

virtual microbial eco-evolutionary dynamics
evolution predictable???

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TODAY

focus on metabolism

(1) de novo evolution in artificial metabolic universe
(possible reactions cf KEGG but much smaller)

what will happen when the tape is played N times?

(a) from identical initial conditions in a constant environment

(b) from identical initial conditions in a strongly variable environment
and brought in lab conditions (LTEE)

(2) What should we expect to evolve after WGD of YEAST
using its metabolic network as starting condition

what will happen when the tape is played N times?

Compare to what happened in the unique case of YEAST on earth

(3) "causal drift": what changes in metabolic rates cause diabetes?

“Virtual Microbes”

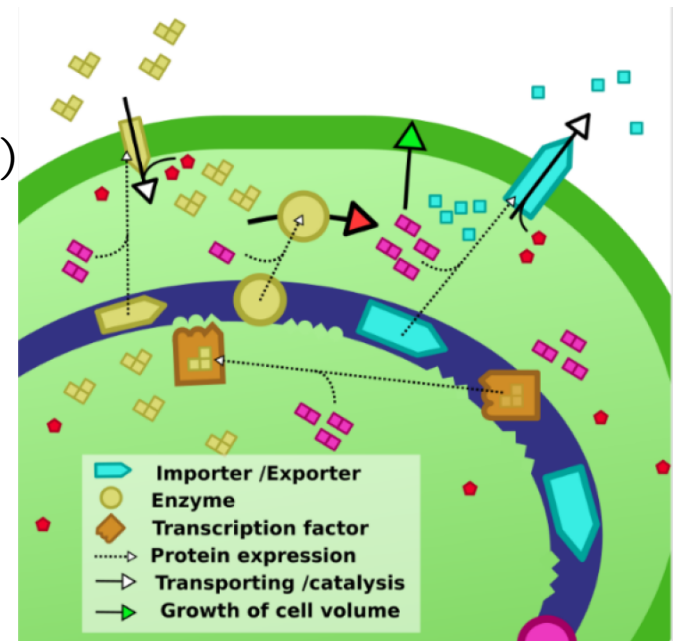
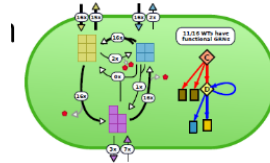
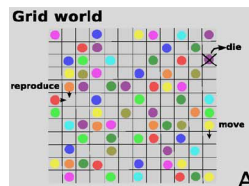
a paradigm system for bottom up modeling of multiple modes of adaptation in biological-like complex complex adaptive systems

Thomas Cuypers and Bram van Dijk

Cell with
Genome with
genes (TFs, pumps, enzymes) with
parameters (V_{max} , K , binding)
metabolism
grow and divide
Mutate
(duplication/deletions, HGT, par. changes)

In 'universe'
potential metabolic
reactions
Resource influx
space

NO PREDEFINED FITNESS



Metabolic universe

{ Meijer , v. Dijk & H.
2020

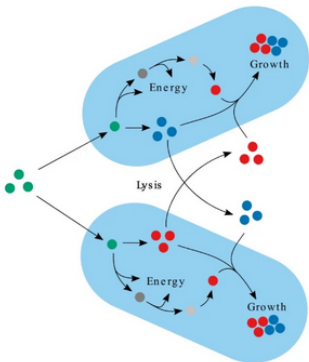
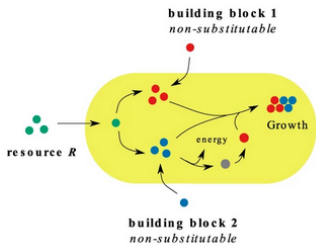
Table 1 Metabolic universe: Using VirtualMicrobes, we generated an artificial biochemistry comprised of 9 metabolites, 8 importers, 8 exporters and 43 conversion reactions, with two non-substitutable building blocks and a single energy molecule.

From: [Contingent evolution of alternative metabolic network topologies determines whether cross-feeding evolves](#)

Metabolite	Class	Diffusion rate	Degradation rate	Toxicity	Mass	Influx rate
R	Resource	0.011	0.0003	0.077	9	0.002
B1	Building block	0.011	0.0100	0.058	8	0
B2	Building block	0.012	0.0100	0.103	8	0
E	Energy carrier	0.050	0.1000	0.065	1	0
M1	-	0.014	0.0006	0.105	6	0
M2	-	0.019	0.0003	0.080	4	0
M3	-	0.013	0.0006	0.079	7	0
M4	-	0.014	0.0014	0.158	6	0
M5	-	0.016	0.0008	0.047	5	0
Reactions						
R → B2 + E	B1 → M4 + 2 E	M1 → M2 + 2 E	M2 + M5 → B1	M3 + M5 → R		
R → B1 + E	B1 → M5 + 3 E	M2 + M3 → B1	M2 + M5 → M1	M3 → M5 + 2 E		
R → M1 + 3 E	B2 → M1 + 2 E	M2 + M3 → B2	2 M1 → R	M3 → M4 + E		
R → M2 + 5 E	B2 → M2 + 4 E	M2 + M5 → M4	2 M2 → M1	M4 → M2 + 2 E		
R → M4 + 3 E	B2 → M3 + E	M2 + M4 → M3	2 M2 → B1	M4 + M5 → B2		
R → M5 + 4 E	B2 → M4 + 2 E	M2 + M5 → M3	2 M2 → B2	M4 + M5 → R		
R → M3 + 2 E	B2 → M5 + 3 E	M2 + M4 → B2	2 M2 → M4	M4 + M5 → B1		
B1 → M1 + 2 E	M1 + M4 → R	M2 + M5 → B2	2 M2 → M5	M5 → M2 + E		
B1 → M2 + 4 E	M1 + M2 → M3	M2 + M5 → R				
Importers: one for each non-energy metabolite (8 total)						
Exporters: one for each non-energy metabolite (8 total)						

De Novo evolution in a constant environment (1 resource)

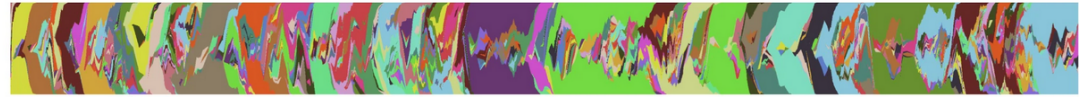
Meijer, v. Dijk & T
2020



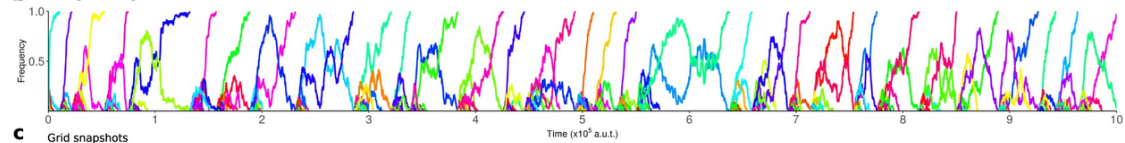
a Metabolically autonomous community (21/60 simulations)

Muller plot of metabolic genotypes

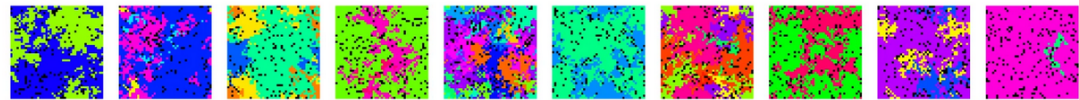
Community 23



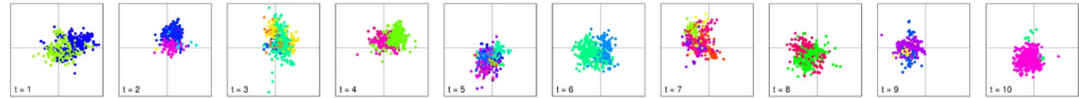
b Lineage tracking



c Grid snapshots



d PCA of single-cell proteomics (one dot = one cell)

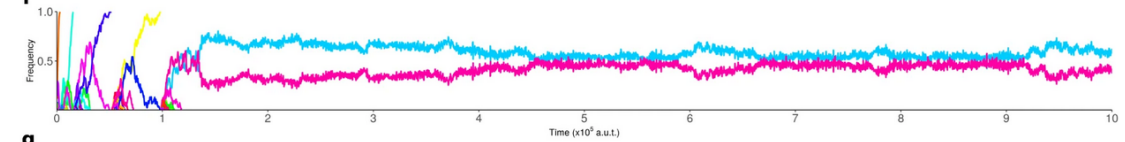


e Cross-feeding community (24/60 simulations)

Community 8



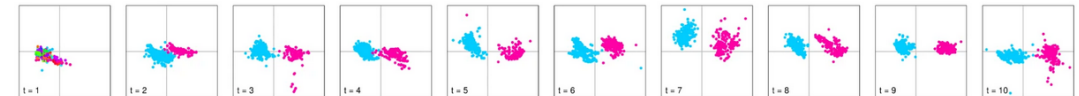
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g



h



2 types of Evolved metabolism generate predictable ecosystems

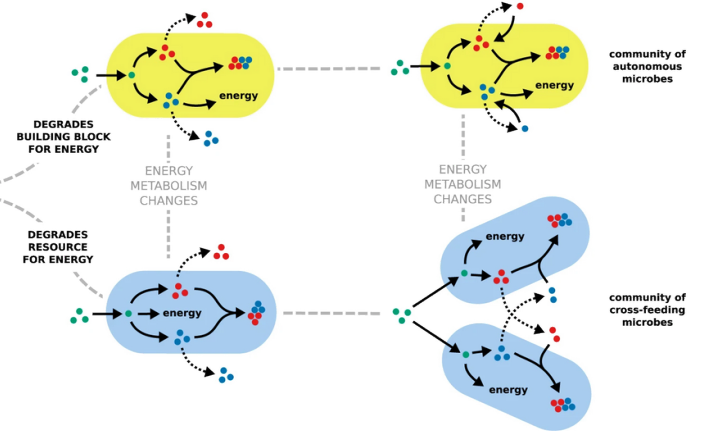
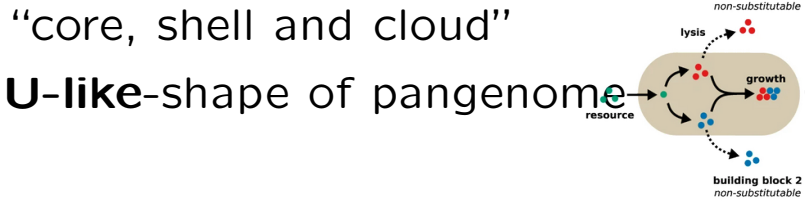
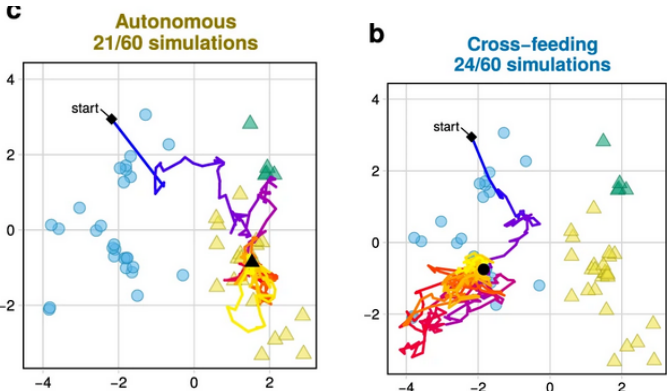
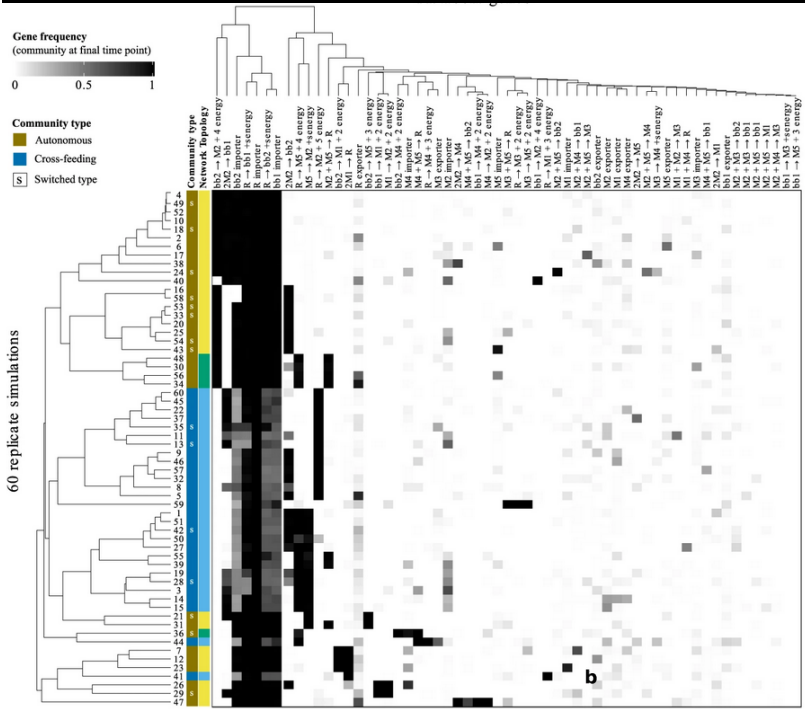
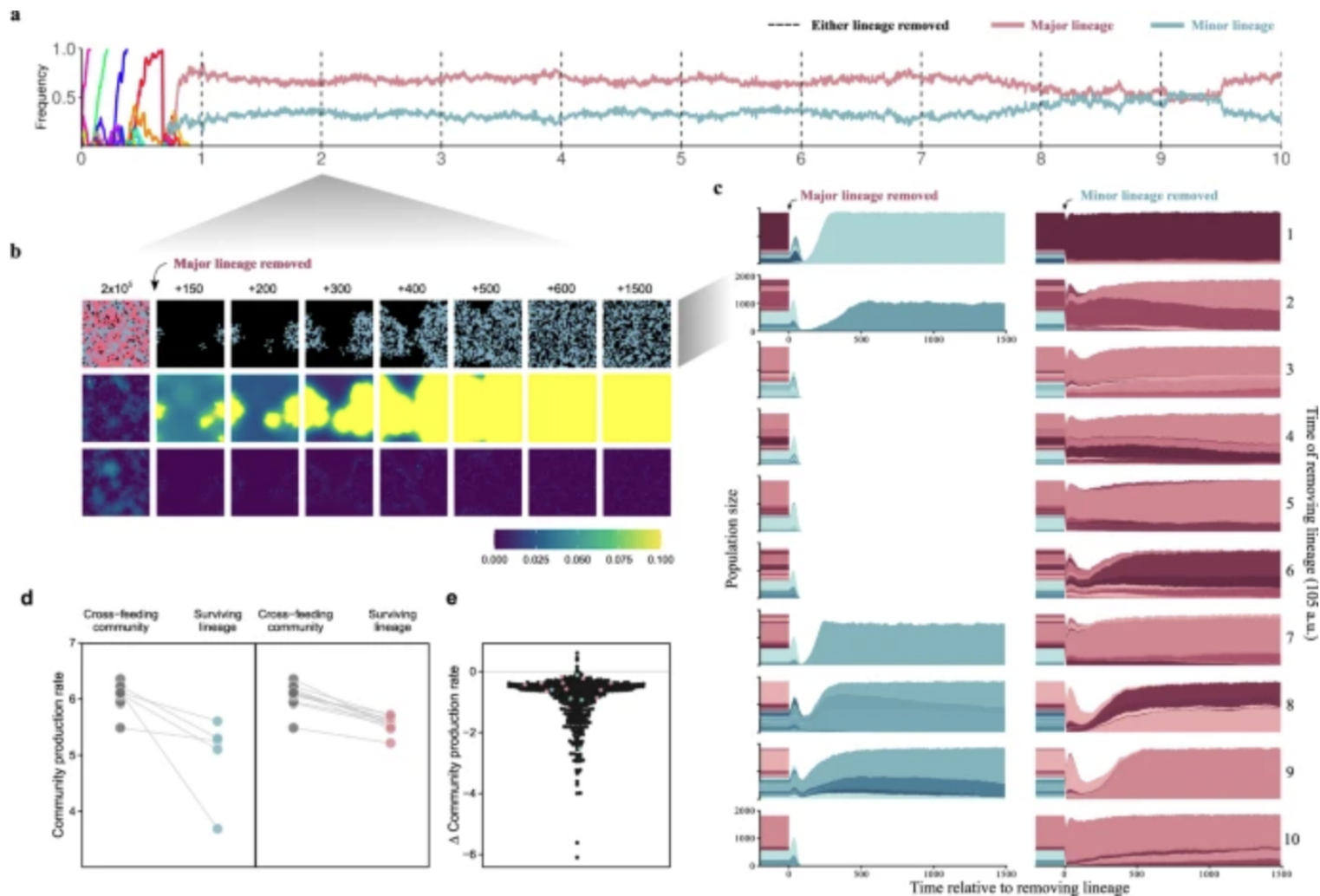
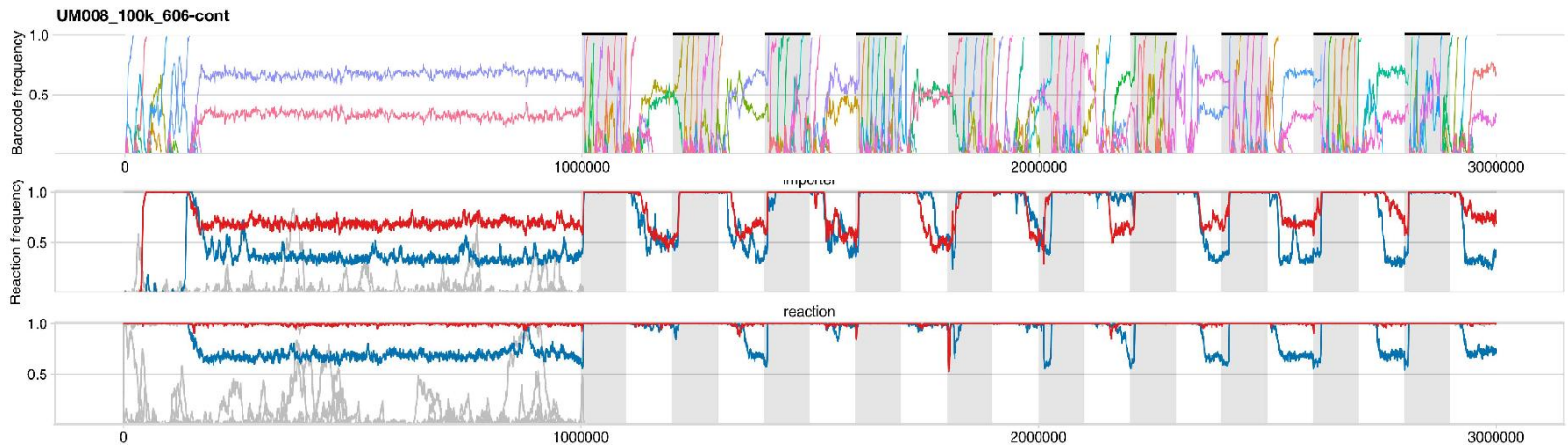
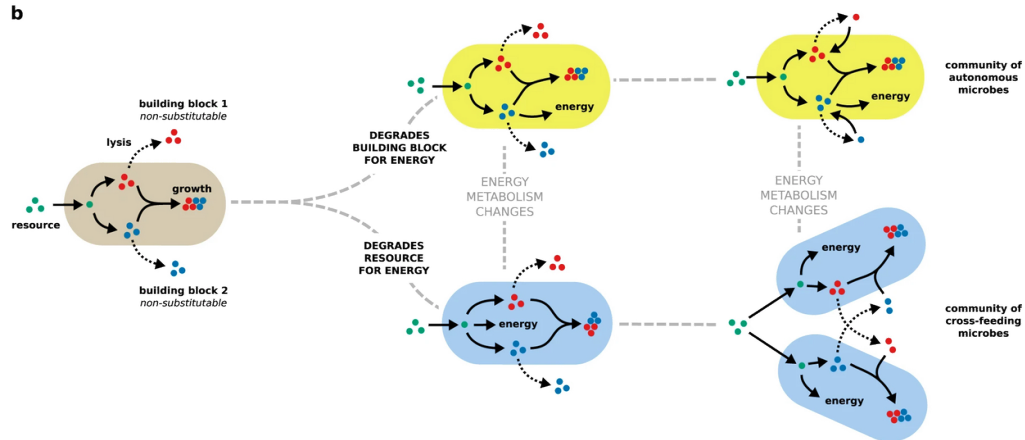


Fig. 3: Metabolic dependencies in cross-feeding communities.



Cross-feeding evolves in 1 of 2 types of metabolism IN SPACE

self-sufficiency regained when mixed (switching)

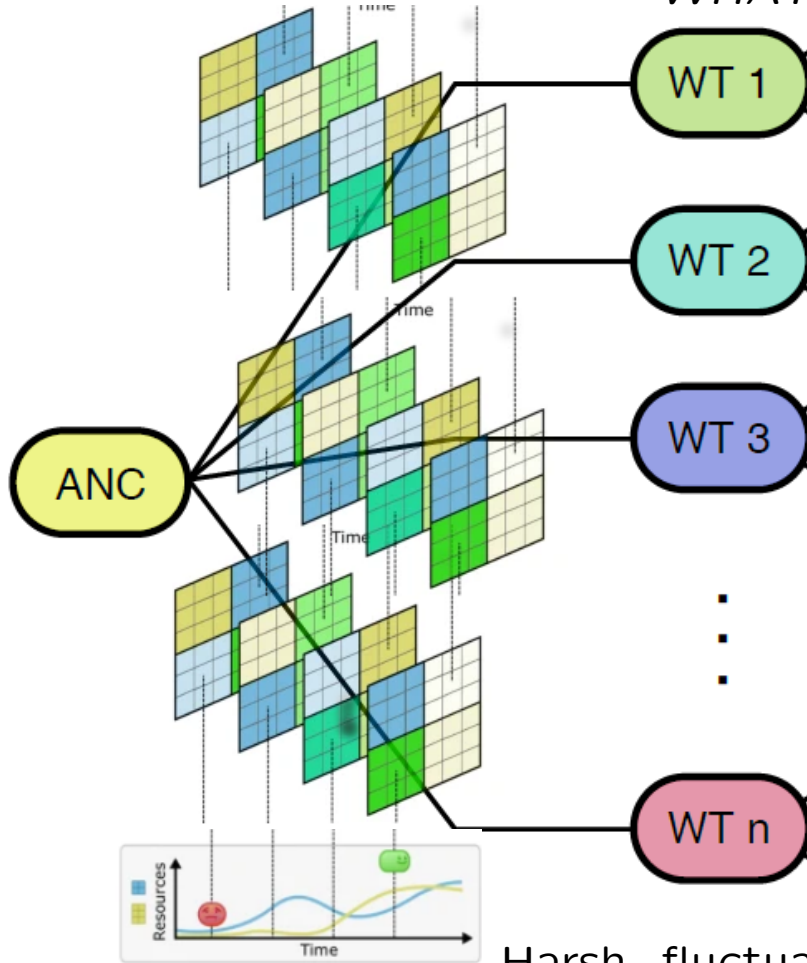


conclusion

- Crossfeeding can evolve in space without explicit costs/tradeoffs or environmental variability
- Selfsufficiency can also evolve in exactly the same "universe"
- Crossfeeding and selfsufficiency contingent outcomes from their LUCA
- Crossfeeding and selfsufficiency are predictable outcomes from evolved metabolism
- selfsufficient mutants exist in crossfeeding ecosystem but do not take over
- switching spatial system (biofilm) to wellmixed lead to switching between crossfeeding and selfsufficiency

De Novo Evolution in variable environment

“WHAT” has evolved?, How to observe?



LCA of evolved population

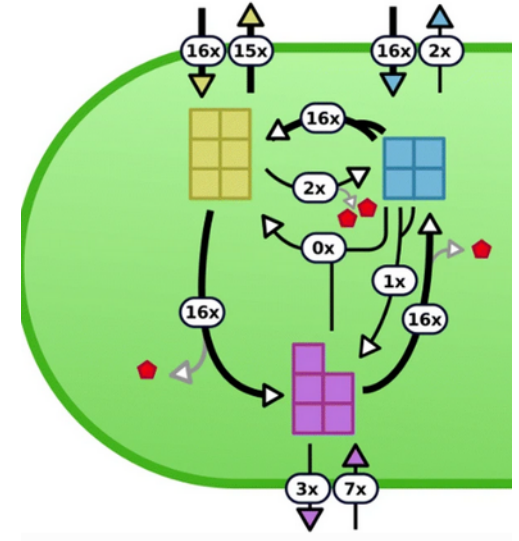
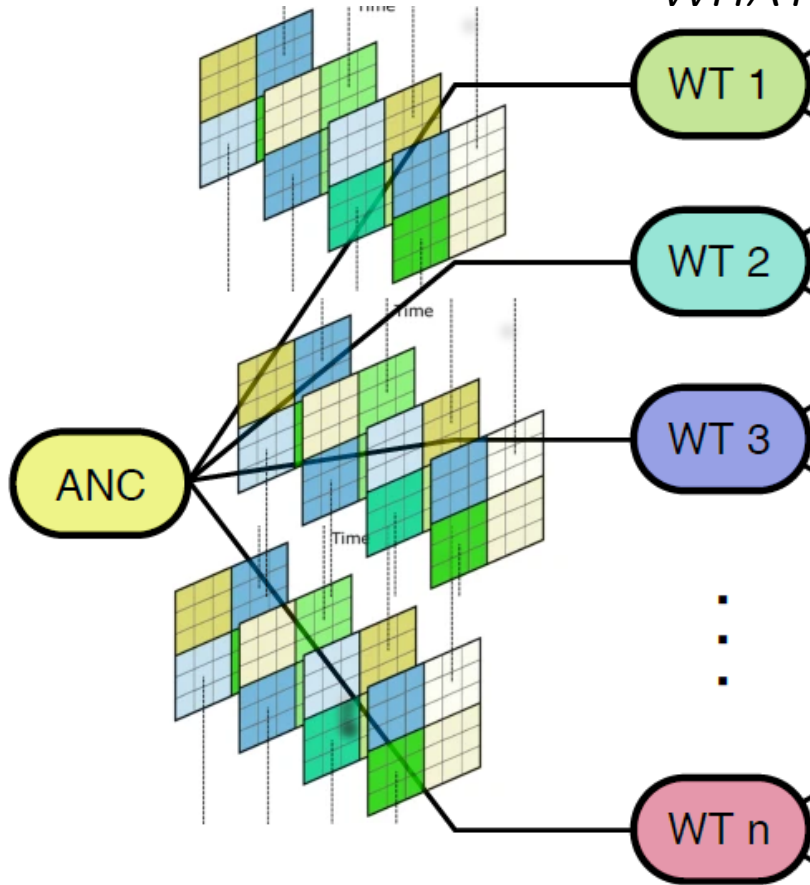
Harsh, fluctuating environment (2 resources)

Identical for all replicates

van Dijk et al 2019

De Novo Evolution in variable environments

“WHAT” has evolved?, How to observe?



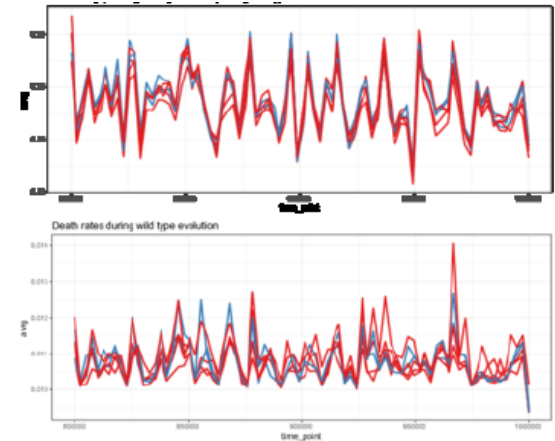
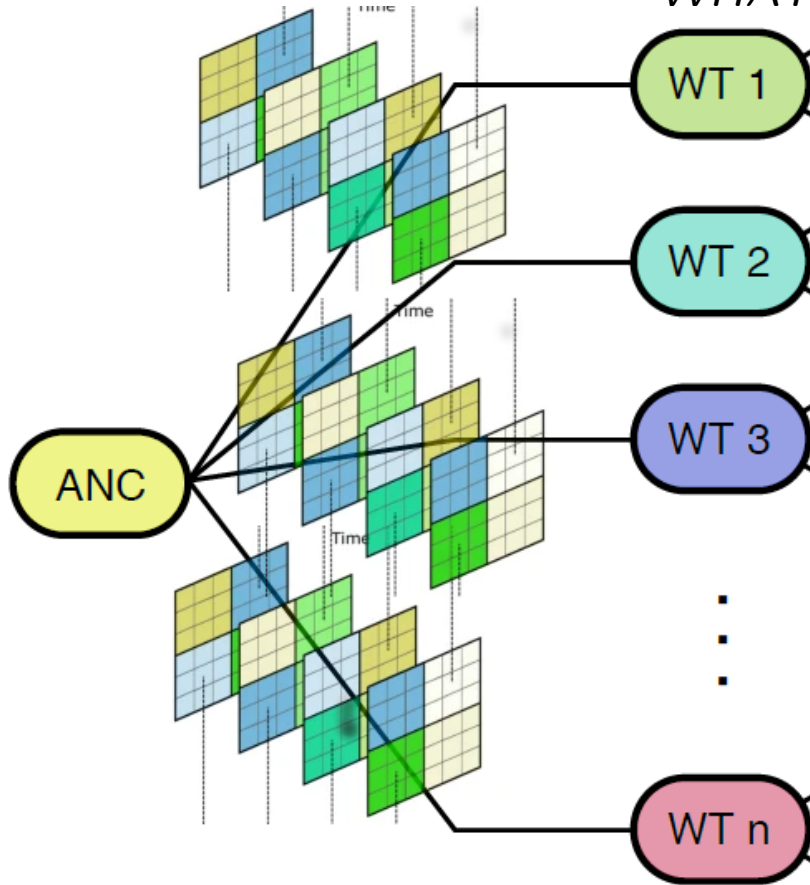
Common metabolic cycle

LCA of evolved population

Harsh, fluctuating environment
Identical for all replicates

De Novo Evolution in variable environments

“WHAT” has evolved?, How to observe?



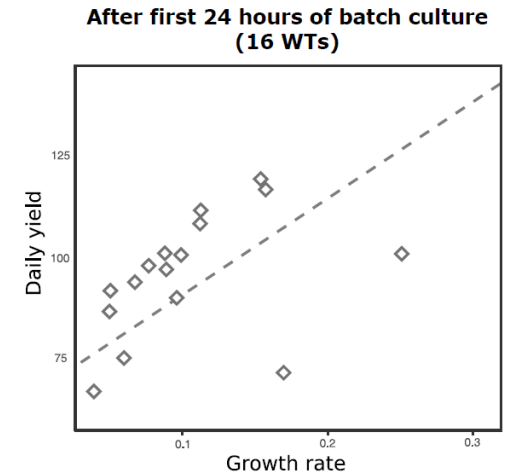
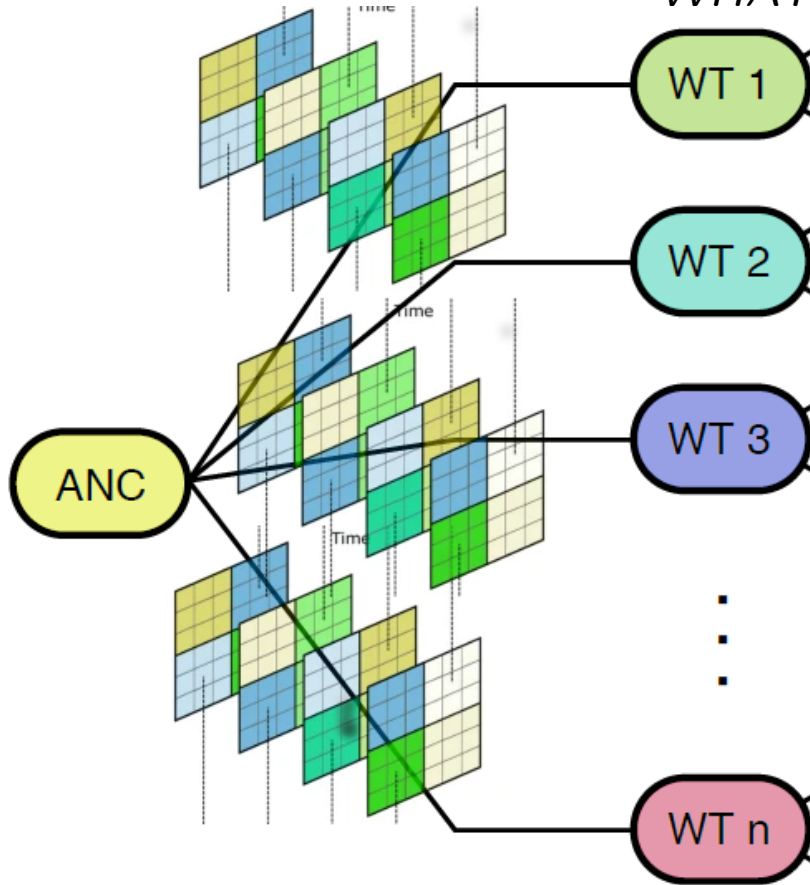
Similar “fitness”

LCA of evolved population

Harsh, fluctuating environment
Identical for all replicates

De Novo Evolution in variable environments

“WHAT” has evolved?, How to observe?

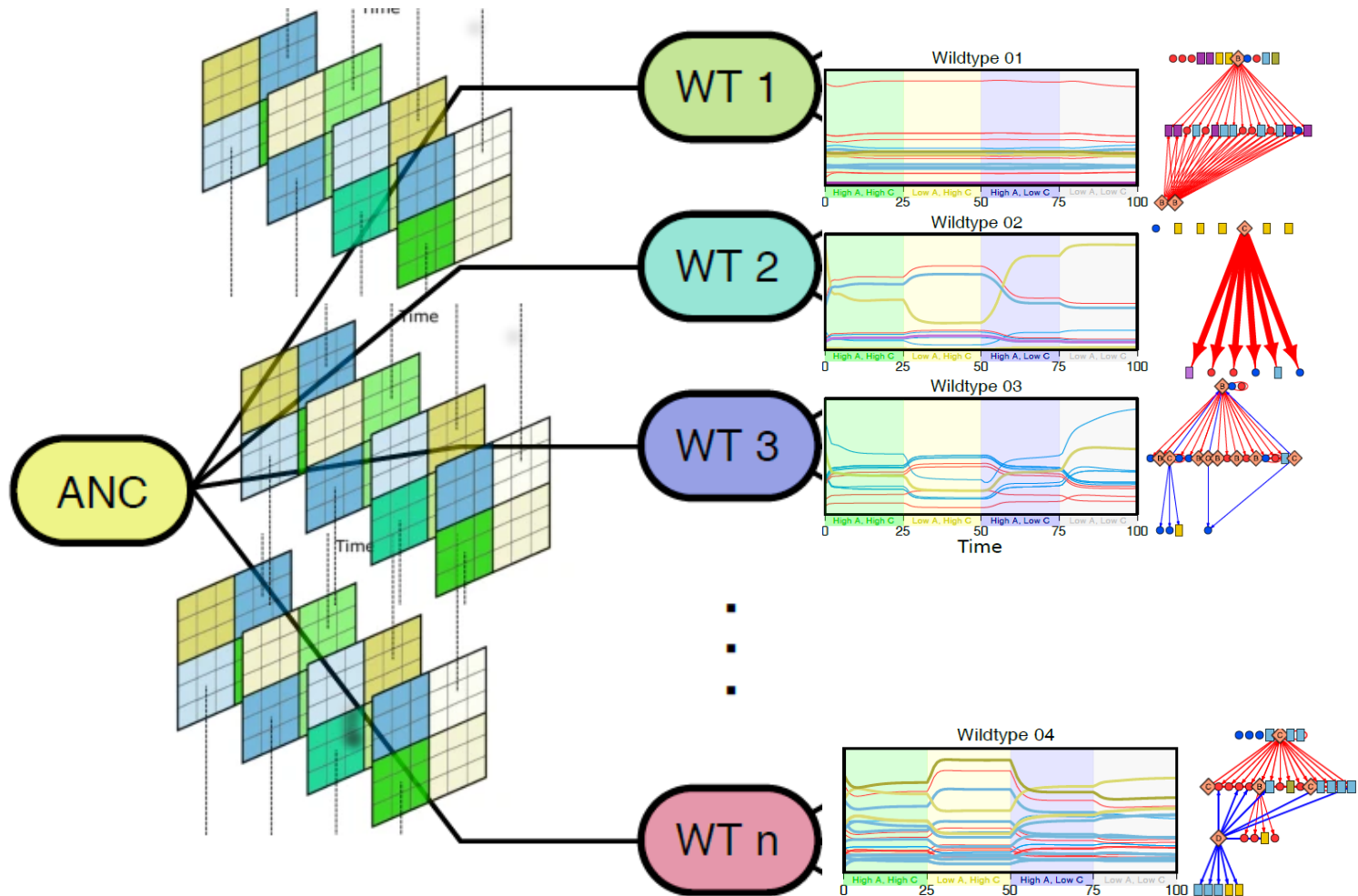


Dissimilar “fitness”

LCA of evolved population

Harsh, fluctuating environment
Identical for all replicates

BUT: very diverse GRN (or none) and metabolic reaction to alternative environments



Experimental evolution: starting with pre-evolved “wildtypes”

Well known example of experimental evolution:

Long term evolutionary experiment (LTEE) (Lensky 1991)
One strain of E.Coli is evolving in lab-conditions since 1988
(>70000 generations) in 12 replicates
in a serial transfer protocol (diluted in new medium every 24 hours)
still adapting (getting “better”)
Continued new ways of observing & new insights

This case study:

In silico evolution of the above pre-evolved “wildtypes” (WT 1-16)
in similar serial transfer protocol

study “generic” features of such an evolutionary process

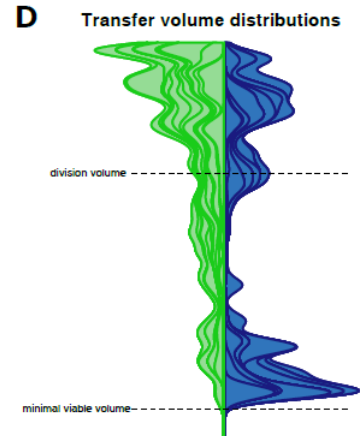
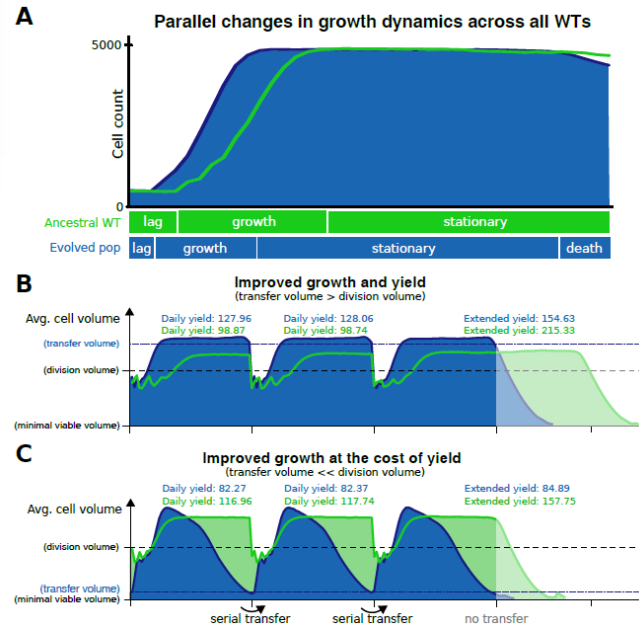
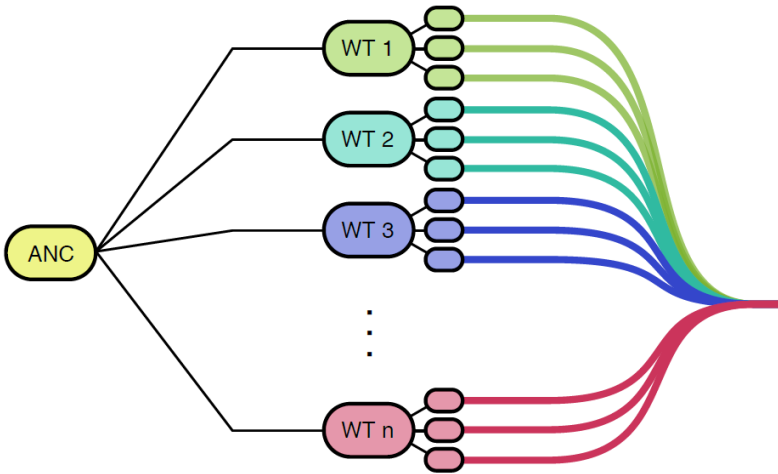
To WHAT does the population adapt?

HOW does it adapt?

Multiple observables

Similarities/differences to E. coli?

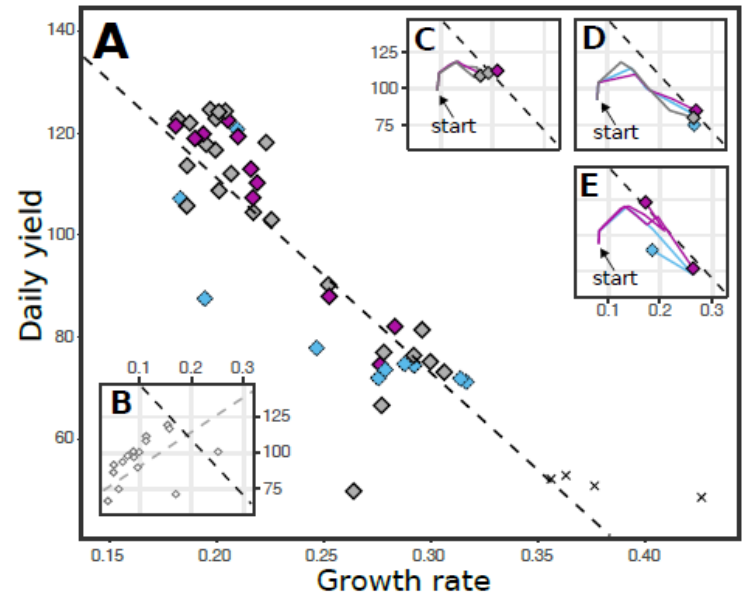
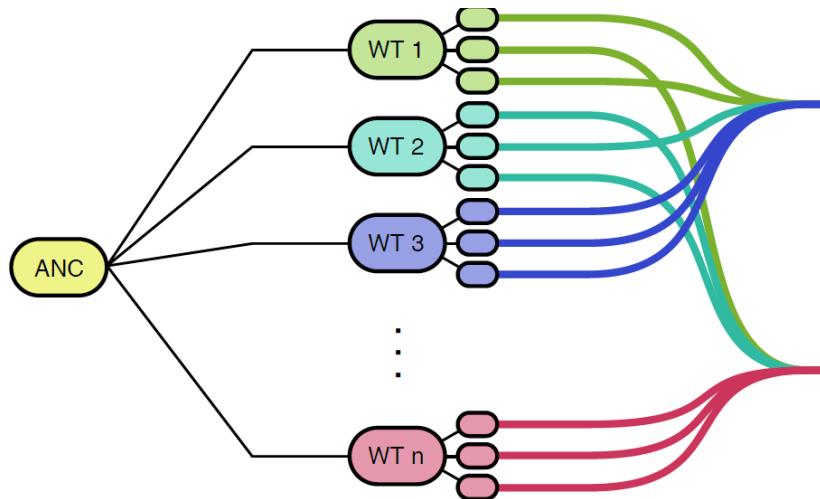
In serial transfer protocol they all evolve to “Trust the hand that feeds them” (anticipate 24 hr cycle)



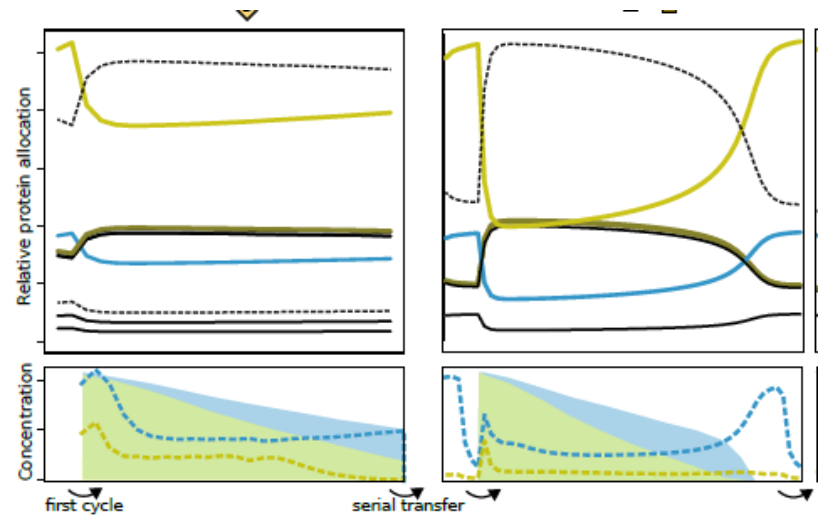
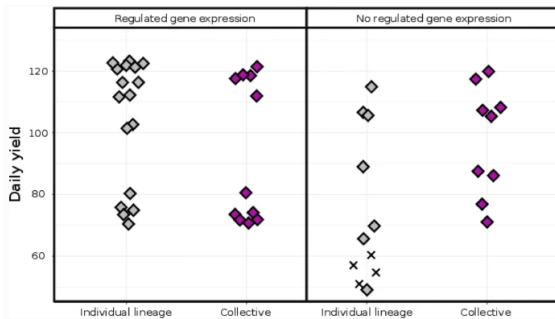
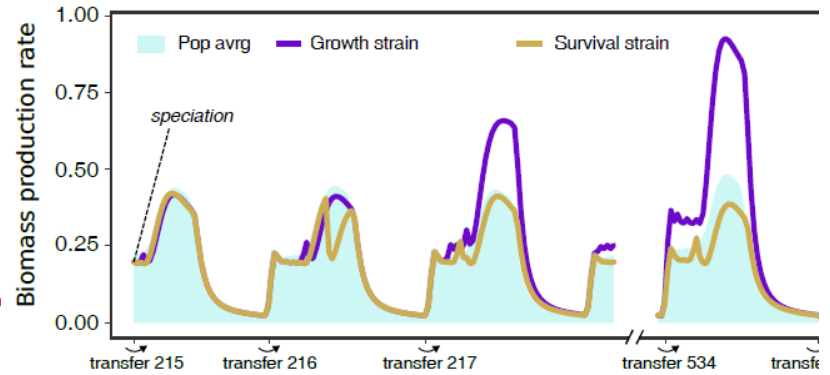
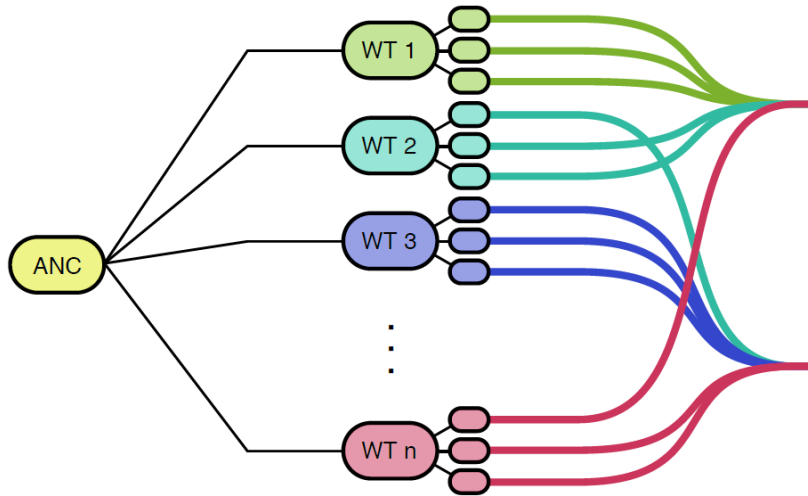
Minimize Lag-phase
Exhaust all food after 24 hours
remaining JUST alive
OR
remaining JUST ready to divide

Maximizing growth rate OR Yield

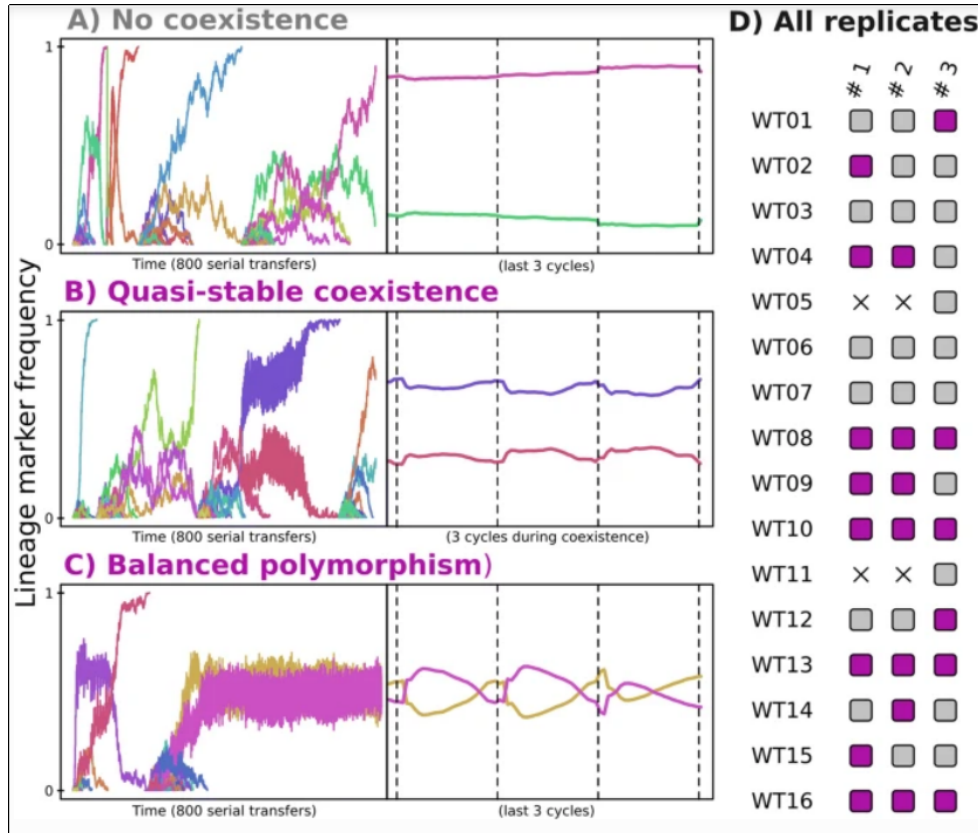
evolved trade-off and distinct strategies



By individual based regulation OR collective tuning



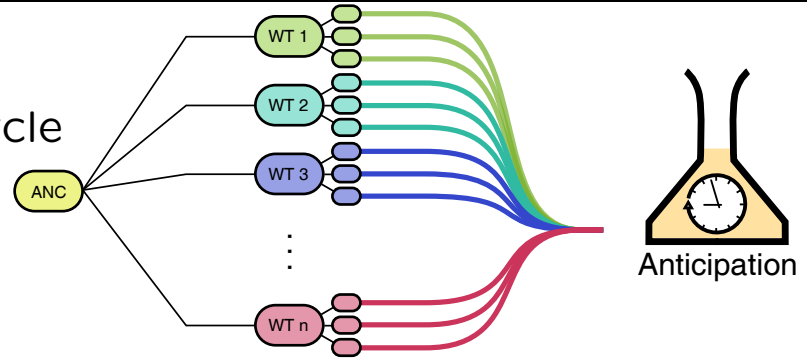
By individual based regulation OR collective tuning



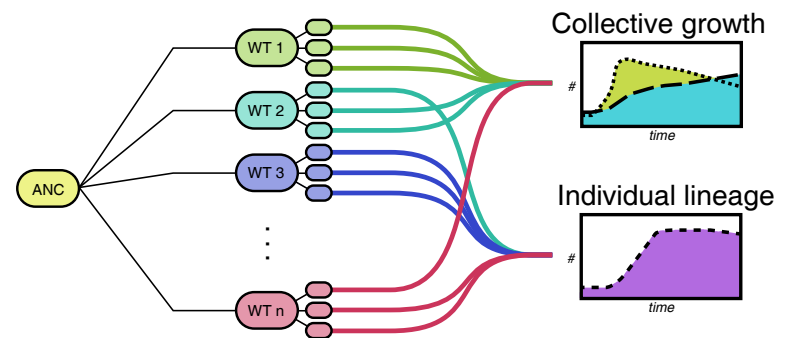
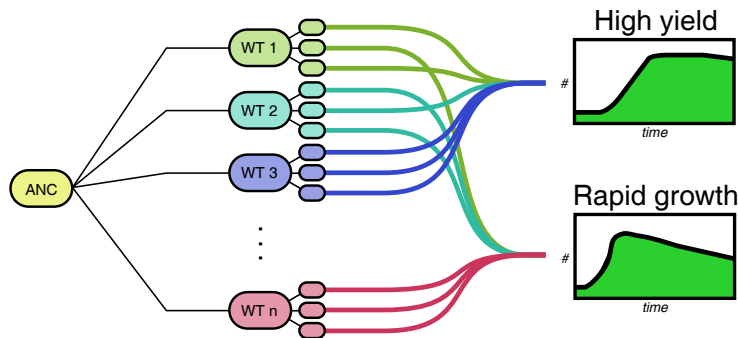
Conclusion

Evolved contingency of predictability combinatorial set of discrete outcomes

Diversified evolved wildtypes
all evolve anticipation of 24 hr cycle
un-predicted predictability

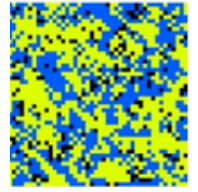
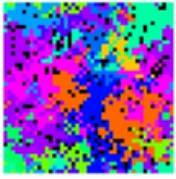


BUT in unpredictable ways



Some WT's adapt in a predictable way , others in very different ways

predictability is an unpredictable outcome of evolution



Conclusions/Observations

- What is fitness / what has evolved not obvious
- Evolutionary attractors can be characterized as a combinatorial set of a limited set of alternatives
- Autonomous and Collective “problem solving” (metabolism)
“easy” alternatives
- Non-autonomy not because of lack of genes...
- Spatial embedding, also without spatial patterns important
- Trade-off’s not innate but evolved properties
- GRN very variable (presence and shape)
- Predictability, even in well defined environments depends on prior evolution
Predictability is an unpredictable outcome of (prior) evolution