

Efficient Bioremediation of Environmentally Persistent Contaminants with Nanomaterial-Fungus Framework



PI: Susie Dai, Texas A&M University

Co PI: Dr. Joshua Yuan, Texas A&M University

Dr. Greg Lowry, Carnegie Mellon University



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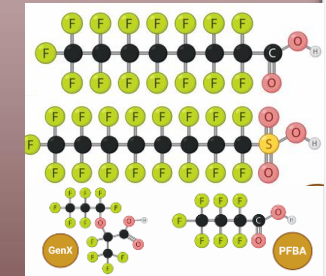
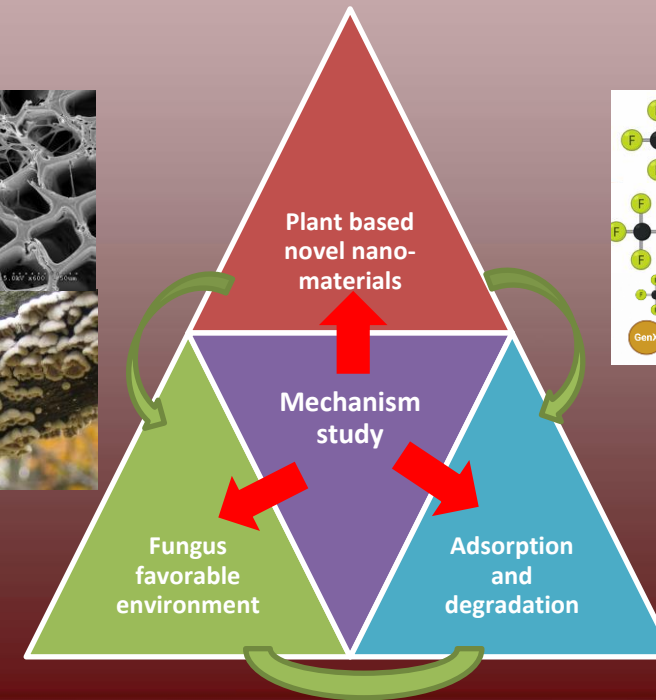
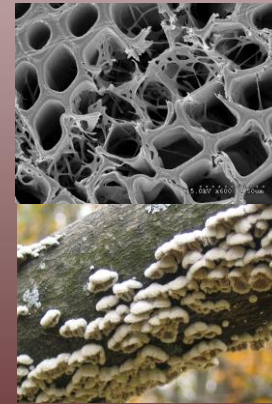
The problem and solution

The Problem:

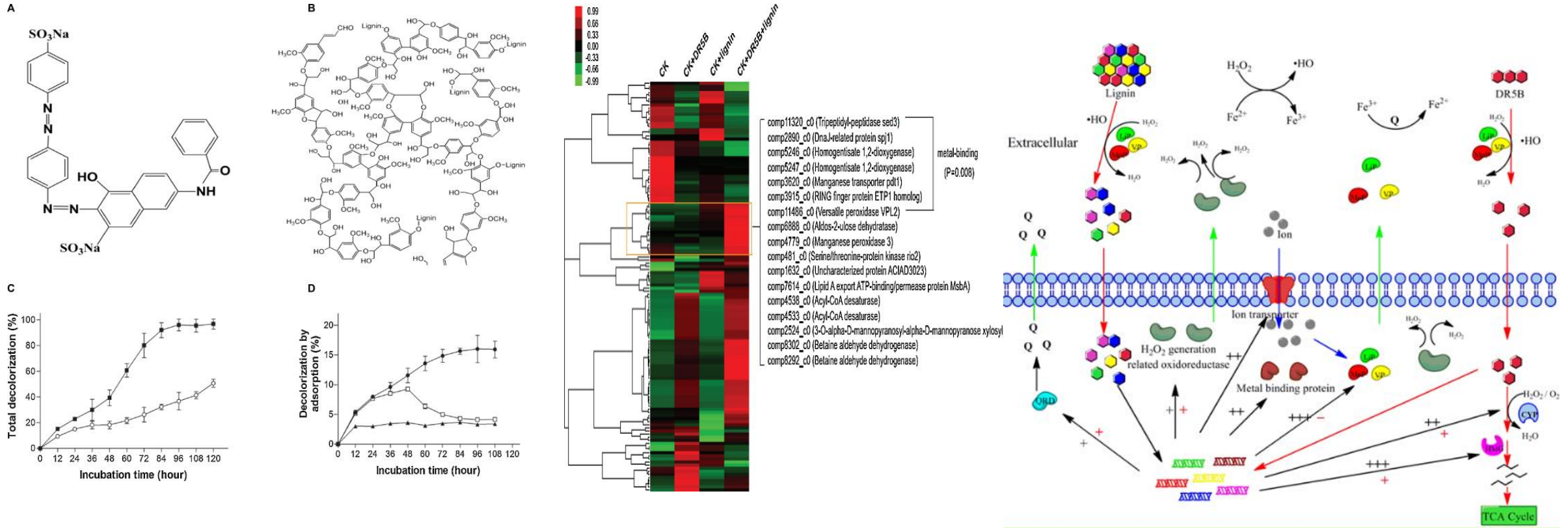
- 1) No sustainable technology to remediate PFAS
- 2) PFAS bioremediation approaches need improvement
 - PFAS bioavailability/toxicity are double-edged swords to the remediation efficiency.
 - Slow growth and PFAS degradation rate
 - Need for continuous nutrient supply for organisms
 - Degradation mechanisms and products are not elucidated

Our Solution:

- ❖ “Efficient Bioremediation of Environmentally Persistent Contaminants with Nanomaterial-Fungus Framework”
- ❖ Integrating material science/engineering/fungal biology/informatics to address the challenges

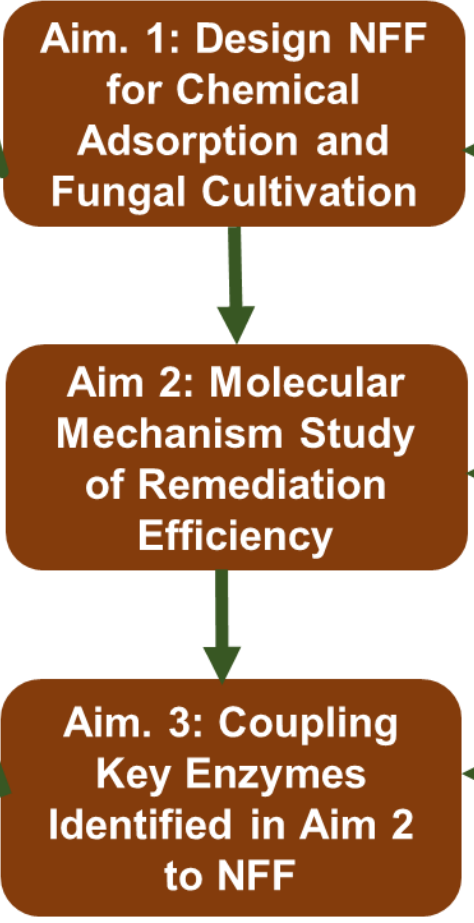


Rational



The fungal degradation of organic dye was significantly enhanced when lignin, the natural fungal substrate was added into the culture media

Technical Approach



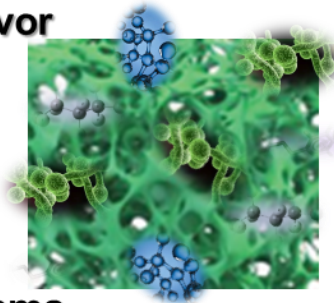
Specific Aims




How material modification can improve chemical adsorption and favor fungal growth?

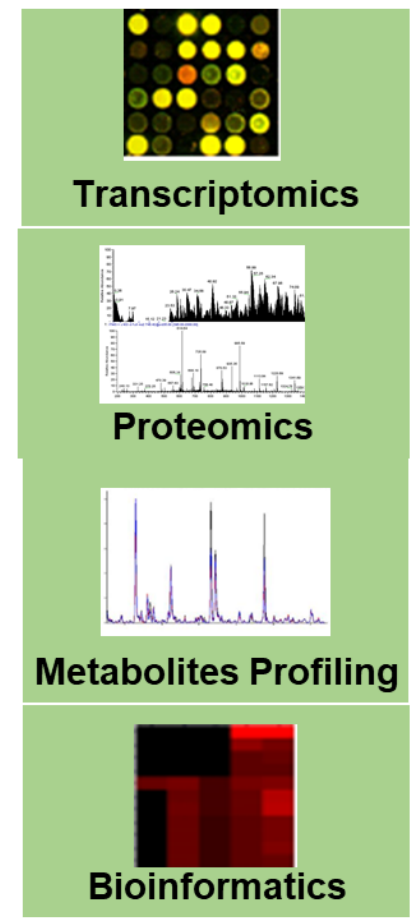
What are the molecular and systems mechanisms of NFF efficiency?

How key genes and enzymes can further improve remediation efficiency?

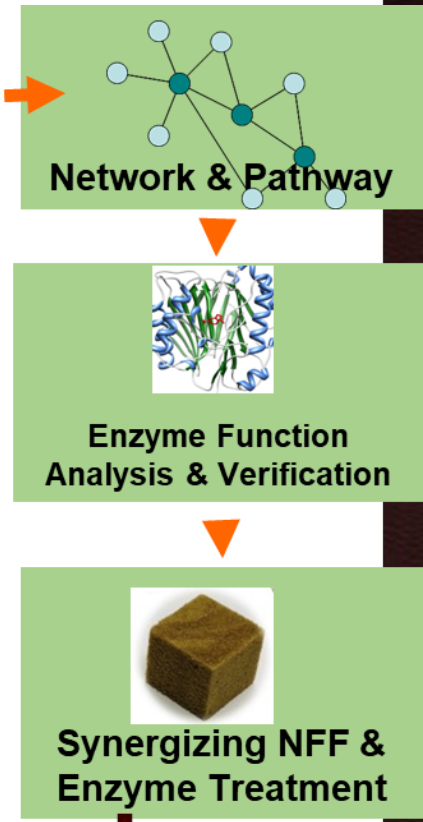
Biological Questions



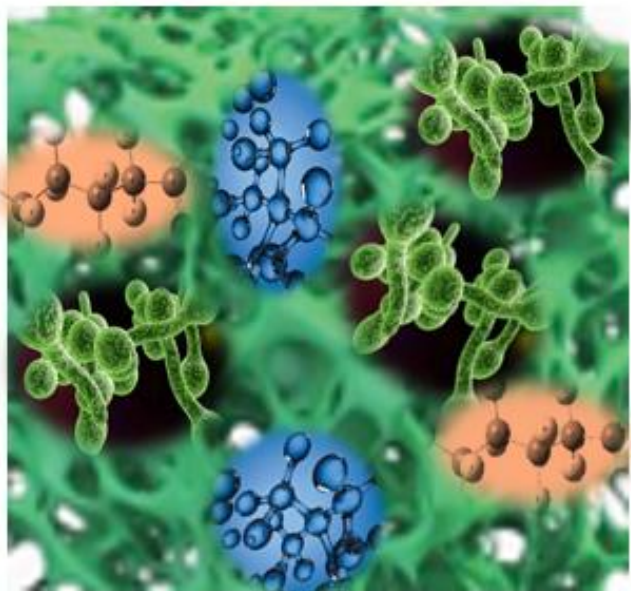
-  Composite
-  Fungus
-  Chemicals



Systems Biology Approach



Aim 1



PFOA



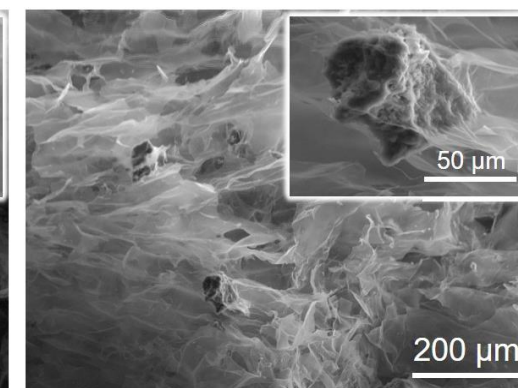
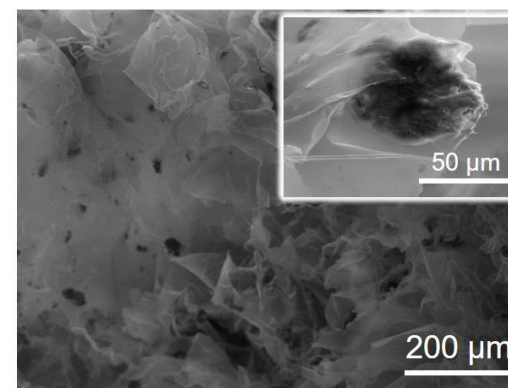
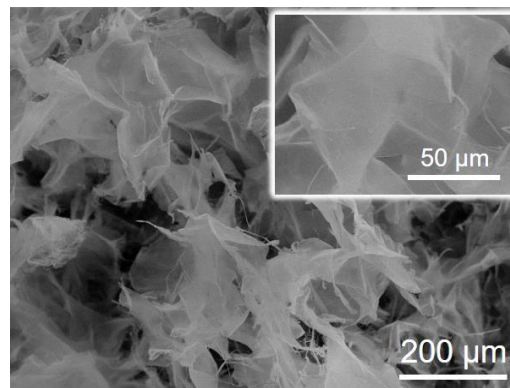
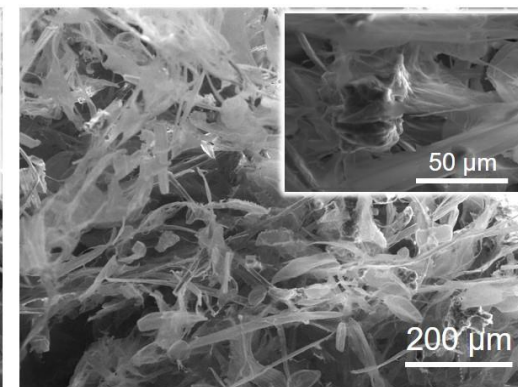
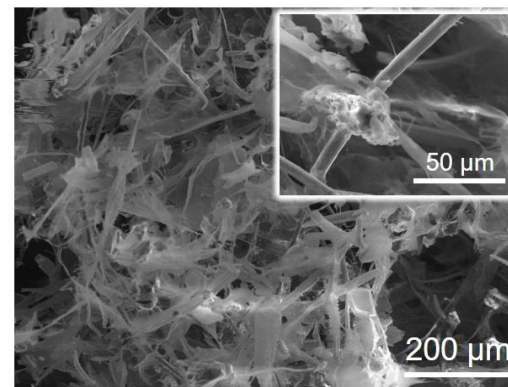
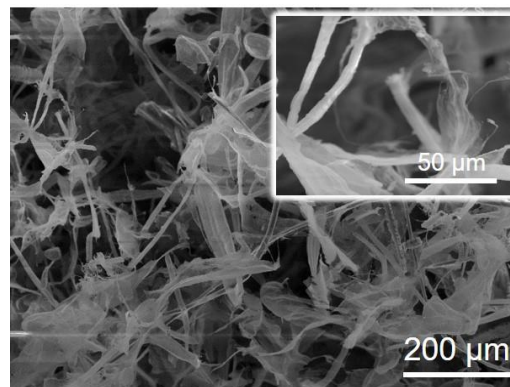
I. lacteus



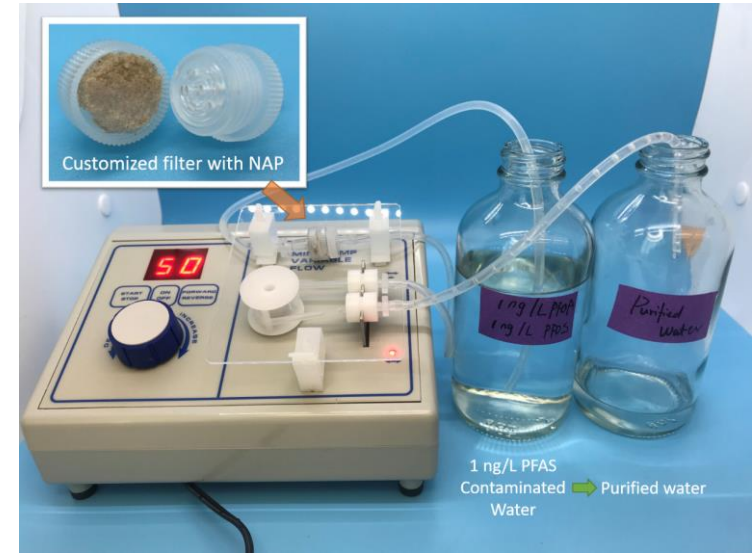
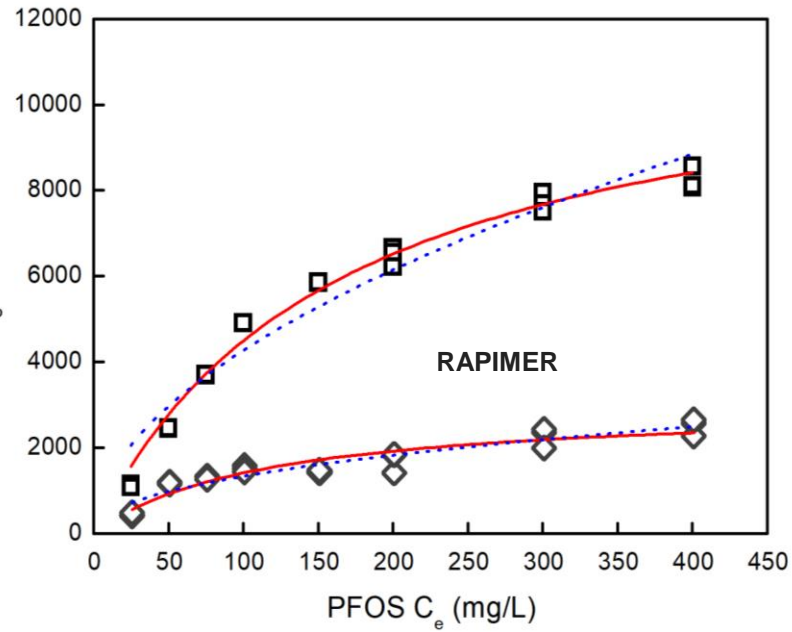
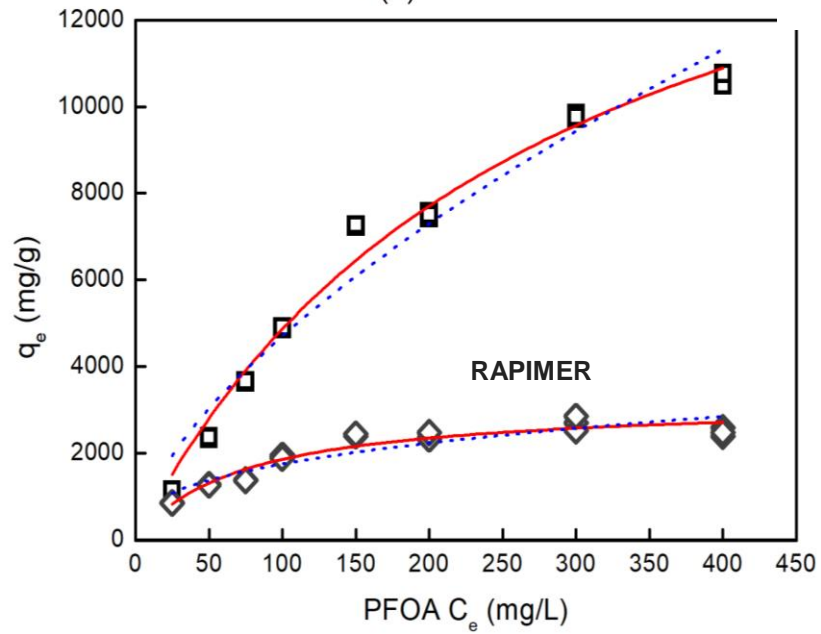
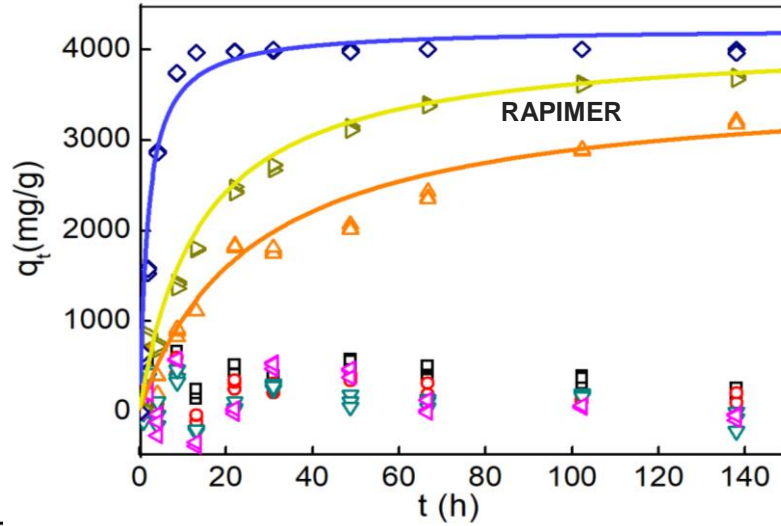
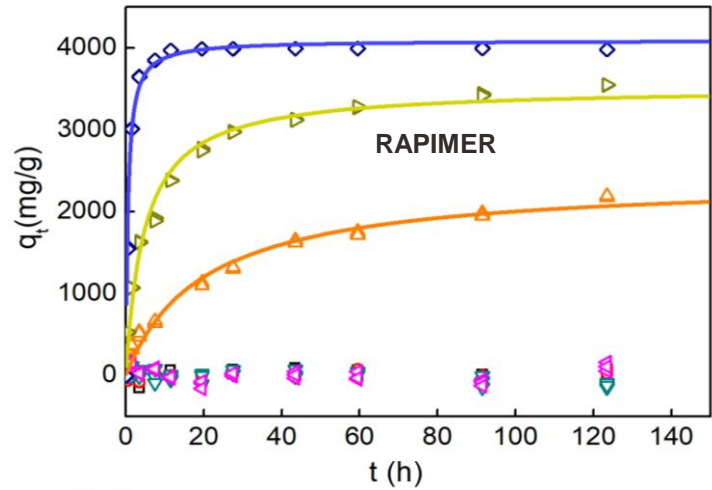
PFOS

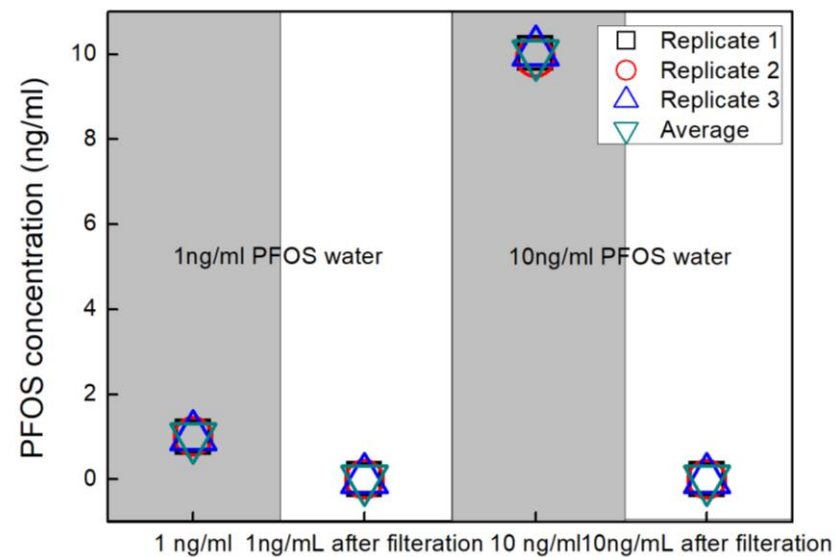
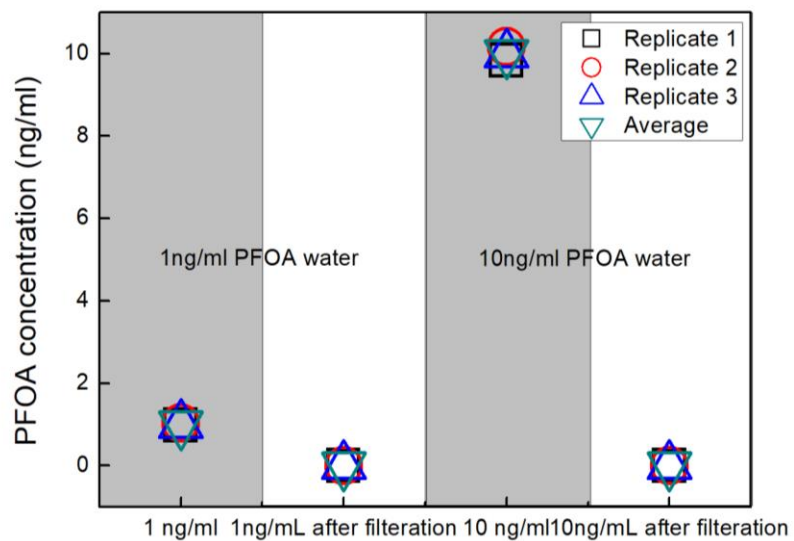
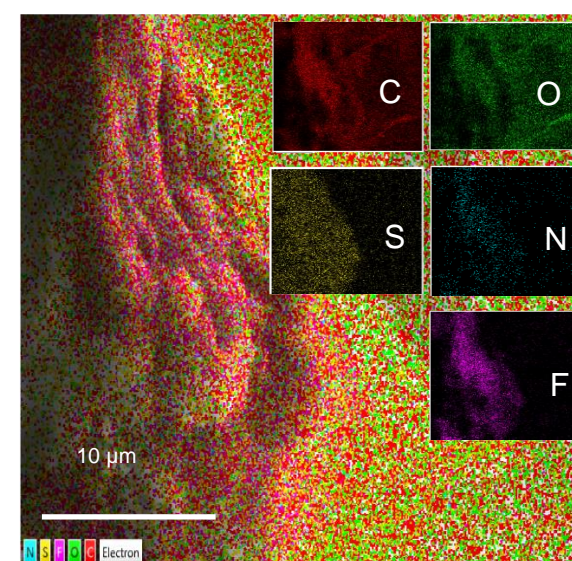
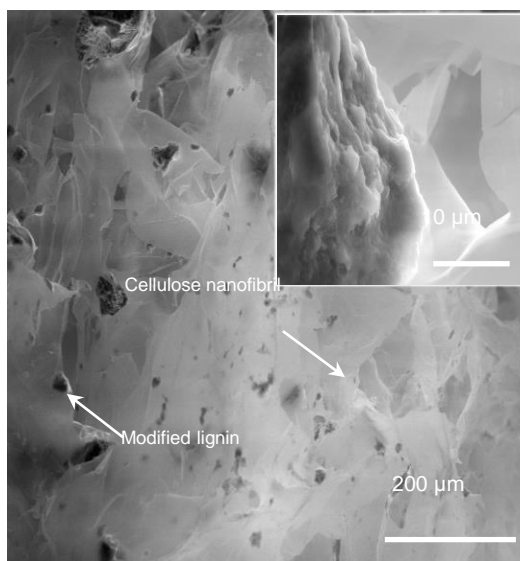
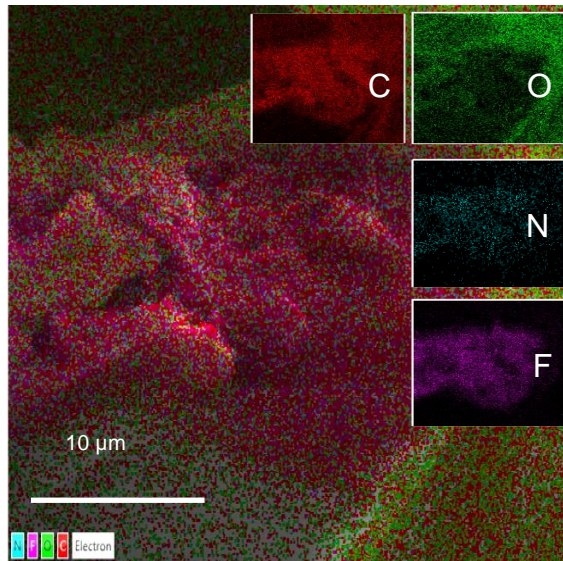
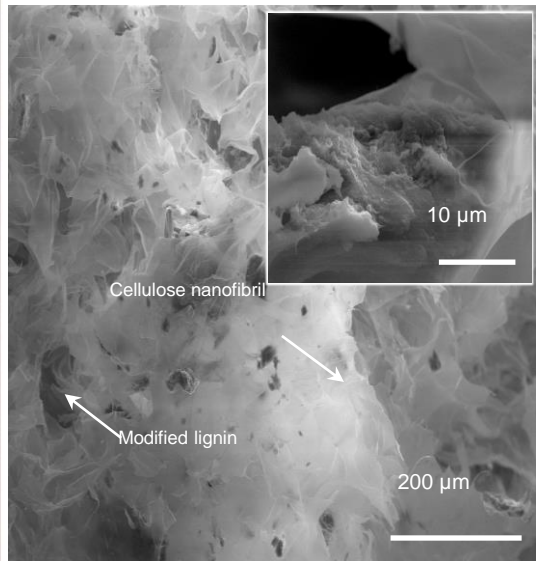


RAPIMER

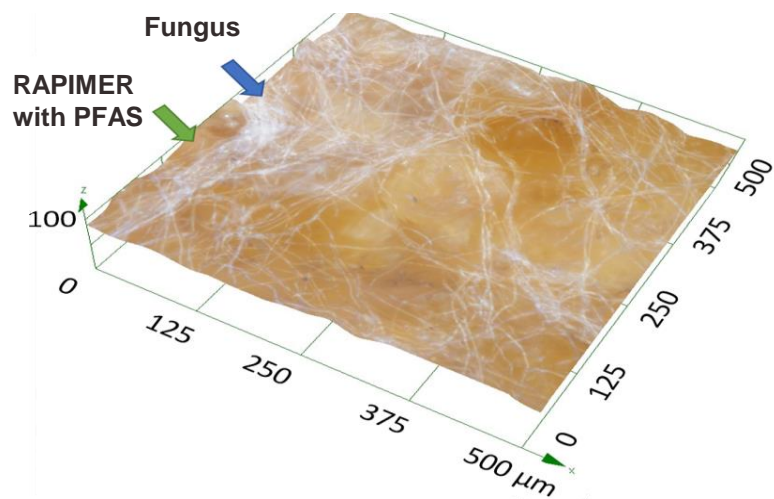
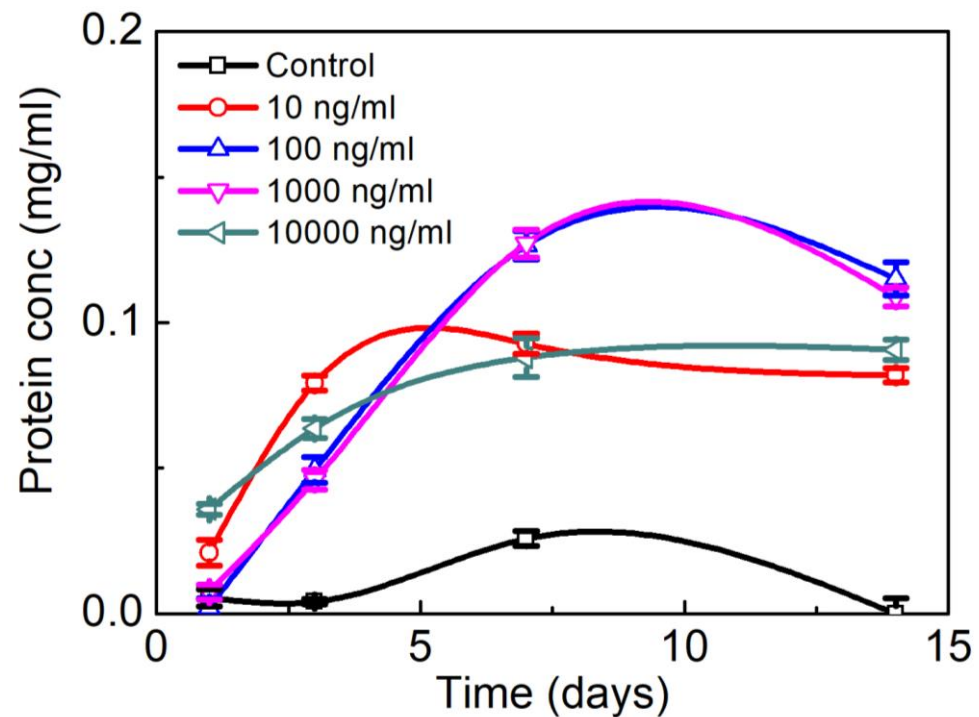
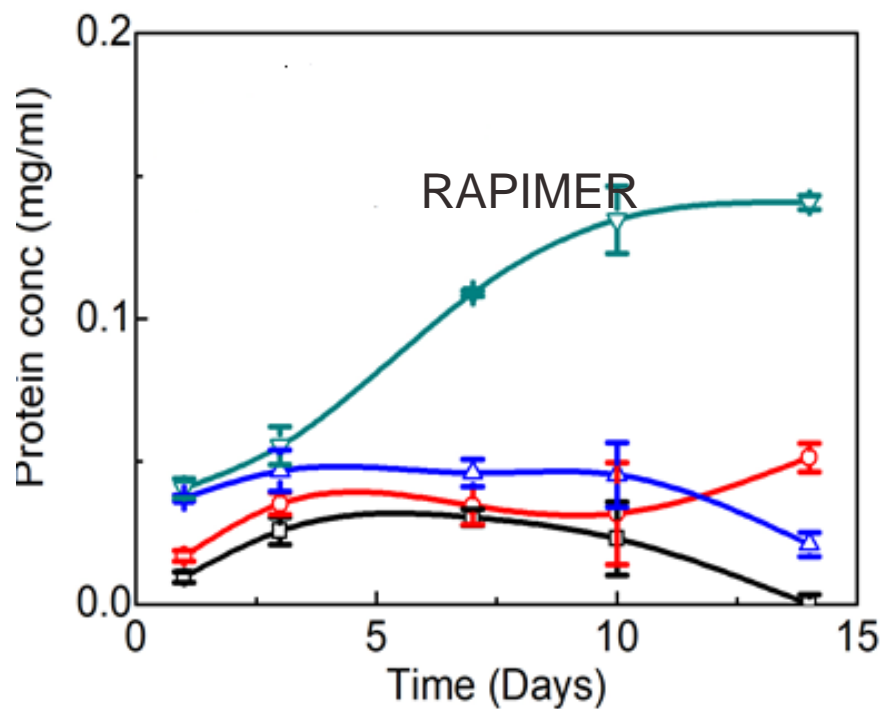


Renewable Artificial Plant for In-Situ
Microbial Environmental Remediation
(RAPIMER)



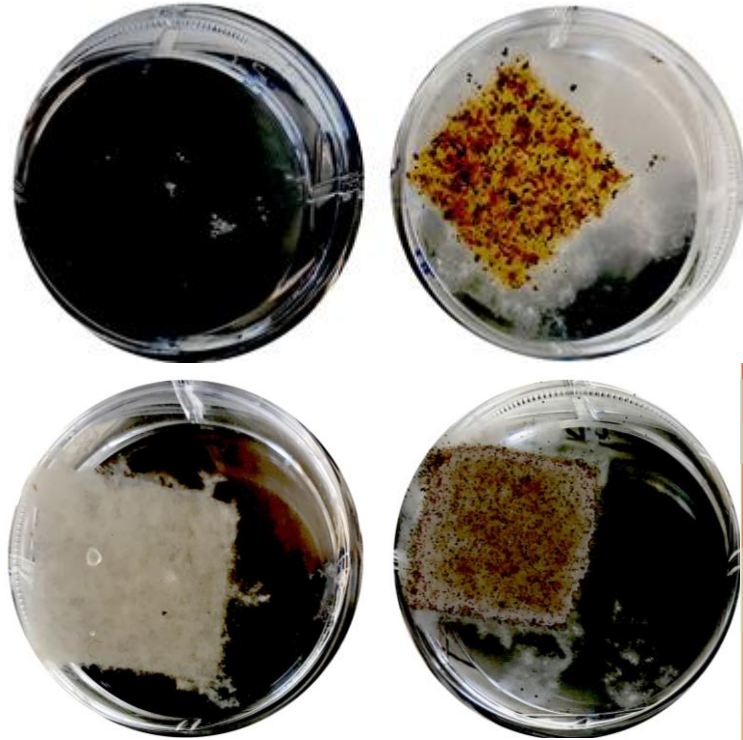


Aim 2



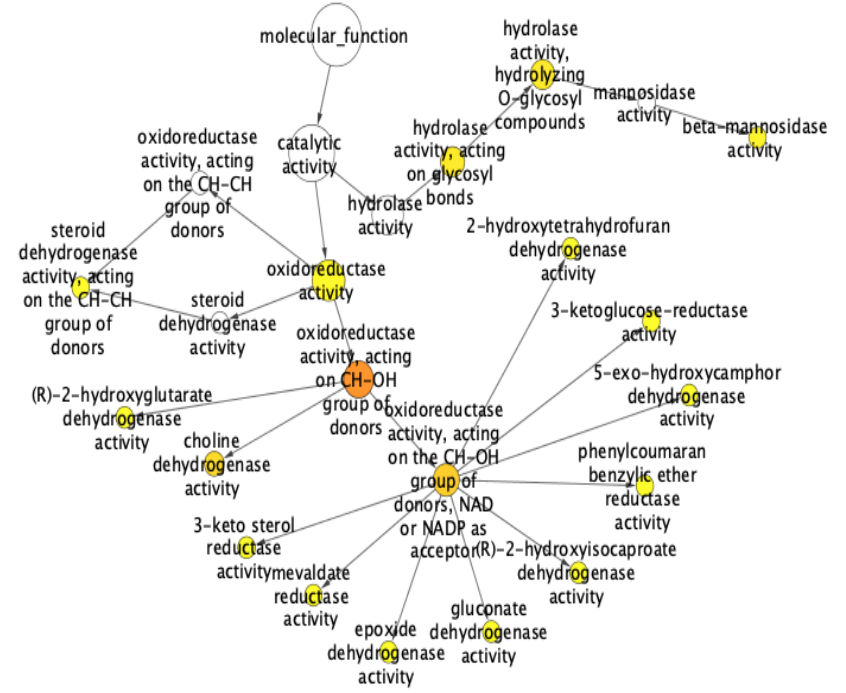
- RAPIMER sustains fungal growth without external carbon source
- RAPIMER can adsorb large amounts of PFAS
- Non-targeted PFAS analysis can help elucidate degradation products

RAPIMER



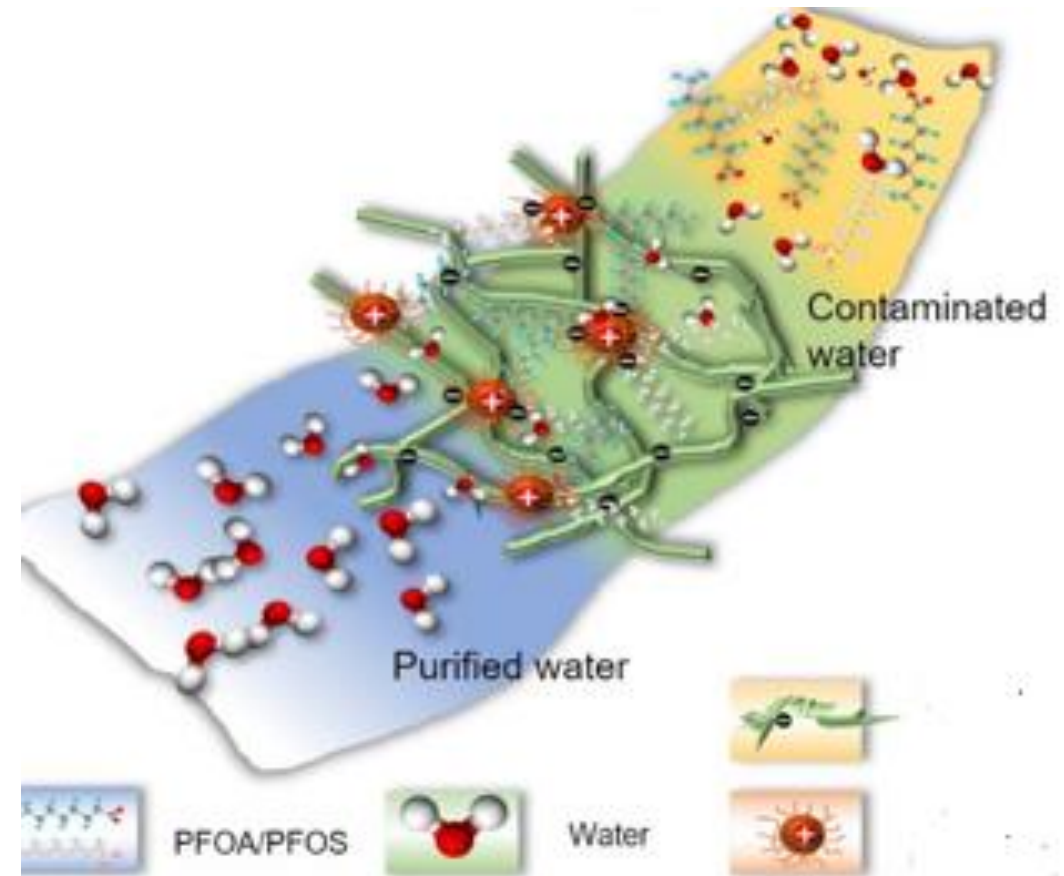
RAPIMER sustains the fungal growth on the PDA plate

- 5.52 Cellulose 1,4-beta-cellobiosidase
- 4.71 Mannan endo-1,4-beta-mannosidase
- 5.60 Hexosyltransferases
- 4.41 GMC oxidoreductase family
- 3.98 Glycoside hydrolase, family 5
- 4.29 Metallopeptidase
- 2.38 Thioredoxin
- 3.41 GMC oxidoreductase family
- 4.47 Endo-1,4-beta-xylanase
- 3.36 Acylglycerone-phosphate reductase
- 2.99 Cellulose-binding domain, fungal
- 4.07 Flavonol reductase/cinnamoyl-CoA reductase
- 4.11 Voltage-gated shaker-like K+ channel, subunit beta
- 2.95 Reductases with broad range of substrate specificity
- 3.09 Function unknown
- 3.24 Glycerol 2-dehydrogenase (NADP+)
- 3.23 Beta-glucosidase
- 3.43 Glycoside hydrolase, family 5
- 3.05 With NAD(+) or NADP(+) as acceptor
- 2.95 60s ribosomal protein L23
- 2.72 Function unknown
- 2.55 Voltage-gated shaker-like K+ channel, subunit beta/KCNAB
- 2.14 Voltage-gated shaker-like K+ channel, subunit beta/KCNAB
- 2.50 Cytochrome P450 CYP4/CYP19/CYP26 subfamilies
- 2.57 Peptidyl-prolyl cis-trans isomerase



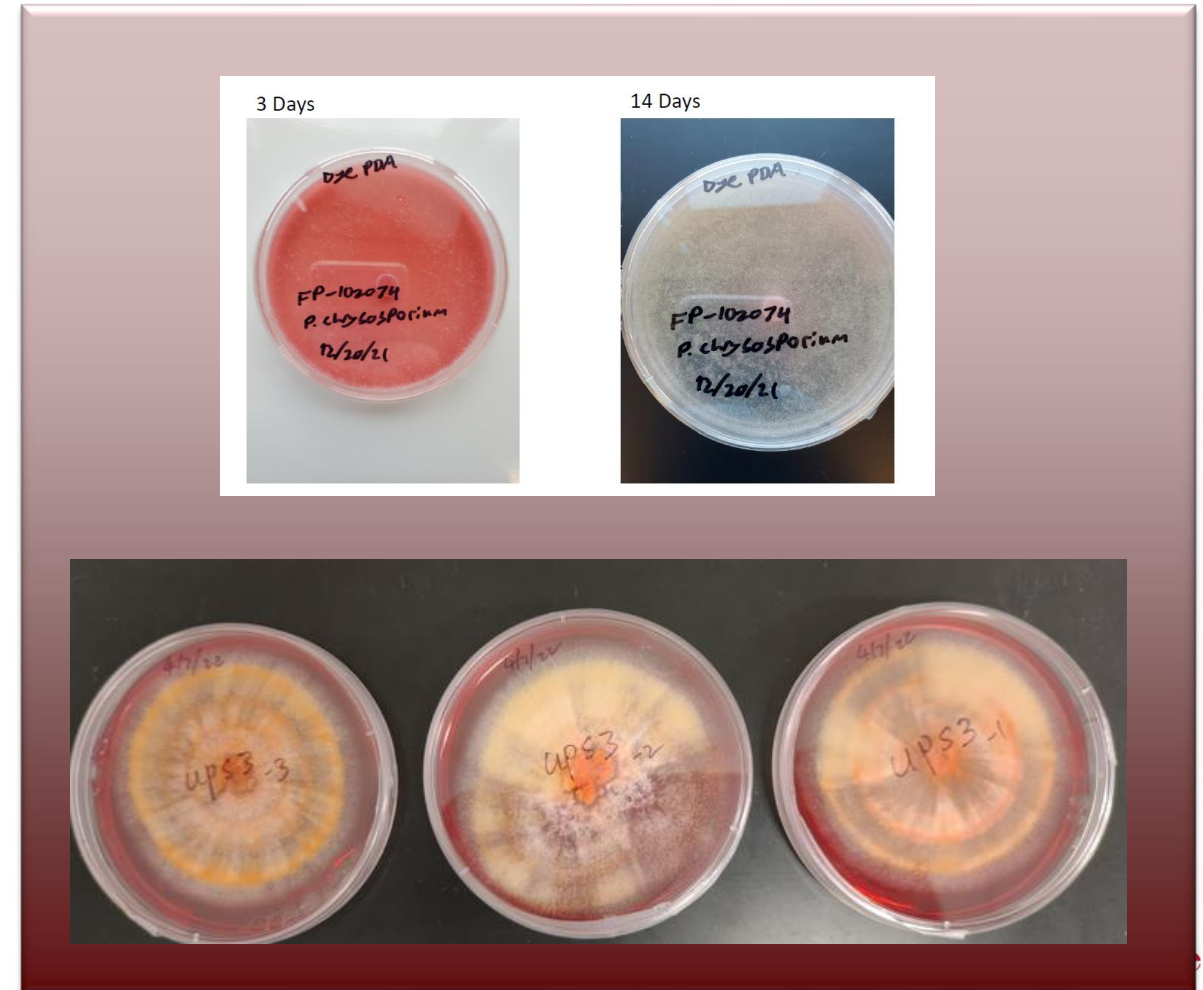
Summary

- The prototype of NFF-RAPIMER has been developed
- Highly efficient sorbent for PFAS and organic dye
- Sustains fungal growth as the carbon source and enables PFSA biotransformation
- The fungal *I. lacteus* strain degrades RAPIMER and PFAS, as revealed by the proteomics and metabolite analysis



Future work

- Screen a large fungus library and identify PFAS tolerant fungi species
- Further optimize RAPIMER for larger scale applications and more matrices
- Non-targeted analysis for PFAS degradation products
- Proteomics for important genes and pathways



Acknowledgement

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