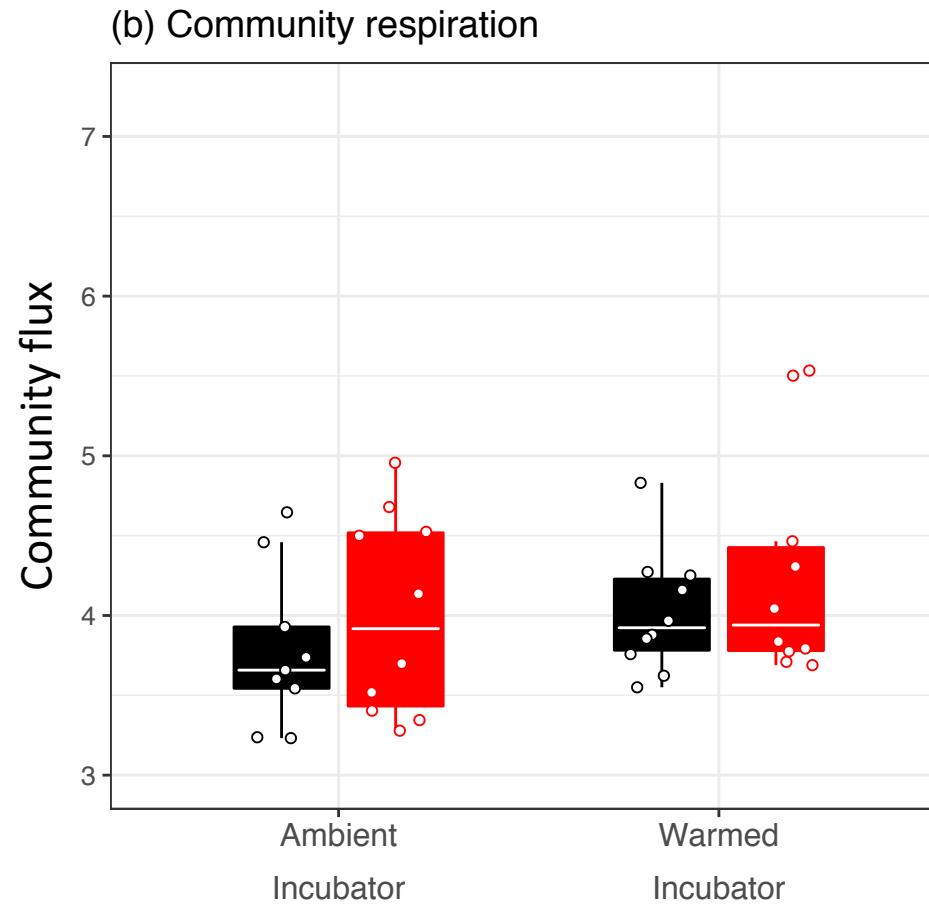
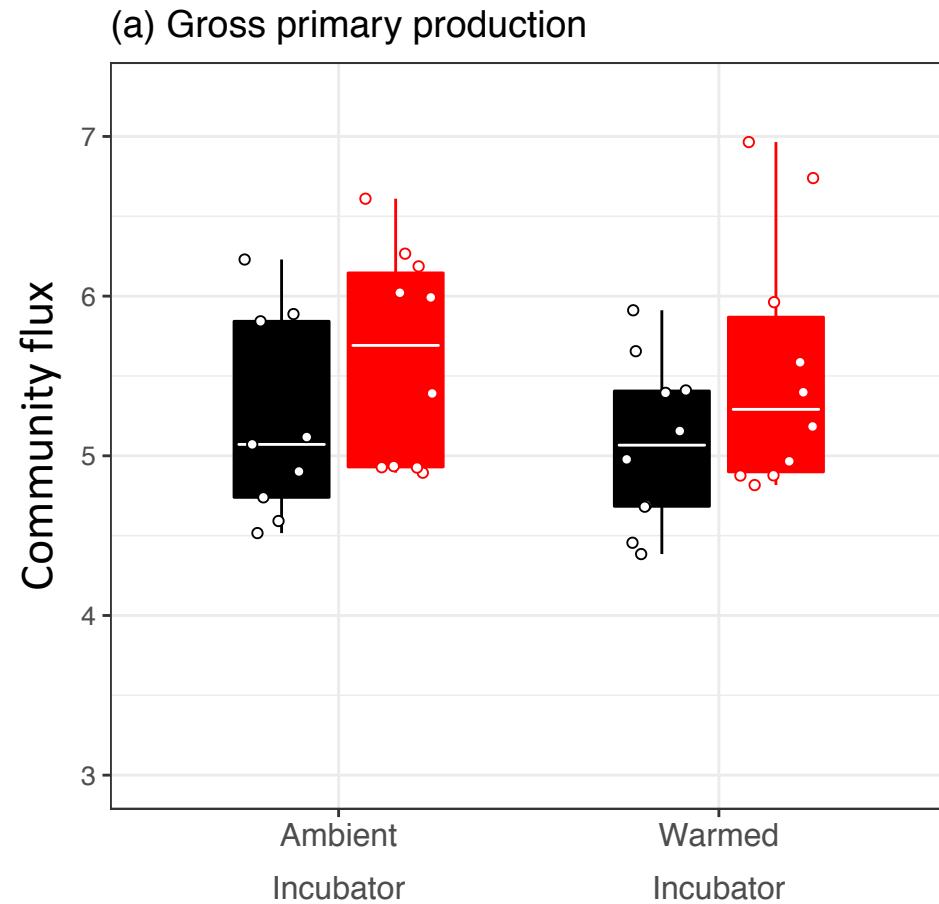
Long-term warming changes communities



No difference in measured raw metabolism



No difference in measured raw metabolism



Model

$$B_j(T) = \sum_{i=1}^{n_{tot}} b_i(T_c) m_i^a e^{E(\frac{1}{kT_c} - \frac{1}{kT})}$$

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$b_i(T_c)$ - the individual
normalisation constant

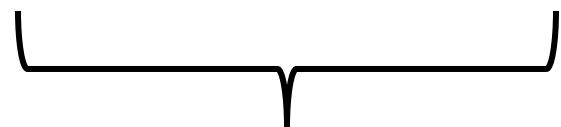
Model

a - how rates change
with body size

$$B_j(T) = \sum_{i=1}^{n_{tot}} b_i(T_c) m_i^a e^{E(\frac{1}{kT_c} - \frac{1}{kT})}$$

Model

$$B_j(T) = \sum_{i=1}^{n_{tot}} b_i(T_c) m_i^a e^{E(\frac{1}{kT_c} - \frac{1}{kT})}$$



E - how rates change
with temperature

Model predicts size and temperature-dependence

parameter	units	estimate	95% confidence interval
E_{GPP}	eV	0.741	0.196 - 1.286
E_{CR}	eV	1.417	0.853 - 1.982
α_{GPP}	-	0.887	0.567 - 1.174
α_{CR}	-	1.101	0.743 - 1.412
$\ln GPP(T_c)$	$\mu\text{mol O}_2 \text{ L}^{-1} \text{ hr}^{-1}$	-3.426	-6.335 - -0.989
$\ln CR(T_c)$ (ambient mesocosm)	$\mu\text{mol O}_2 \text{ L}^{-1} \text{ hr}^{-1}$	-2.717	-5.943 - -0.150
$\ln CR(T_c)$ (warm mesocosm)	$\mu\text{mol O}_2 \text{ L}^{-1} \text{ hr}^{-1}$	-3.110	-6.126 - -0.650

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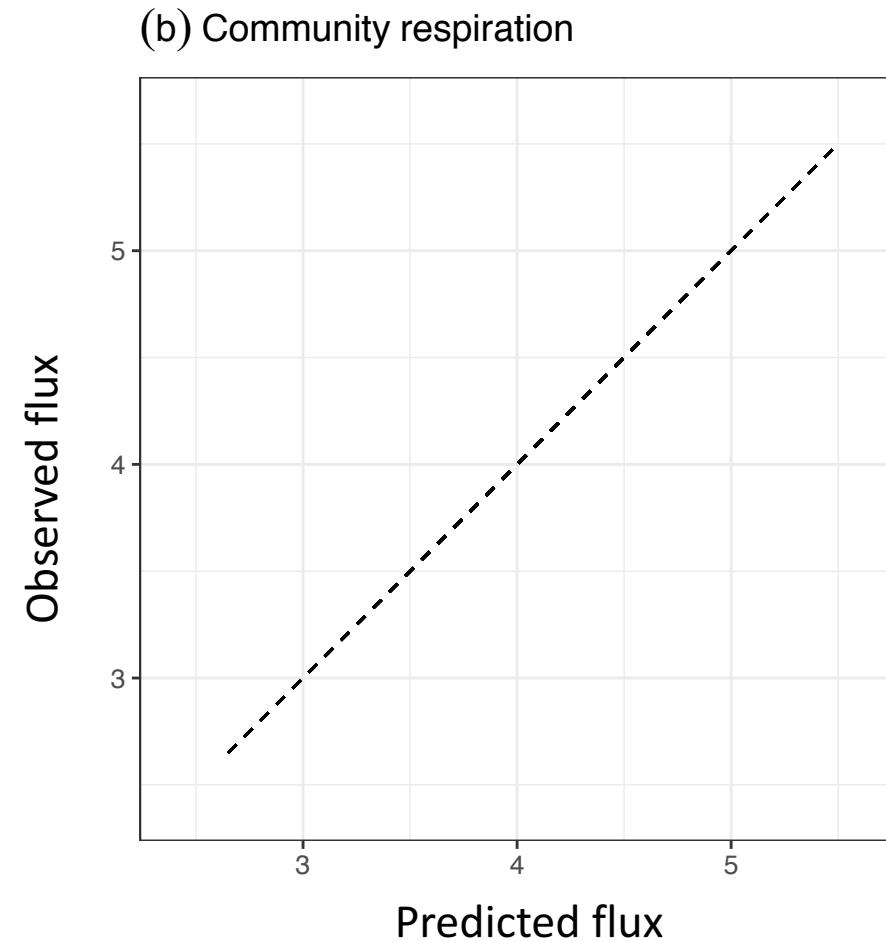
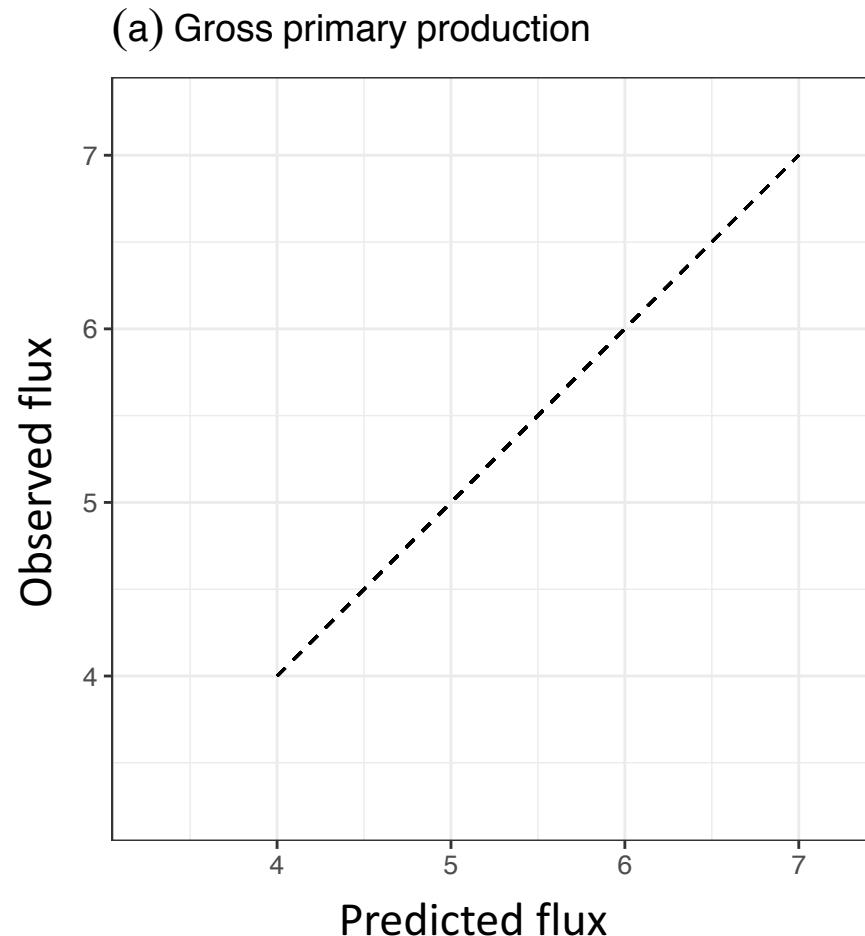
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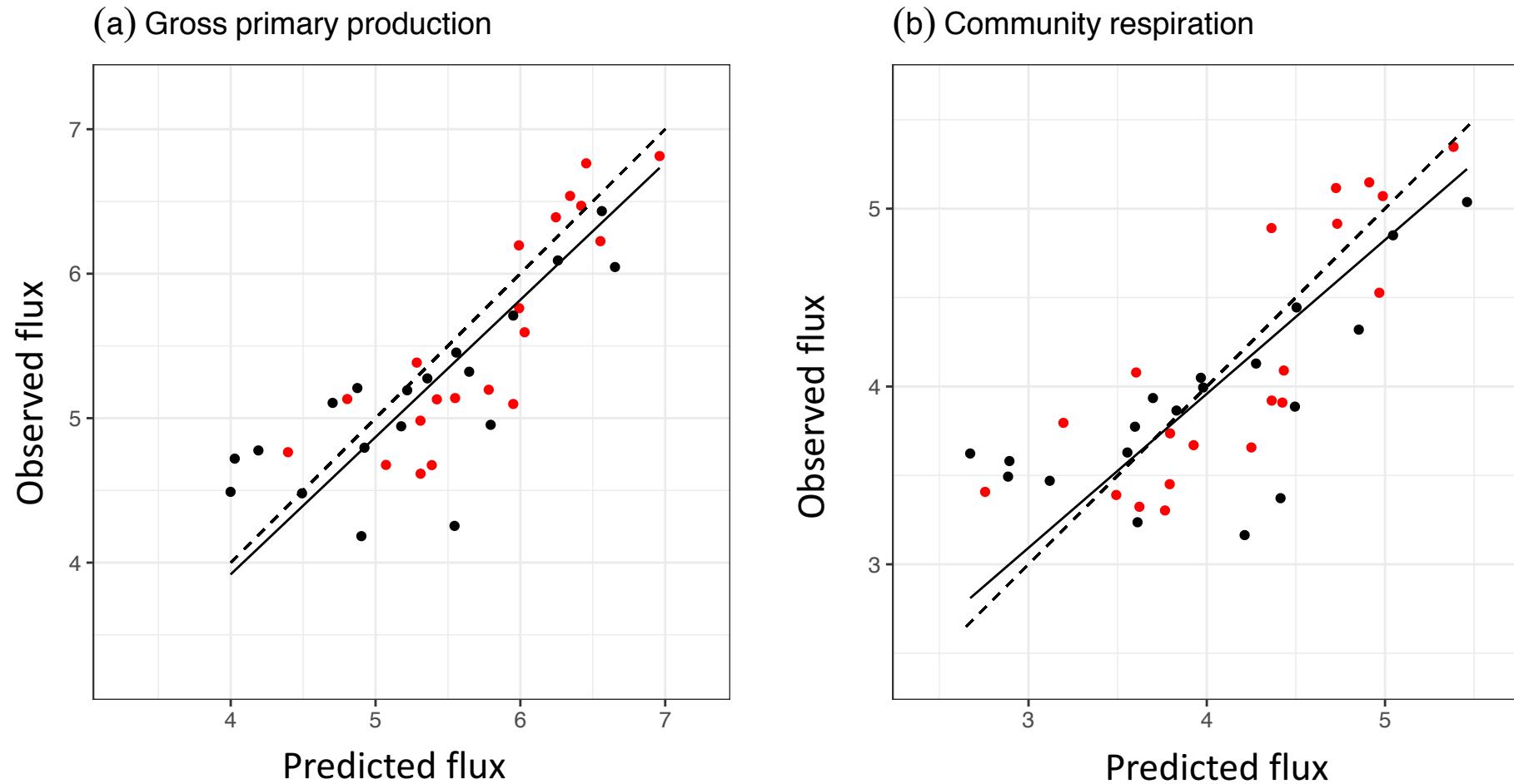
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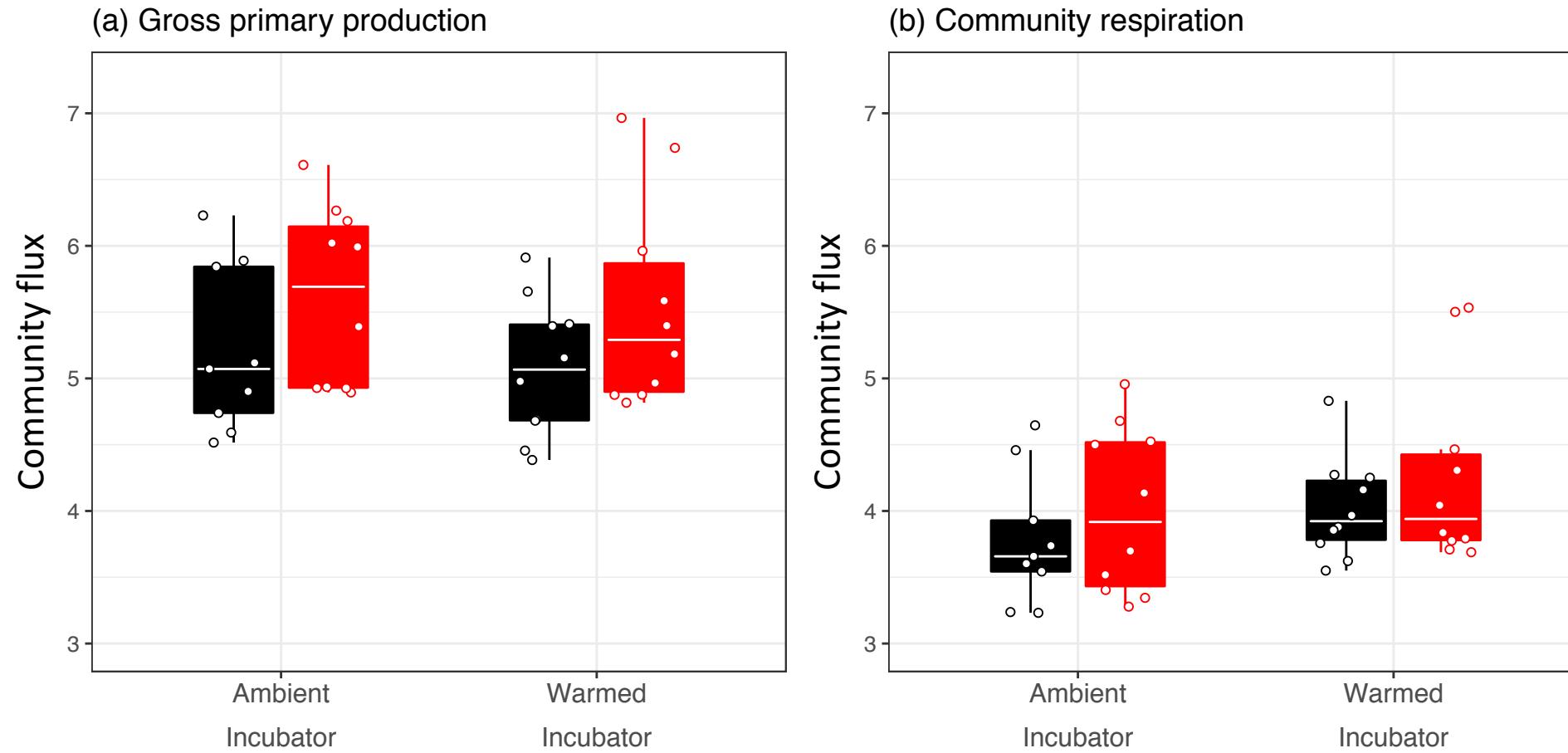
Model does a pretty good job!



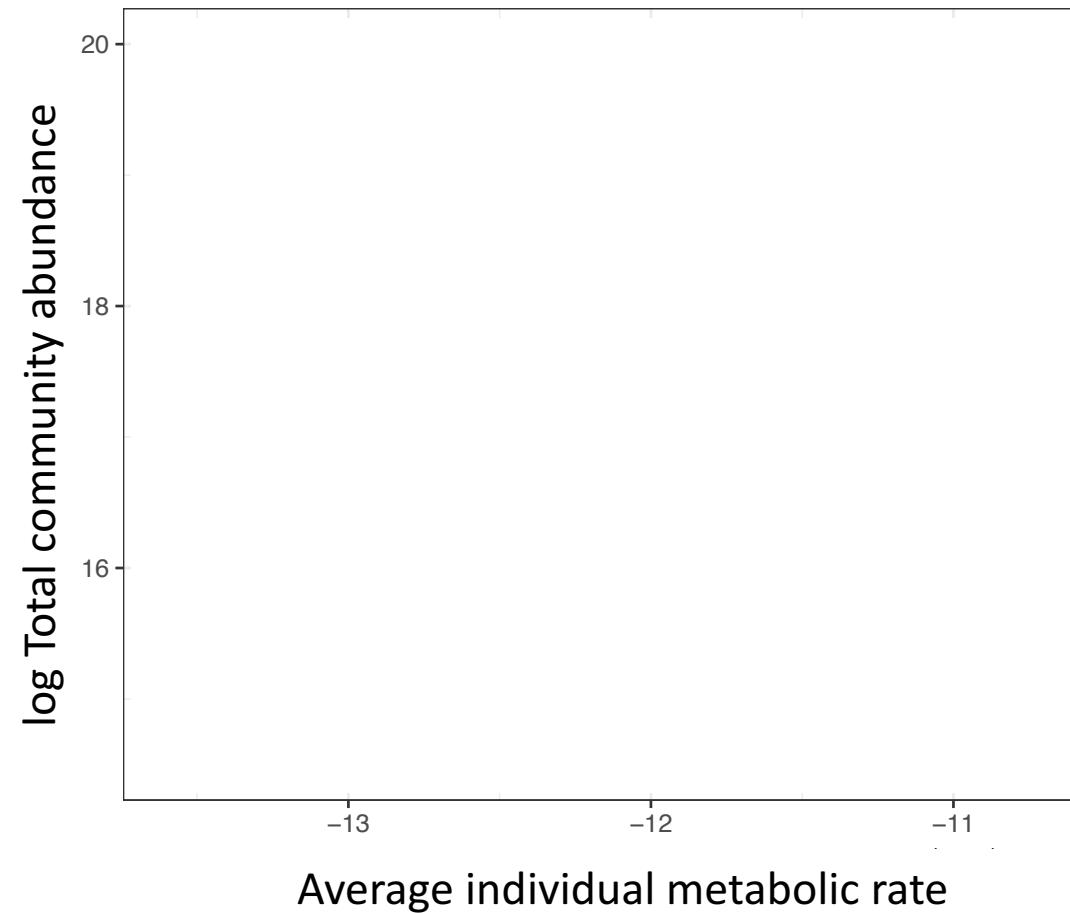
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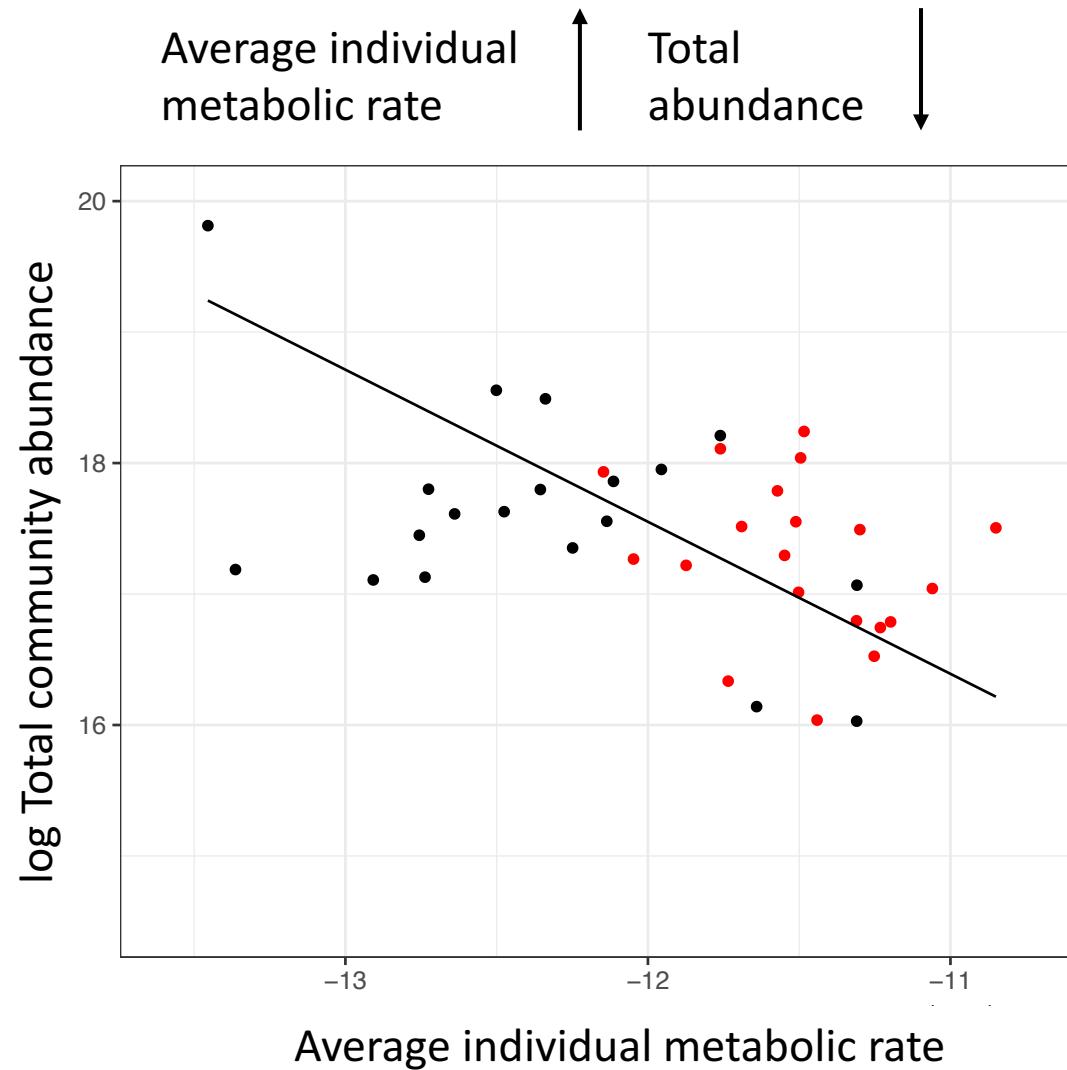
Paradoxical result?



Zero-sum dynamics



Zero-sum dynamics



Conclusions

- Linked the metabolic rates of whole communities to the individual size distribution
- Flow cytometry routinely taken in scientific ship voyages
- As a bonus, validated some key assumptions of metabolic scaling theory

Thanks for listening!