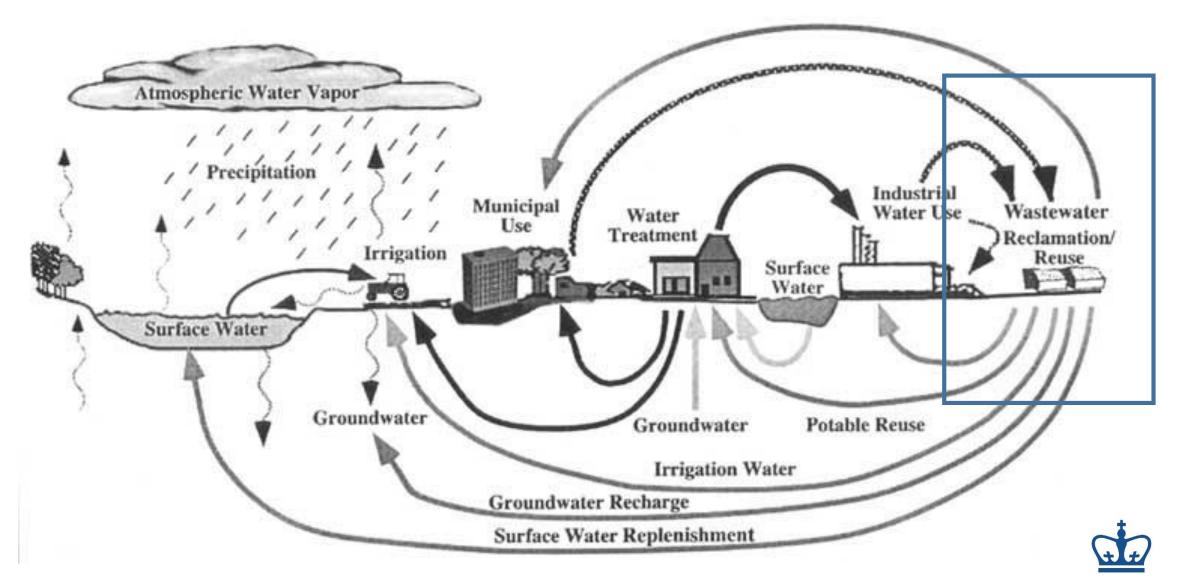
Meta-omics of the Engineered Water Cycle (Nitrogen Cycling in Wastewater Treatment)

> Medini Annavajhala, PhD Student Chandran Laboratory | Columbia University <u>http://www.columbia.edu/~kc2288/</u> Ion World Tour | NYC | September 2015

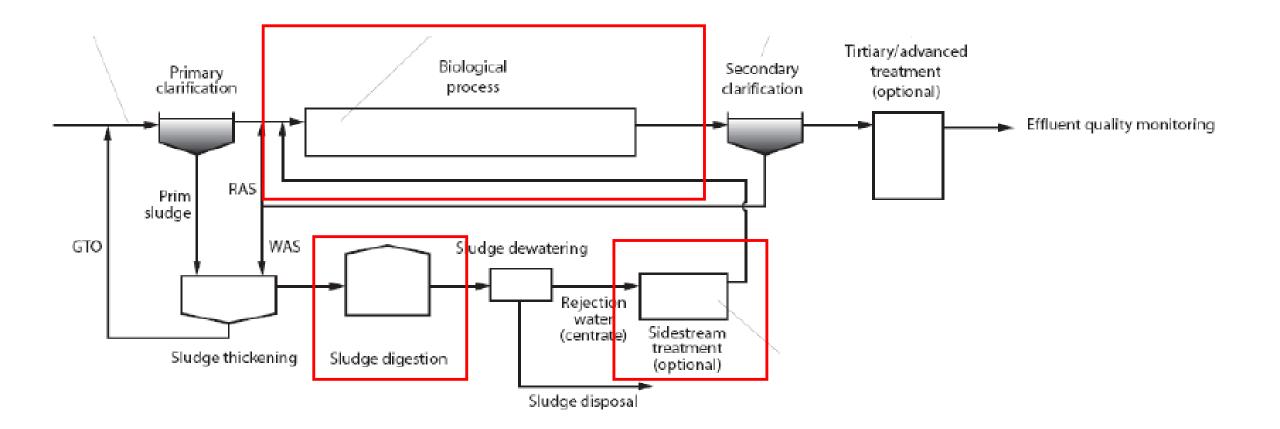


What is the Engineered Water Cycle?



United Nations University - Work in Progress Newsletter - Volume 15, Number 2, 1998

What is the Engineered Water Cycle?



Operation and Control of BNR Facilities, Nutrient Compendium, WERF, 2009

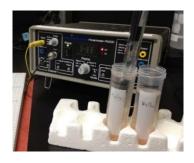


M. Annavajhala, June 2013



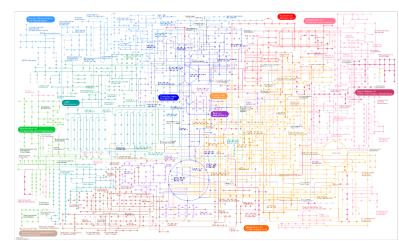
A. C. Brotto









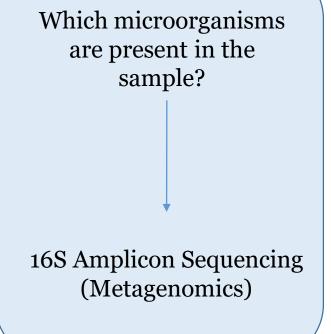


M. Annavajhala and A. C. Brotto, April 2015

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Day 1	Day 7	Day 17	Day 38	[Pseudomonas] Nos

M. Annavajhala and A. C. Brotto, April 2015





Which metabolic & functional pathways are active in the sample?

Shotgun DNA Sequencing (Metagenomics), RNA-Seq (Metatranscriptomics) How are genes being differentially expressed across samples?

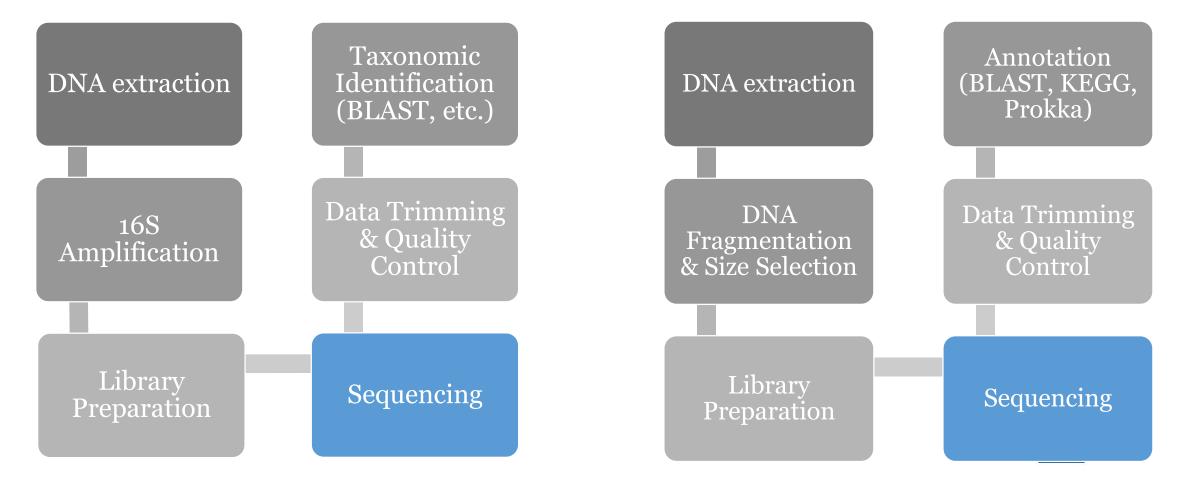
RNA-Seq (Metatranscriptomics)



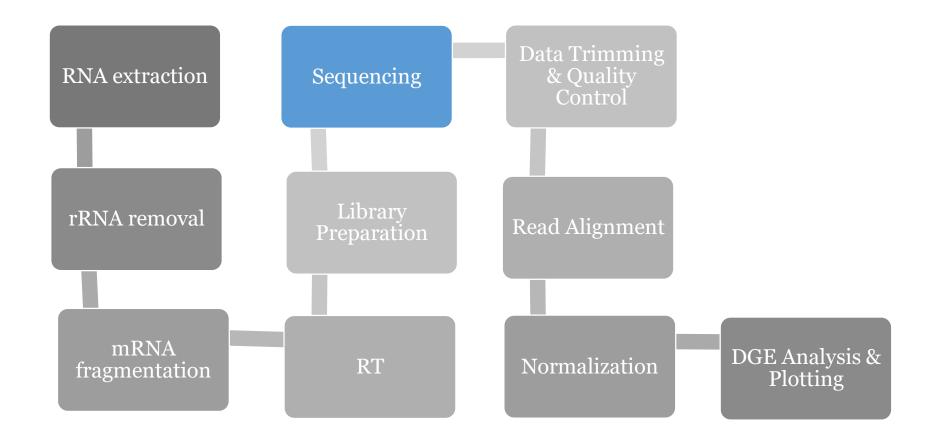
(Meta)genomics

Identifying the Community (16S Amplicon Sequencing)

Identifying Functional Pathways (Shotgun Sequencing)



(Meta)transcriptomics (RNA-Seq)





Library Preparation & Sequencing Kits

16S Amplicon Sequencing

N

In-house developed fusion primers (target: 16S V4 region)

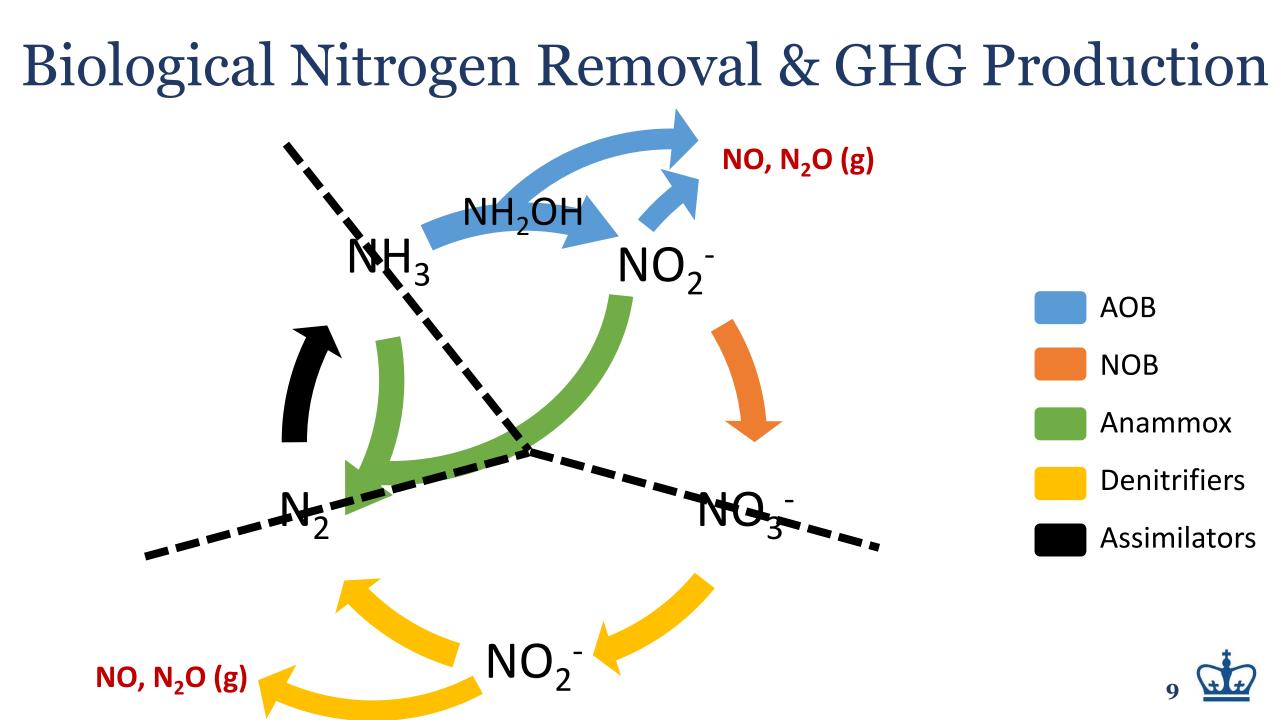
Shotgun	NEB Next Fragmentation Kit		
U	Ion Xpress Plus Fragment		
letagenomics	Library Kit		
	Ion Xpress Barcode Adaptors		

lon OT2 & PGM

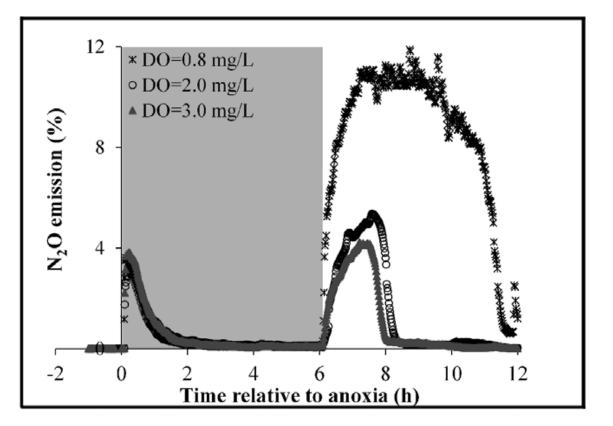
KAPA Library Quantification Kit
Ion Quantitation Kit
Ion PGM 400 OT2 Kit
Ion PGM 400 Sequencing Kit
Ion PGM Hi-Q OT2 Kit
Ion PGM Hi-Q Sequencing Kit
Ion 316v2, 318v2 Semiconductor Chips

RNA-Seq/ Metatranscriptomics Ion RNA-Seq Kit v2 Ion Xpress RNA Barcode Adaptors





Oxygen Cycling: Goals

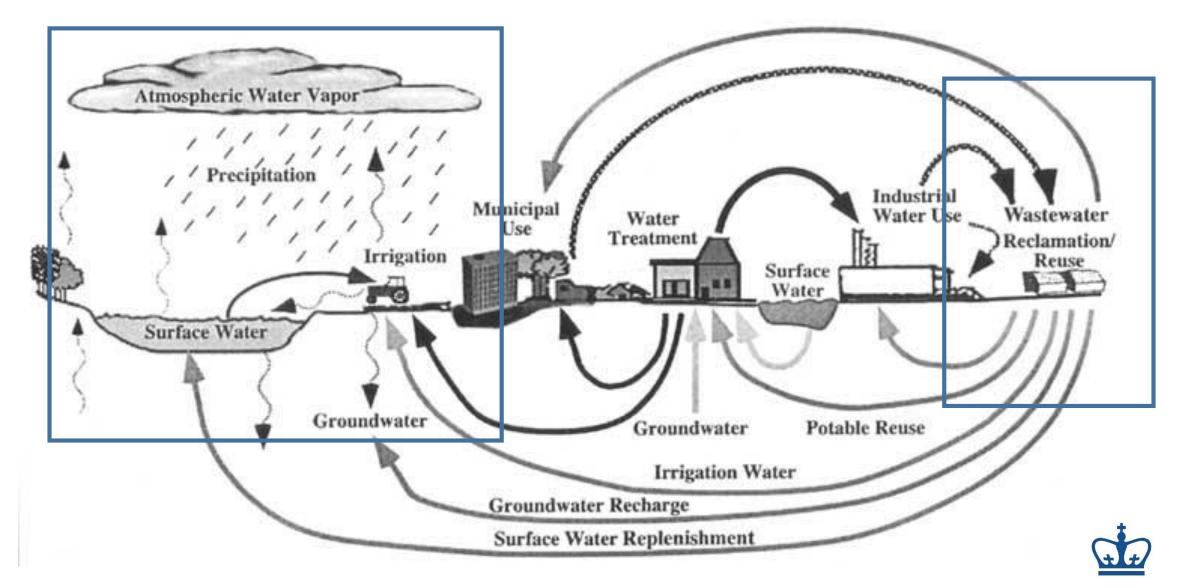


- Identify pathways of NO, N2O production in full nitrification lab-scale reactor under transiently oxic conditions
- Connect chemical & gaseous data to NGS findings
- Find parallels/interactions between N, C, and other cycles
- Track changes in pathways/gene expression over time



Chandran, K (2011). Biochem. Soc. Trans. 39, 1832-1837

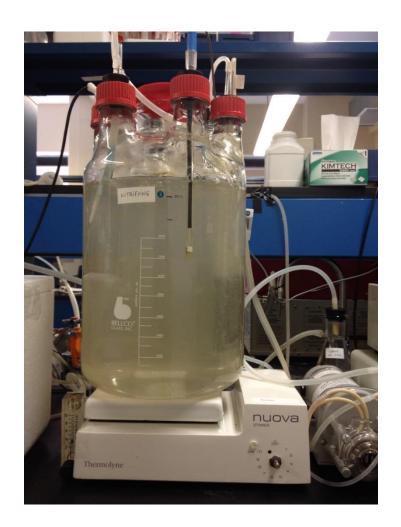
What is the Engineered Water Cycle?



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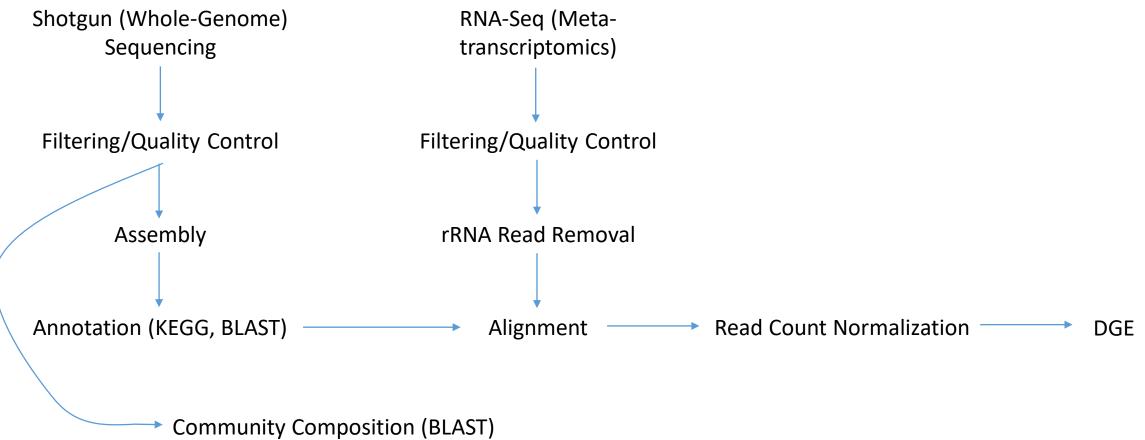
Oxygen Cycling: Experimental Design

- Mimic full nitrification in labscale Sequencing Batch Reactor (SBR)
- Impose transient anoxic-oxic conditions
- Collect chemical and gaseous nitrogen-cycle data
- Collect biomass at specific timepoints for NGS

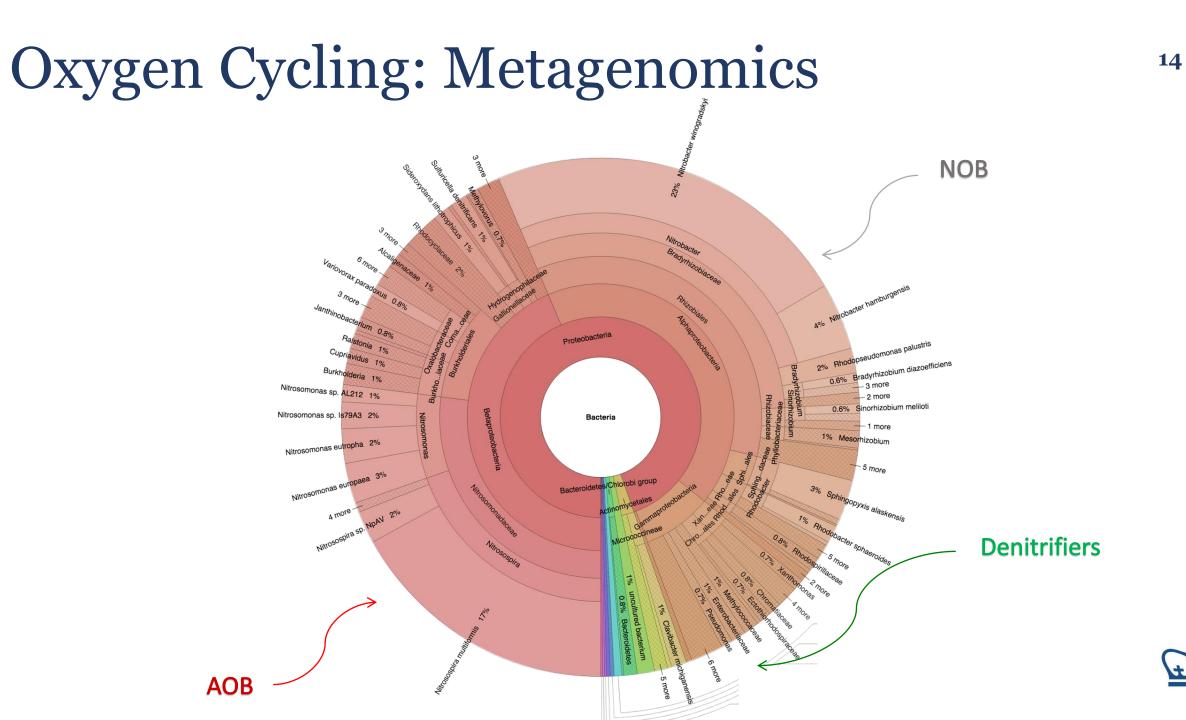


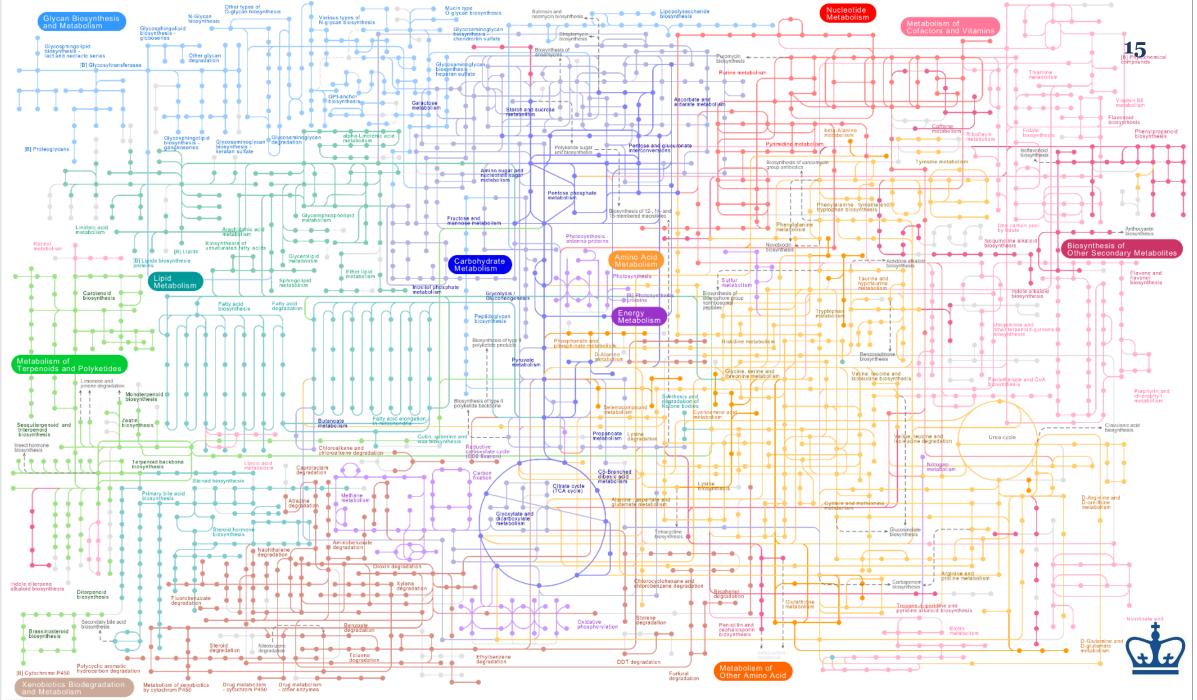


Oxygen Cycling: NGS Workflow



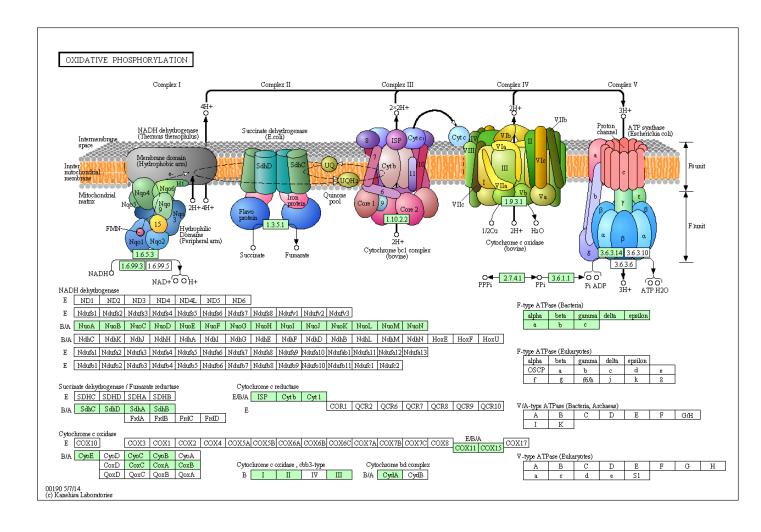


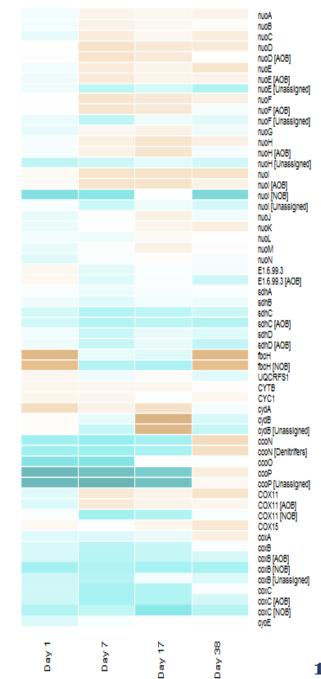




01100-3/11/14 (c) Katelicia Laboratorius

Metatranscriptomics & DGE





Challenges & Lessons Learned

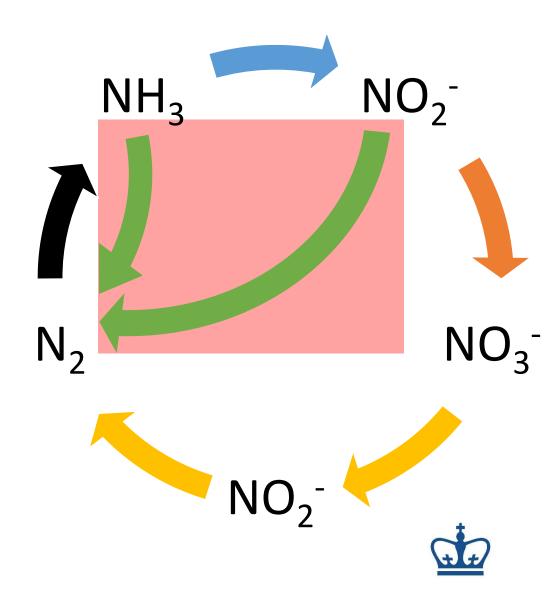
- "Who is present/active" and "What are the system's capabilities/functions" are much easier to answer than "Who is doing what?"
- Extremely well annotated reference sequences (whether already available or created as part of your NGS workflow) are crucial to meaningful RNA-Seq results
- Complex microbial communities make collection (coverage) & interpretation (resolution) of NGS data difficult, as expected
- Meta-omics capabilities are available, but still require troubleshooting on both library preparation and bioinformatics ends
- Experimental design should be driven by end goals; large amounts of data can lead to limits in capacity/experience/analysis



Other NGS Projects & Implications

- Anammox work
- Pure culture studies
- Carbon Cycle & Alternate Endpoints
- Persistent Organic Pollutants

Similar questions can be answered using NGS, but with varied applications and implications



Acknowledgments

- Ariane C. Brotto, PhD Student
- Dr Kartik Chandran, PhD Advisor
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Water Environment Research Foundation Collaboration. Innovation. Results.

