

Predation, Parasitism, and Performance within EBPRs

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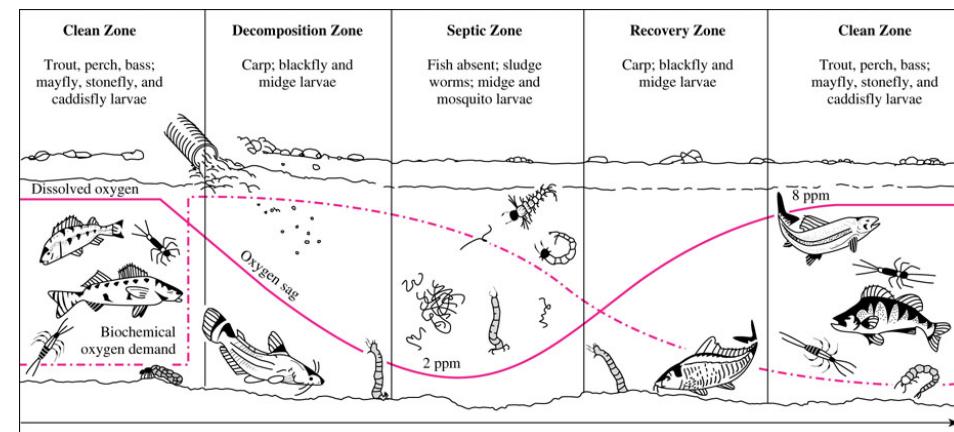
Motivation*

- Release of excess nutrients can lead to eutrophication
 - Hypoxia → fishkills

Raw Wastewater	~ 12 mg P/L
Secondary Treatment	~ 6 mg P/L
Protected Watersheds	1 mg P/L



- Enhanced biological phosphorous removal (EBPR) technologies offer a solution

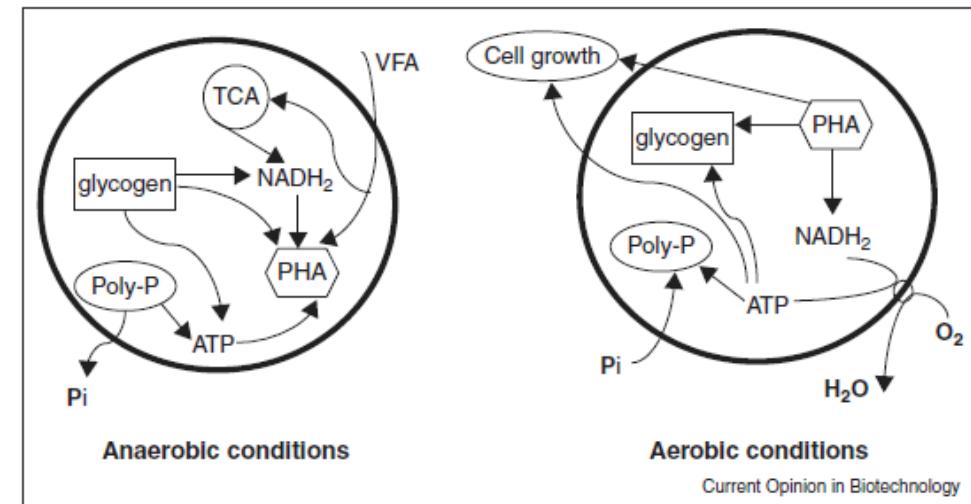


Enhanced Biological Phosphorous Removal

Phosphate Accumulating Organisms (PAO)

- Accumulate 2-5 times as much P as typical heterotrophic bacteria

Anaerobic \longleftrightarrow Aerobic

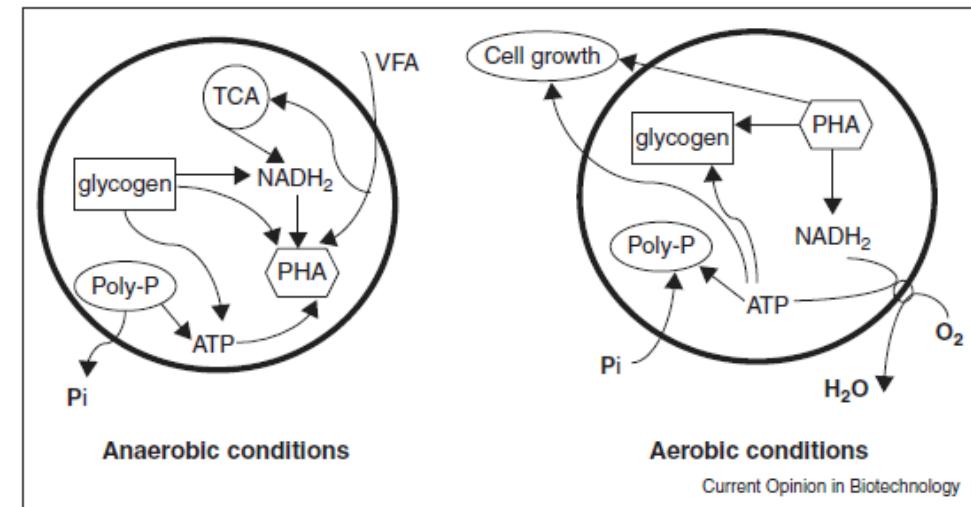


Enhanced Biological Phosphorous Removal*

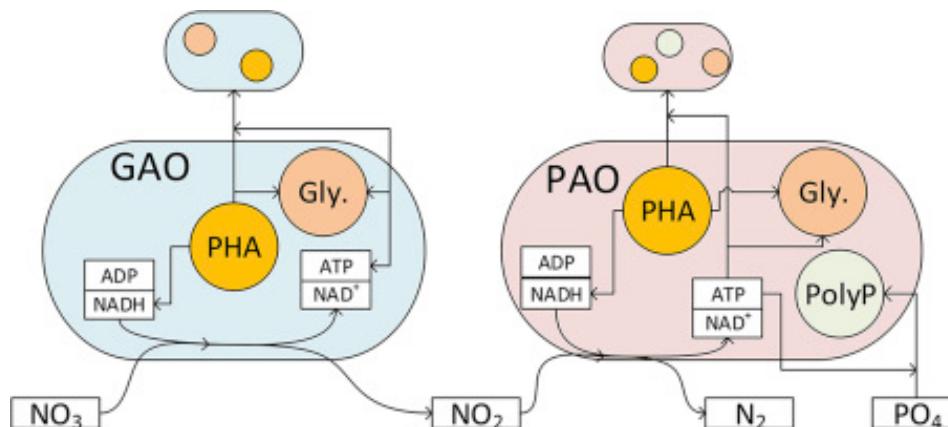
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Current Opinion in Biotechnology

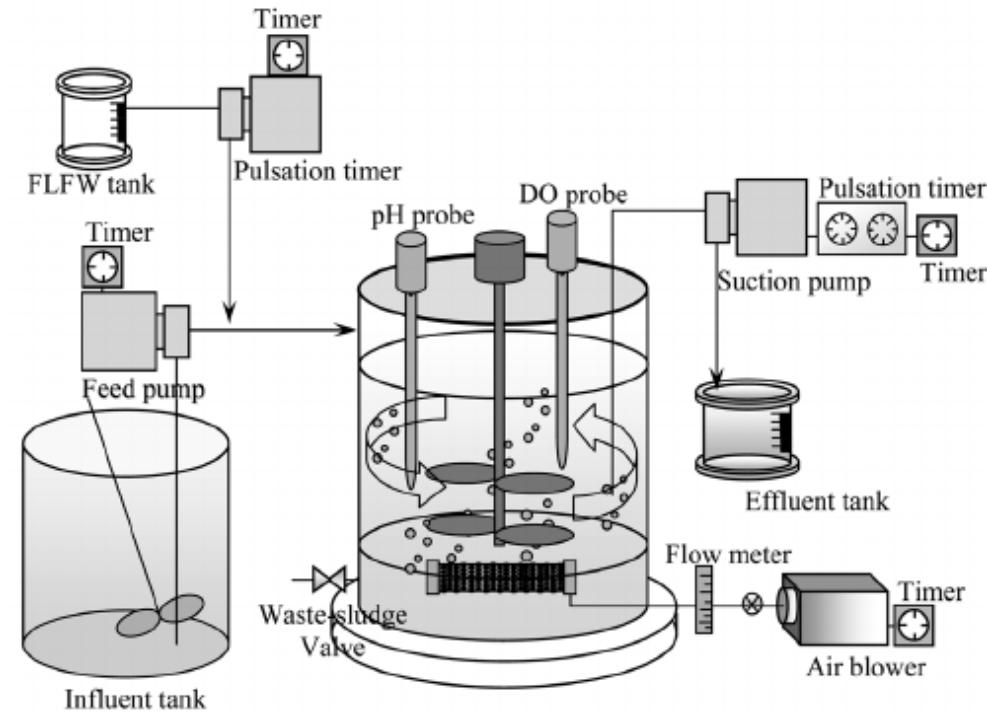


Glycogen Accumulating Organisms (GAO)

- Cooperate?
- Compete?
 - (for VFA)

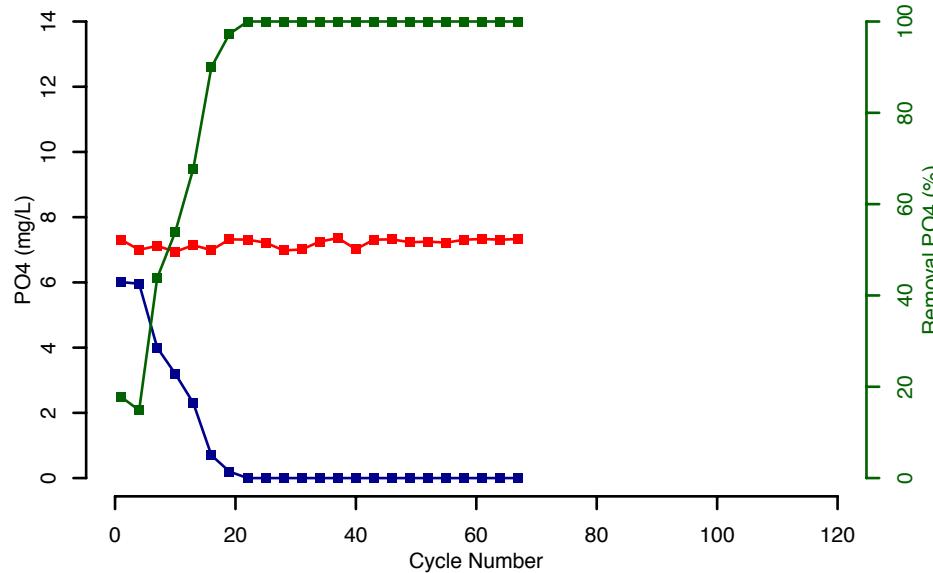
Experimental Approach / Methods*

- 2 x SBRs
 - Experiment S
 - 23 days = 67 cycles
 - Experiment O
 - 40 days = 120 cycles
- 16S rRNA
 - Experiment S
 - 2 sampling events (beginning/end)
 - Experiment O
 - 7 sampling events
- Statistical analysis
 - Ordination (PCoA)
 - Clustering (UPGMA)
 - Differential Abundance Testing



Kinetic Behavior

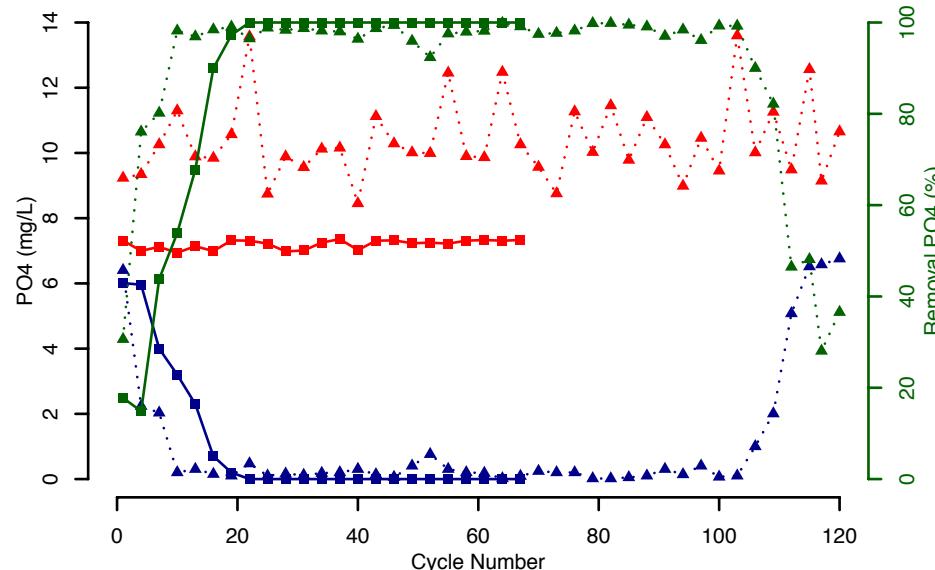
- Experiment S
 - Ideal startup and operation
 - Near complete removal after 7 days (21 cycles)



- Phosphate
 - Initial
 - Final
 - Removal (%)

Kinetic Behavior

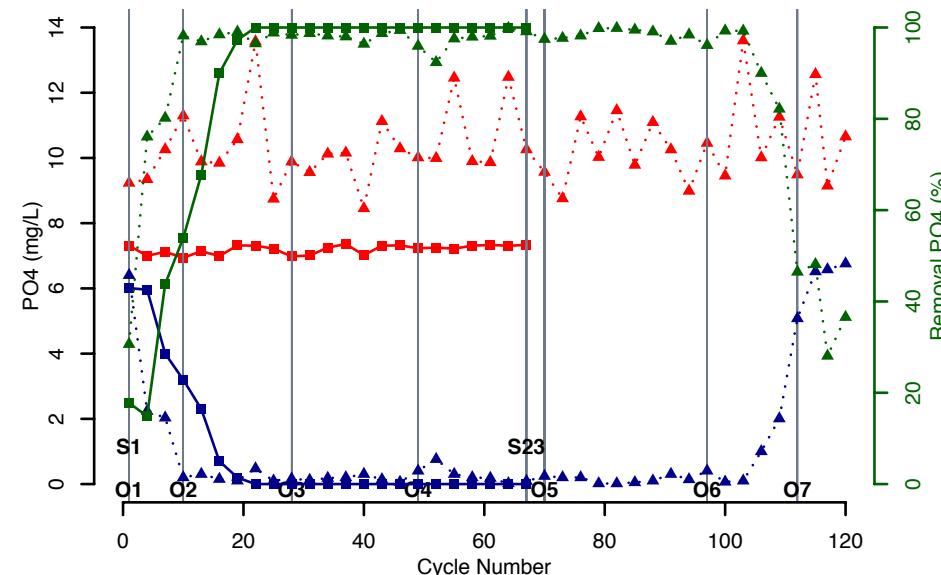
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- Experiment O
 - Similar startup
 - Crash in performance on 35th day



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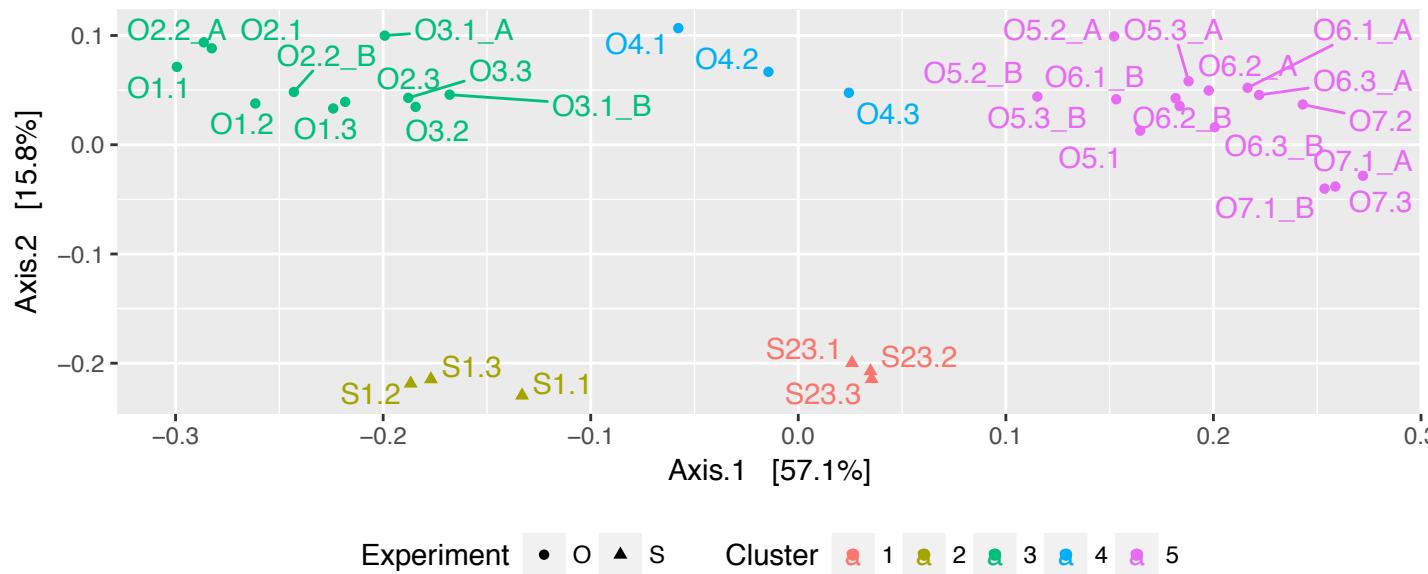
Kinetic Behavior*

- Experiment S
 - Ideal startup and operation
 - Near complete removal after 7 days (21 cycles)
- Experiment O
 - Similar startup
 - Crash in performance on 35th day
- Use sequencing results to investigate the cause of the crash.



- Phosphate
 - Initial
 - Final
 - Removal (%)

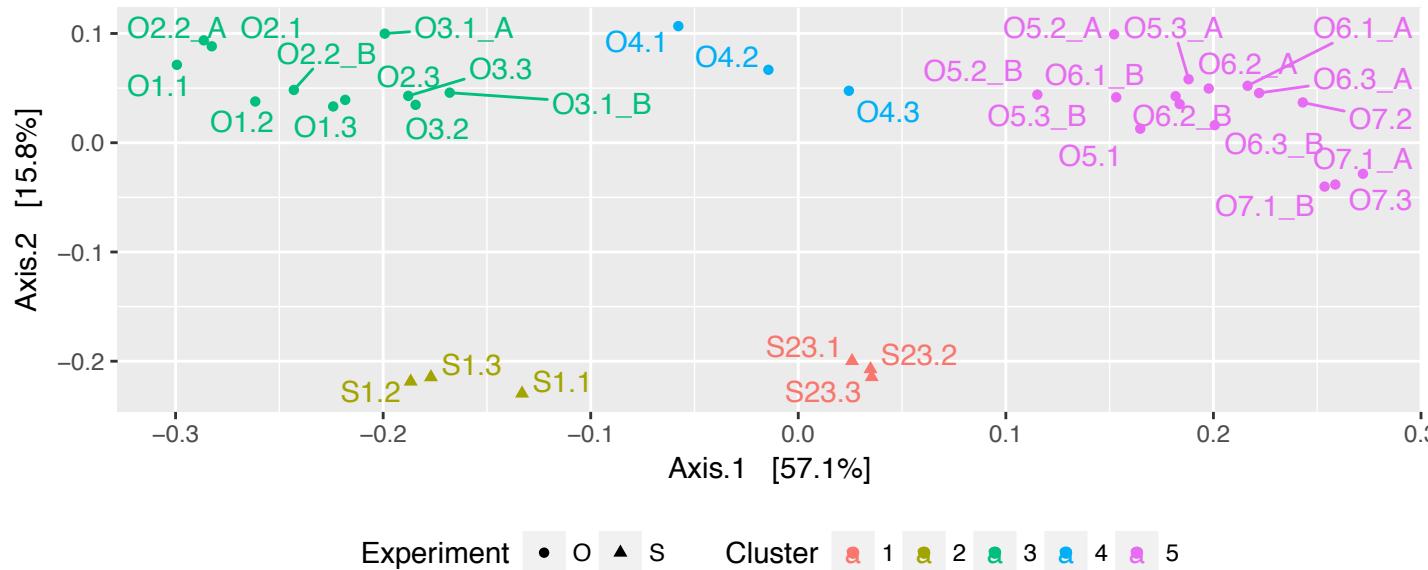
Ordination and Clustering



PCoA + UPGMA

- 5 distinct clusters

Ordination and Clustering*



PCoA + UPGMA

- 5 distinct clusters
- Differential Abundance Testing**
- ~ Experiment + Status

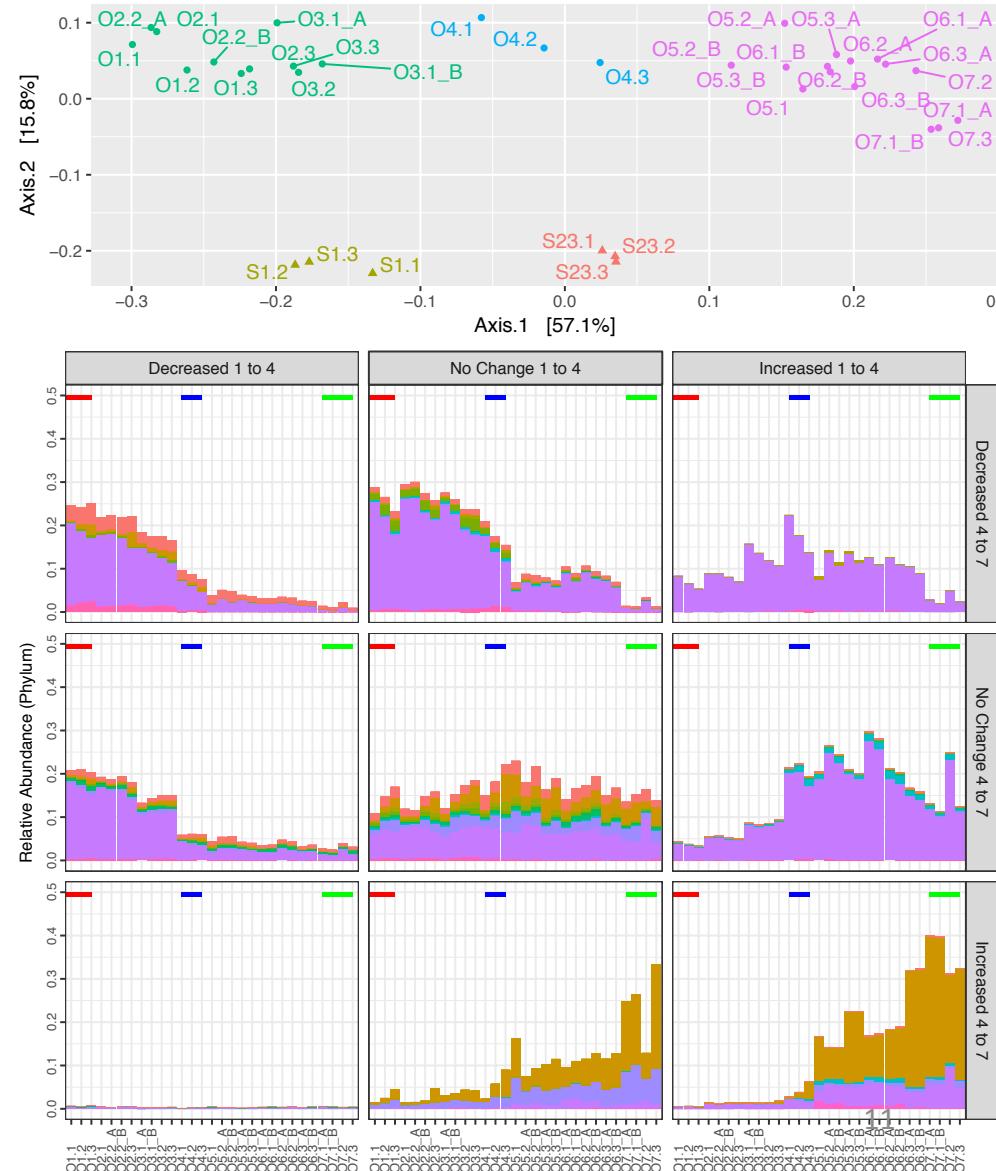
Status	Exp. O	Exp. S
Startup	O1	S1
Attenuation	O2 – O3	
Operation	O4	S23
Failing	O5 – O6	
Crashed	O7	

Differential Abundance & Cohort Plot

Differential Abundance
~ Experiment + Status

Contrasts

- Startup
 - Startup → Operation
- Crash
 - Operation → Crashed
- ASV/OTU State:
 - Decreased
 - No (statistically sig.) change
 - Increased

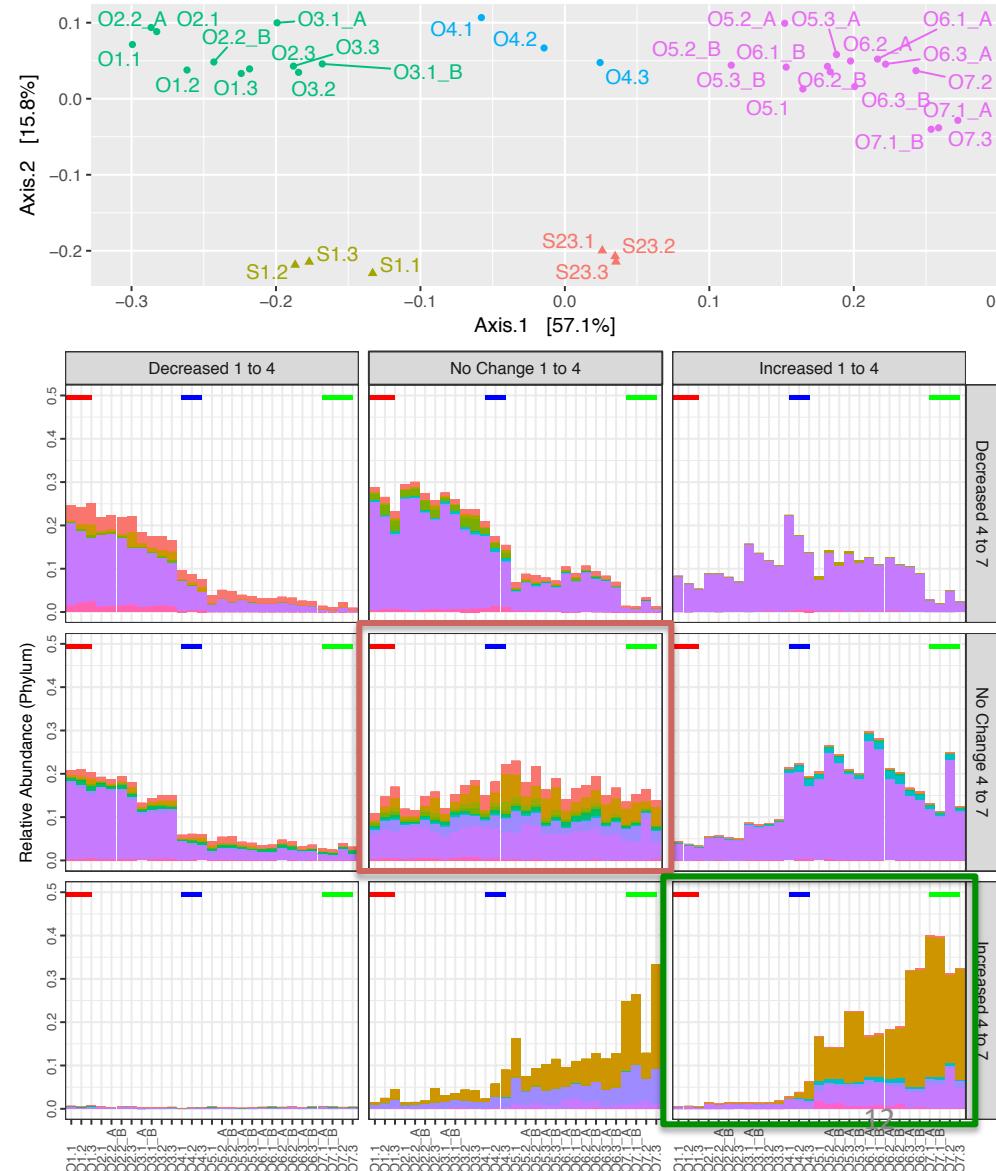


Differential Abundance & Cohort Plot*

Differential Abundance
~ Experiment + Status

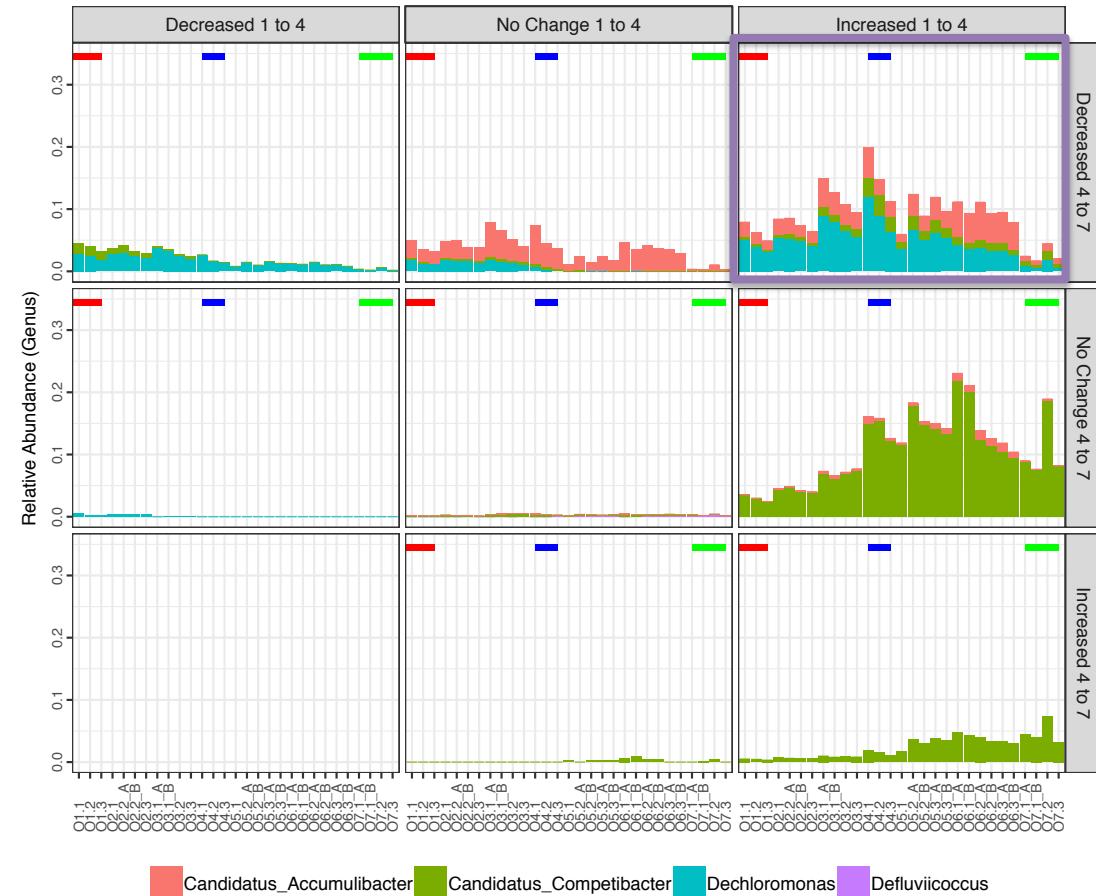
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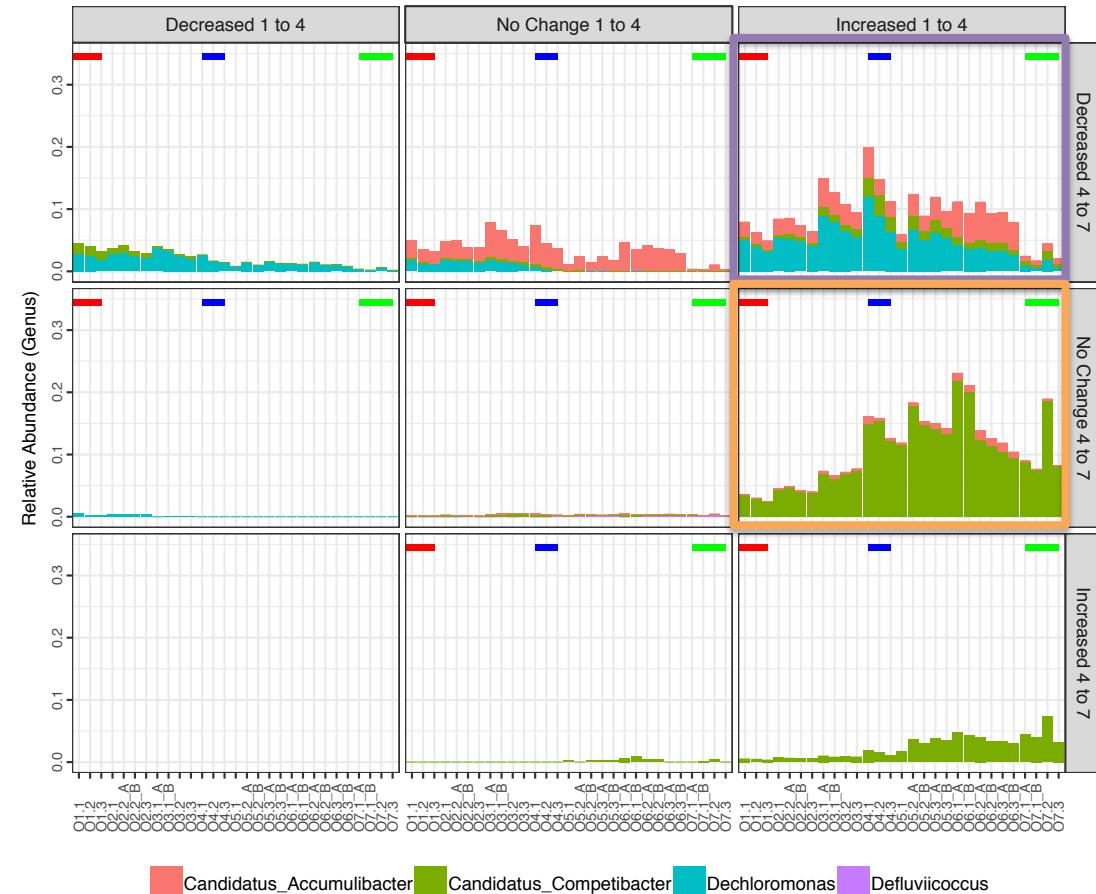
Differential Abundance

- PAO
 - *Ca. Accumulibacter*
 - *Dechloromonas* spp.
- GAO
 - *Ca. Competibacter*
 - *Difluviicoccus* spp.



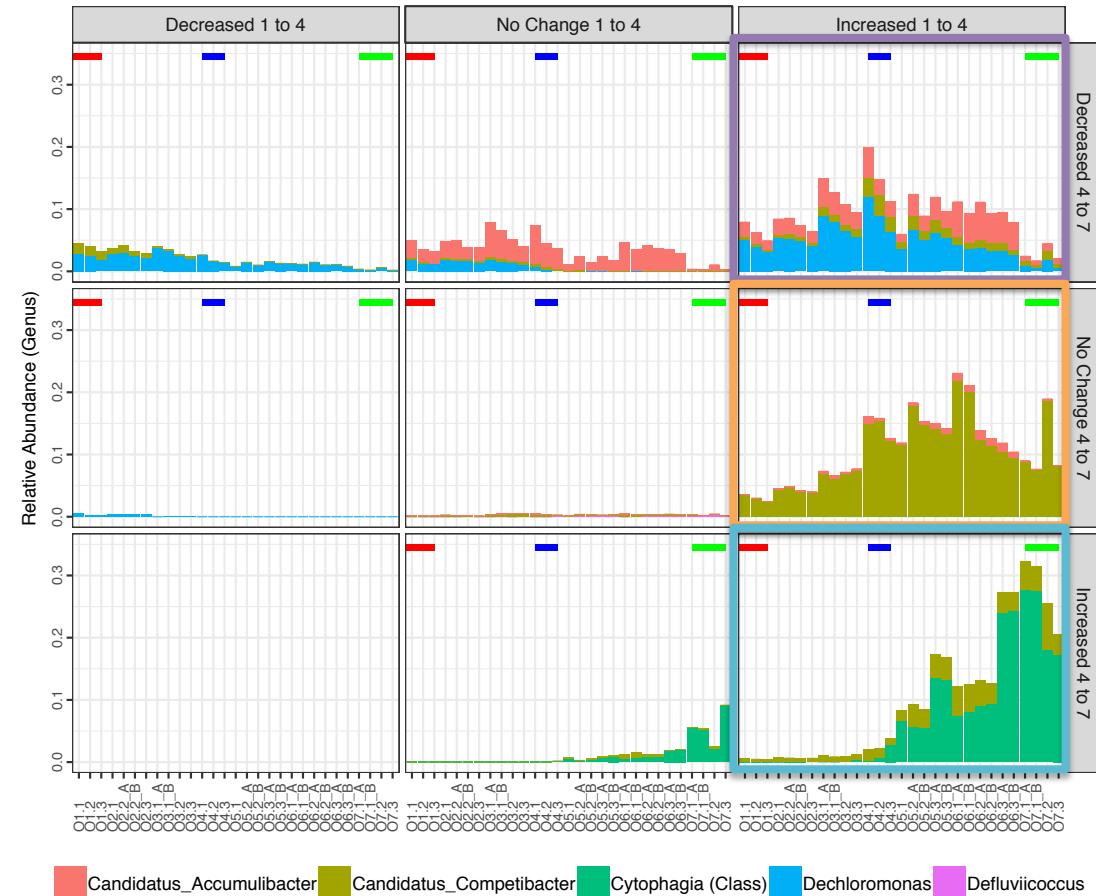
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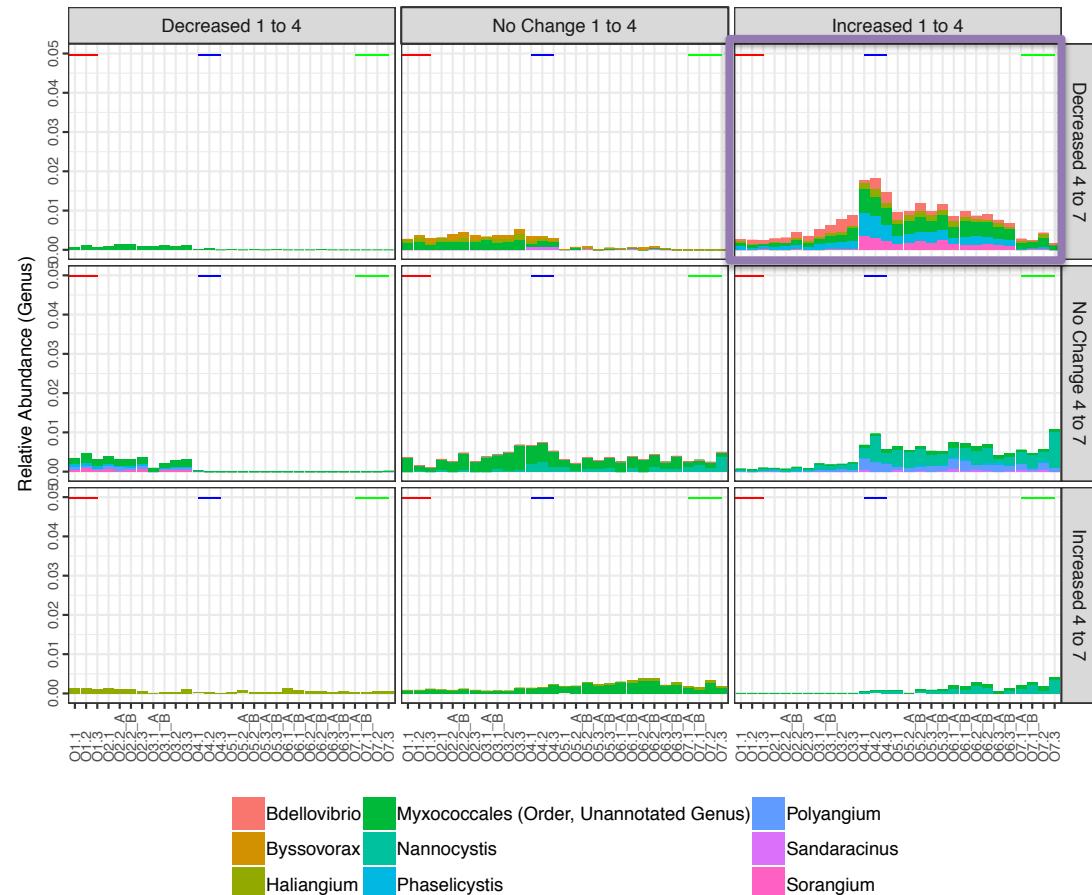
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- Organic C degraders
 - Cytophagales (Class)



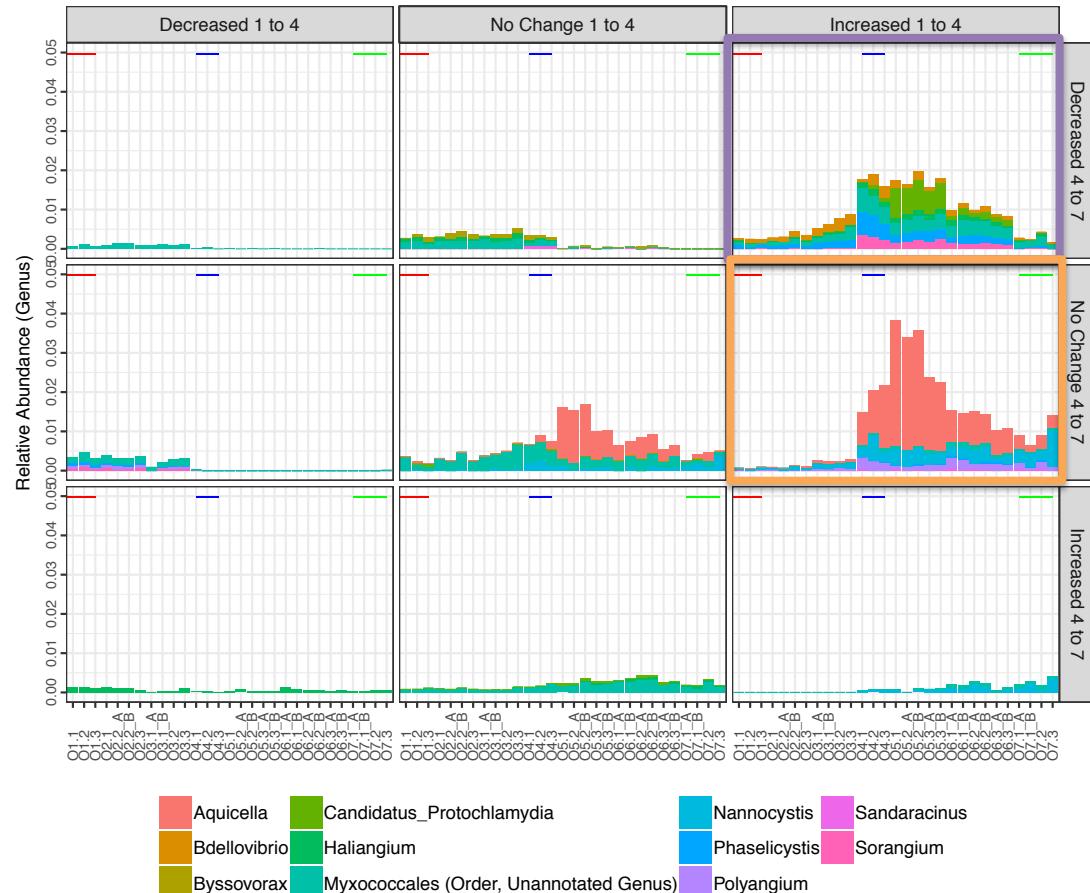
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 - *Difluviicoccus* spp.
- Organic C degraders
 - Cytophagales (Class)
- Predatory bacteria
 - *Bdellovibrio* spp.
 - Myxococcales (Order)



Differential Abundance**

- PAO
 - *Ca. Accumulibacter*
 - *Dechloromonas* spp.
- GAO
 - *Ca. Competibacter*
 - *Difluviicoccus* spp.
- Organic C degraders
 - Cytophagales (Class)
- Predatory bacteria
 - *Bdellovibrio* spp.
 - Myxococcales (Order)
- Protist-associated
 - *Aquicella* spp.
 - *Ca. Protochlamydia*



Discussion & Conclusions

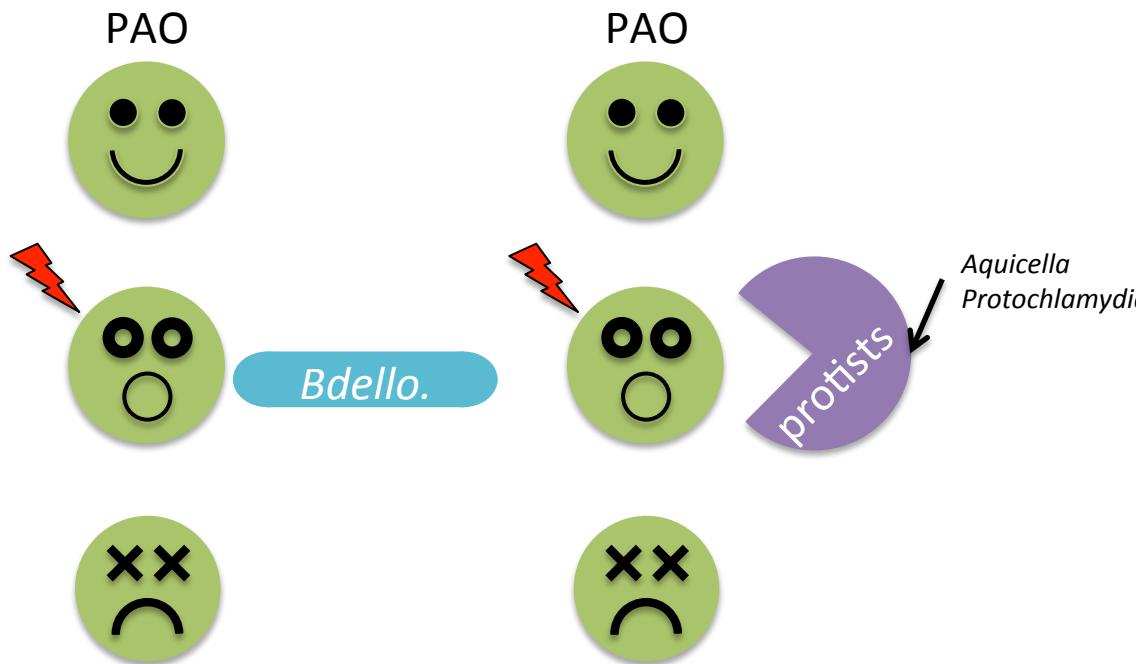
PAO



Bdello.

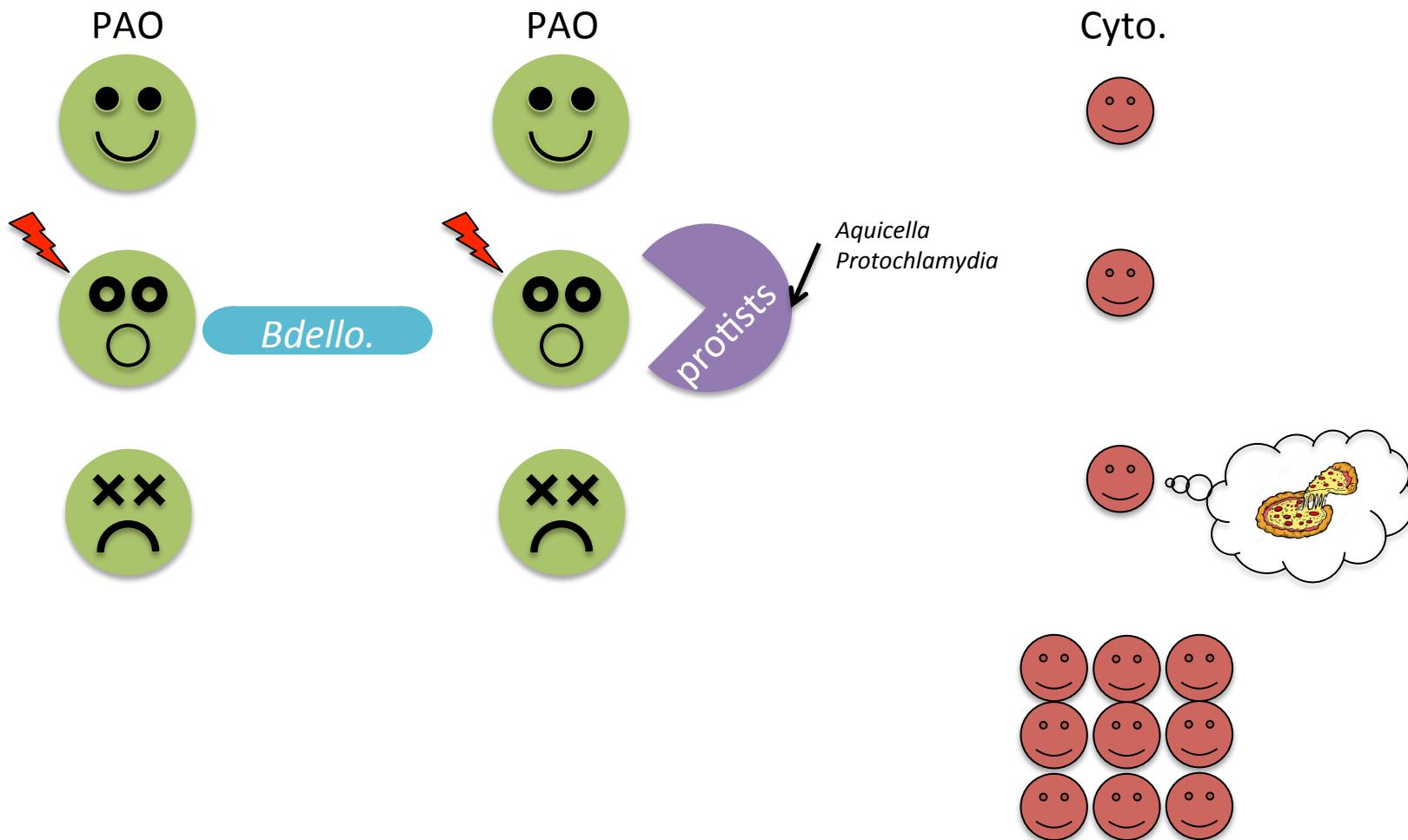


Discussion & Conclusions





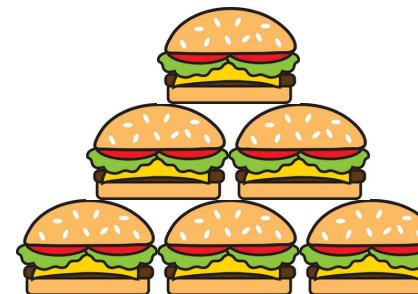
Discussion & Conclusions*





Discussion & Conclusions

PAO



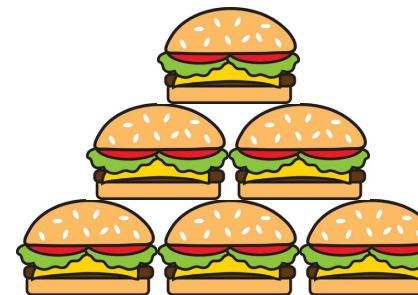
GAO





Discussion & Conclusions

PAO

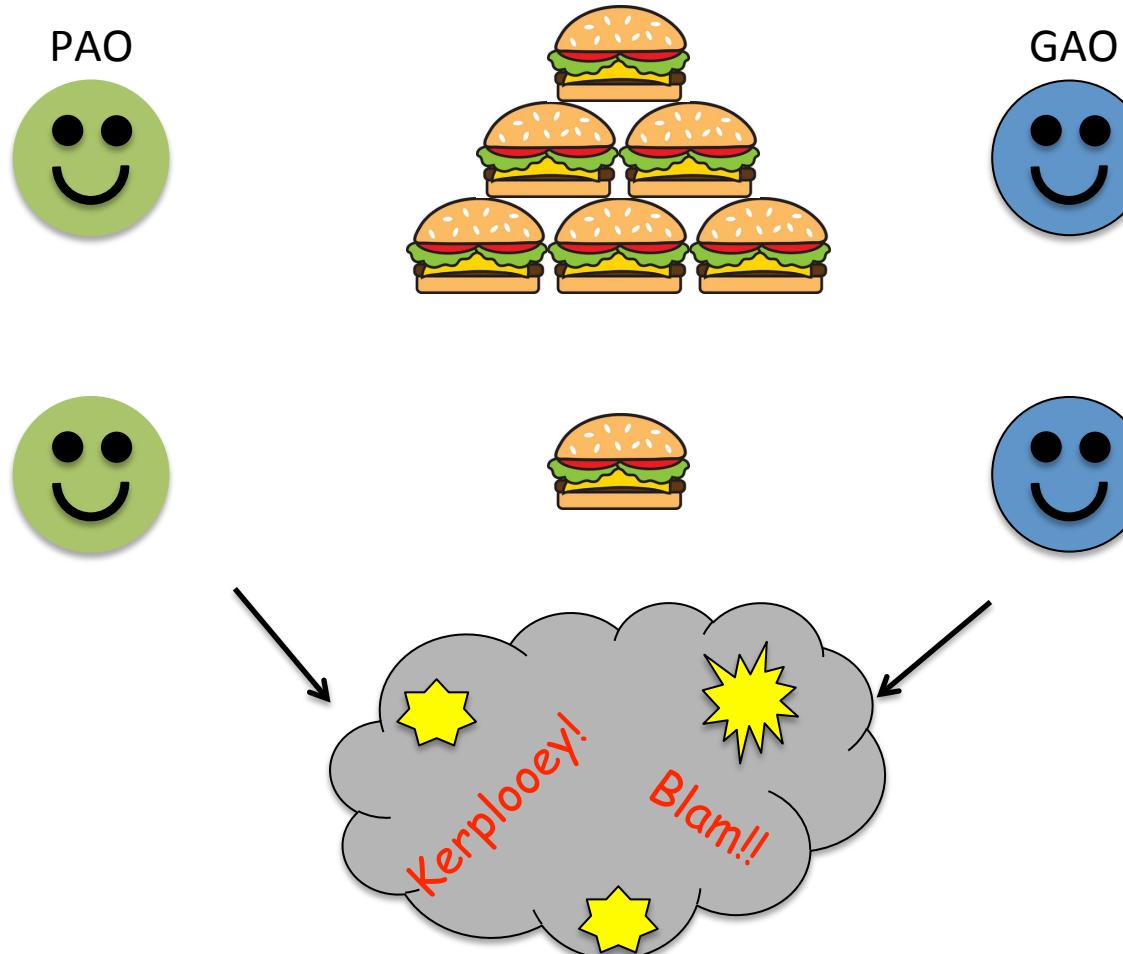


GAO





Discussion & Conclusions**



Future Work*

- Expand differential abundance testing methods
- Metagenomic Sequencing
 - Integrate WGS and Amplicon sequencing results
 - Recover MAGs



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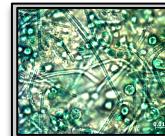
Civil, Architectural, and
Environmental Engineering

College of Engineering

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Sales Laboratory



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- Yaping Nan
 - Yachao Wang
 - Min Cheng
 - Linjiang Yuan



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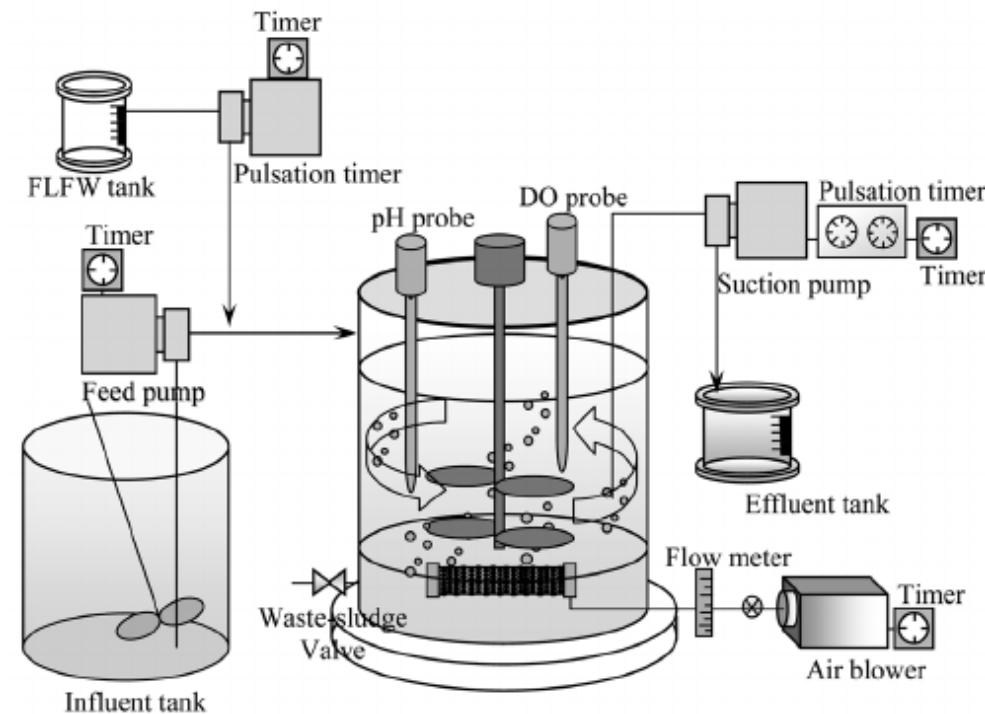
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of
WISCONSIN
MADISON



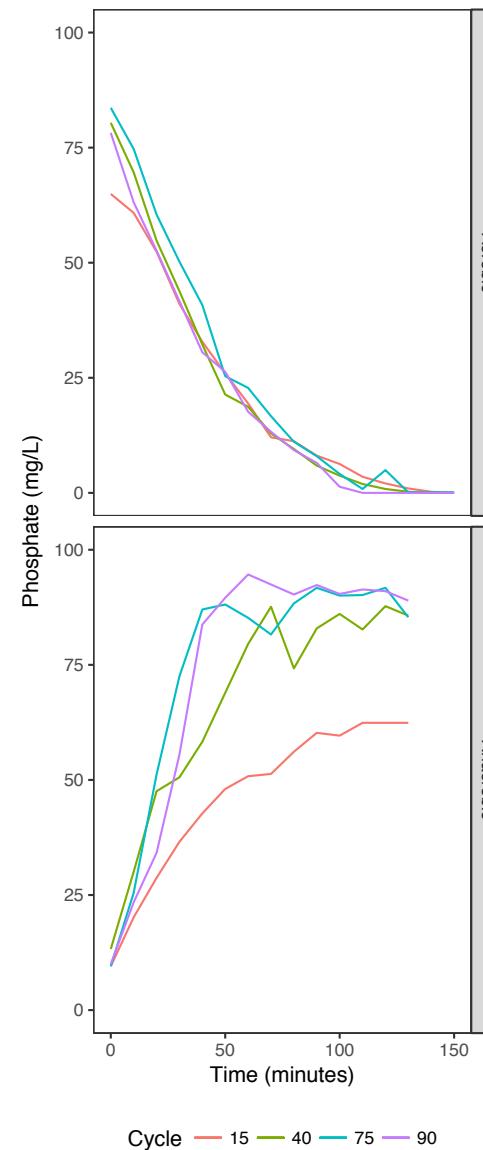
Supplemental

Methods

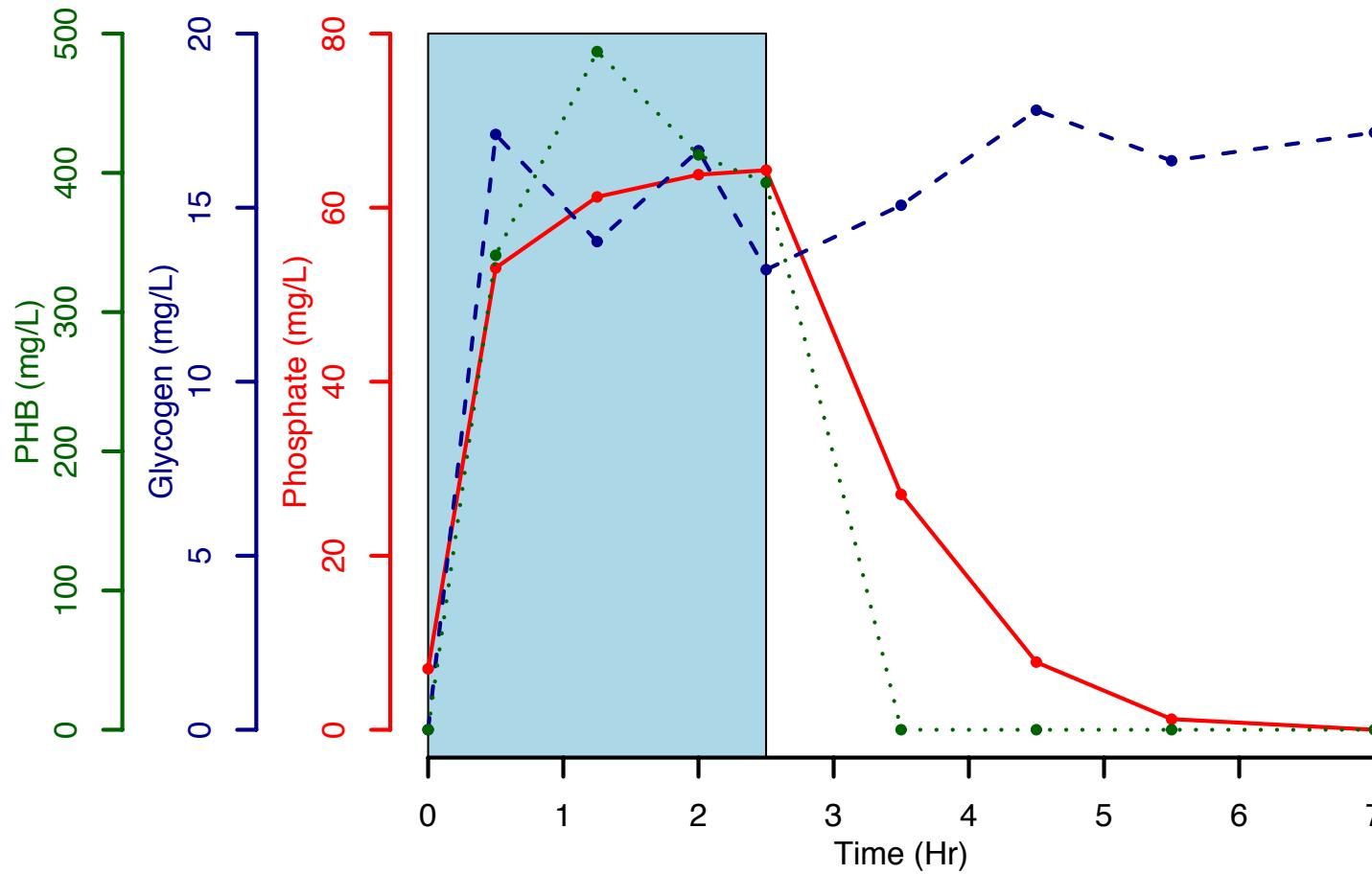
- 2 x SBRs
 - Experiment O
 - 40 days = 120 cycles
 - Experiment S
 - 23 days = 67 cycles
- 8-hour cycles
 - Anaerobic - 2.5 hrs
 - Aerobic – 4.5 hrs
 - Settling – 1 hr
- Synthetic influent
 - C – acetate
 - N – NH_4^+
- Chemical Data
 - PO_4^{3-} , COD, Gly, PHB
- 16S rRNA
 - Experiment O
 - 7 sampling events
 - Experiment S
 - 2 sampling events (beginning/end)



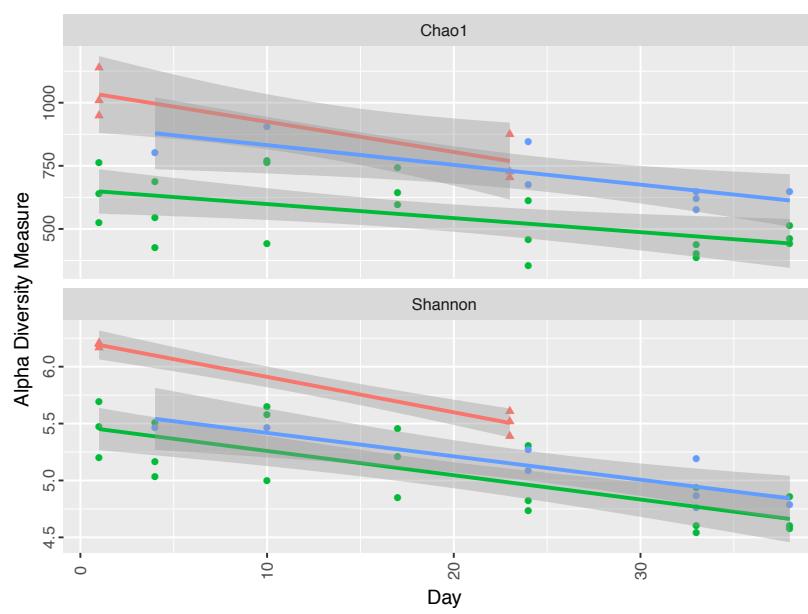
Kinetics



Kinetics



Alpha Div & High-level Abundances



SeqRun

- A
- B
- C

