

# Measuring the microbial biodiversity by single-cell analysis

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## Hot topic...

### Functional Ecology



*Functional Ecology* 2015, **29**, 1341–1349

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### **Bacterial diversity amplifies nutrient-based plant–soil feedbacks**

Shifts in microbial diversity through land use intensity as drivers of carbon mineralization in soil



Vincent Tardy<sup>a</sup>, Aymé Spor<sup>a</sup>, Olivier Mathieu<sup>c</sup>, Jean Lévêque<sup>c</sup>, Sébastien Terrat<sup>b</sup>, Pierre Plassart<sup>a</sup>, Tiffanie Regnier<sup>a</sup>, Richard D. Bardgett<sup>d</sup>, Wim H. van der Putten<sup>e,f</sup>, Pier Paolo Roggero<sup>g,h</sup>, Giovanna Seddaiu<sup>h</sup>, Simonetta Bagella<sup>h,i</sup>, Philippe Lemanceau<sup>a</sup>, Lionel Ranjard<sup>a,b</sup>, Pierre-Alain Maron<sup>a,b,\*</sup>

### **Testing biodiversity-ecosystem functioning relationship in the world's largest grassland: overview of the IMGRE project**

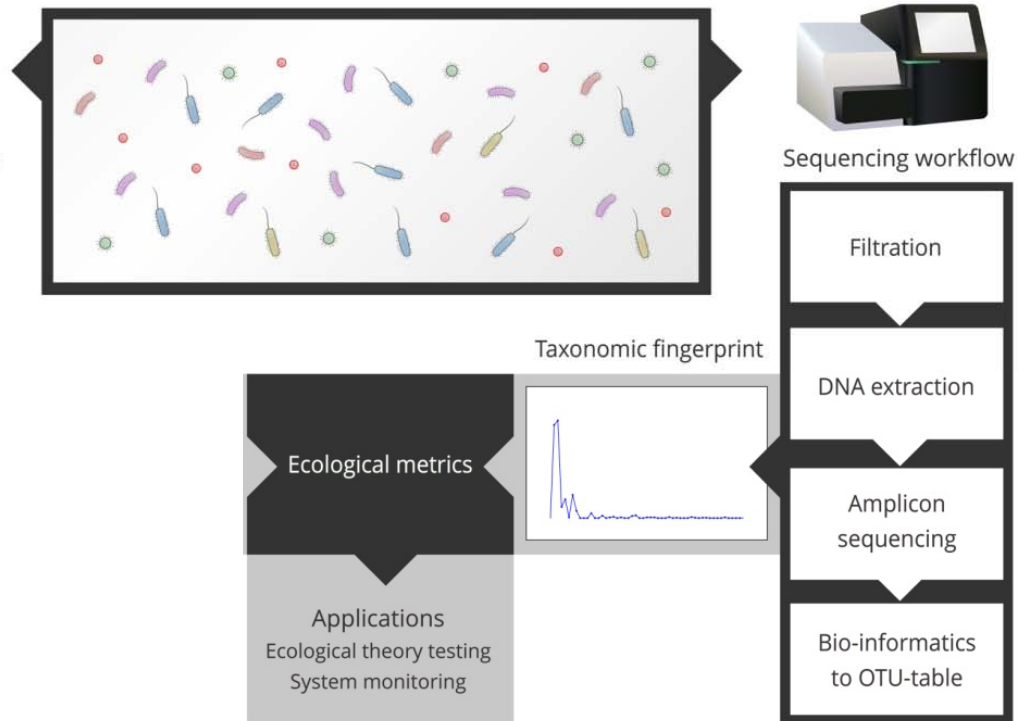
Jianguo Wu · Shahid Naeem · James Elser · Yongfei Bai · Jianhui Huang · Le Kang · Qingmin Pan · Qibing Wang · Shuguang Hao · Xingguo Han

Marine biodiversity and ecosystem function relationships: The potential for practical monitoring applications

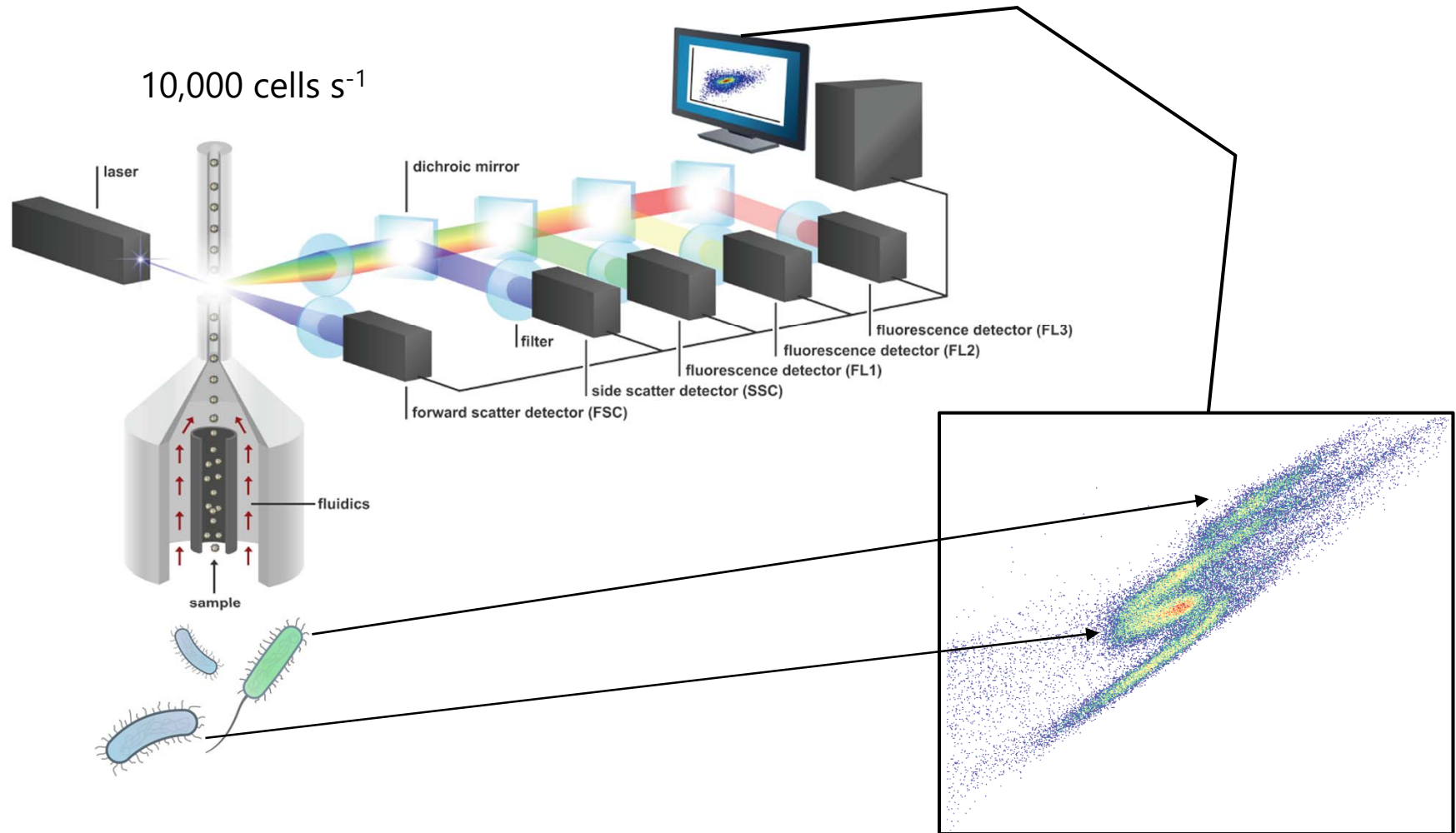
James Asa Strong<sup>a,\*</sup>, Eider Andonegi<sup>b</sup>, Kemal Can Bizsel<sup>c</sup>, Roberto Danovaro<sup>d</sup>, Mike Elliott<sup>a</sup>, Anita Franco<sup>a</sup>, Esther Garces<sup>e</sup>, Sally Little<sup>a</sup>, Krysia Mazik<sup>a</sup>, Snejana Moncheva<sup>f</sup>, Nadia Papadopoulou<sup>g</sup>, Joana Patrício<sup>h</sup>, Ana M. Queirós<sup>i</sup>, Chris Smith<sup>g</sup>, Kremena Stefanova<sup>f</sup>, Oihana Solaun<sup>b</sup>

# Assessing biodiversity

## *The microbial community*

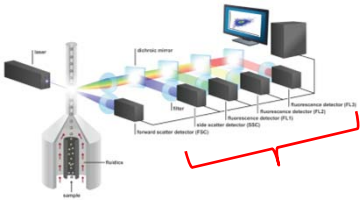


# Flow Cytometry

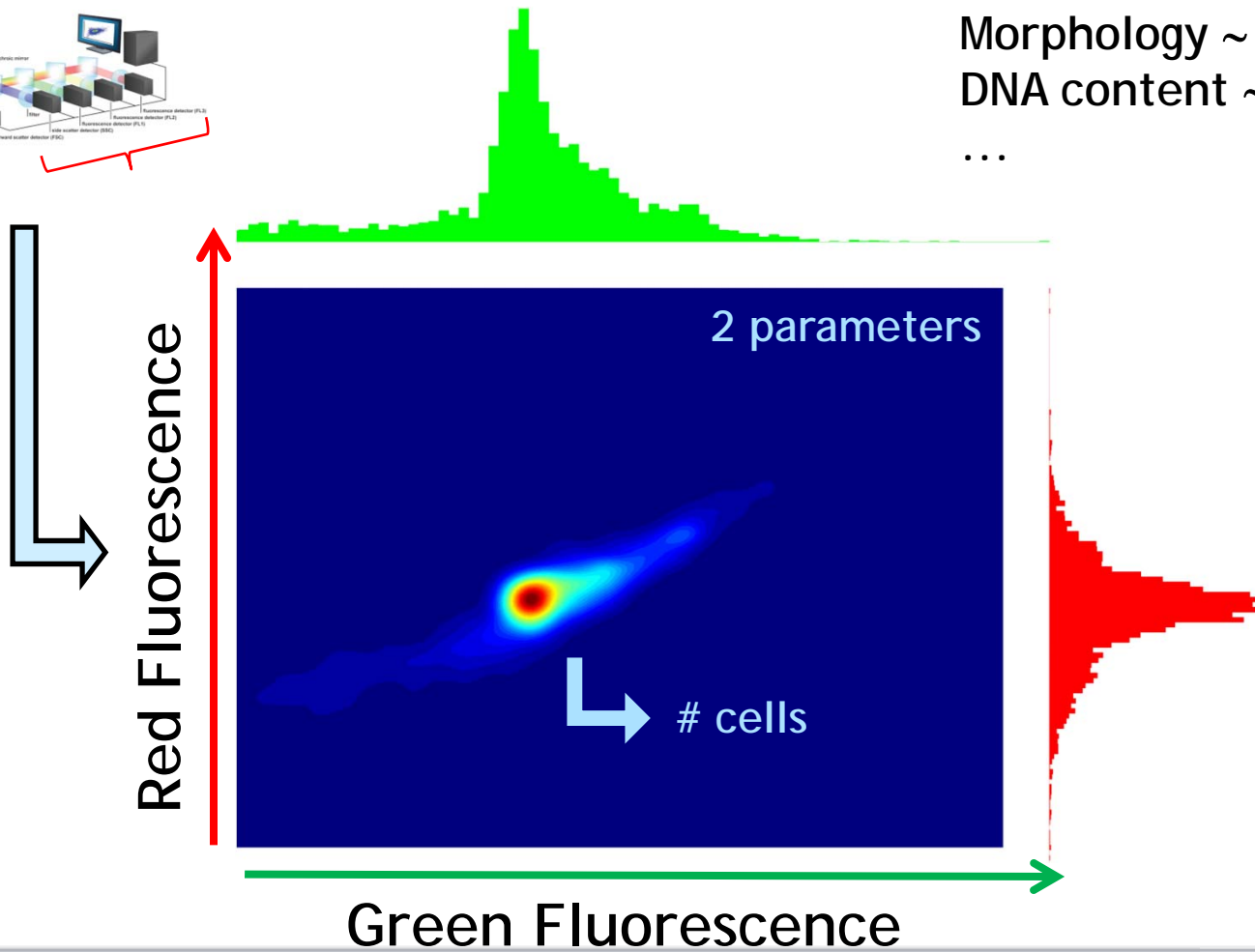


# Phenotypic markers

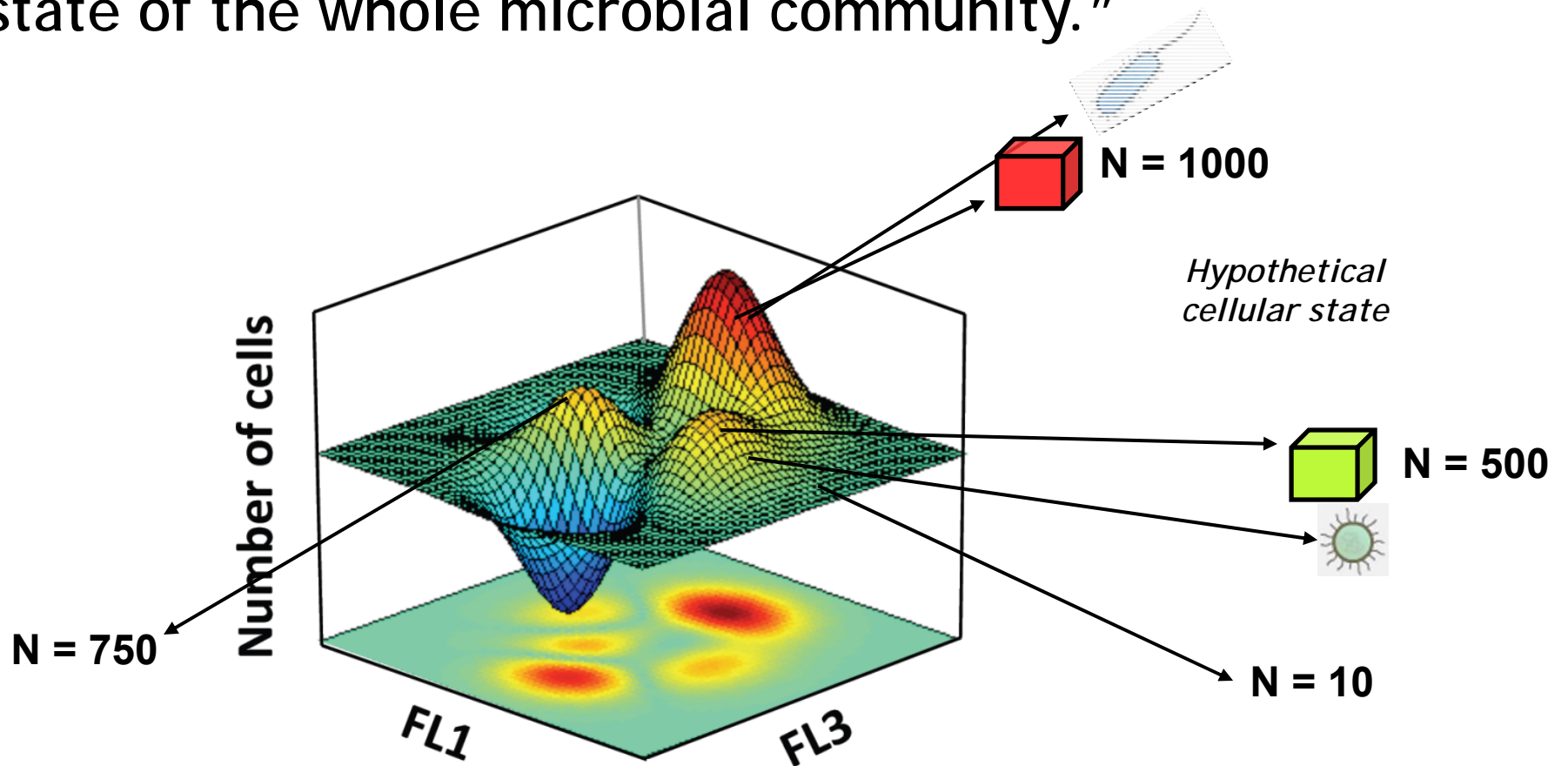
Multivariate data



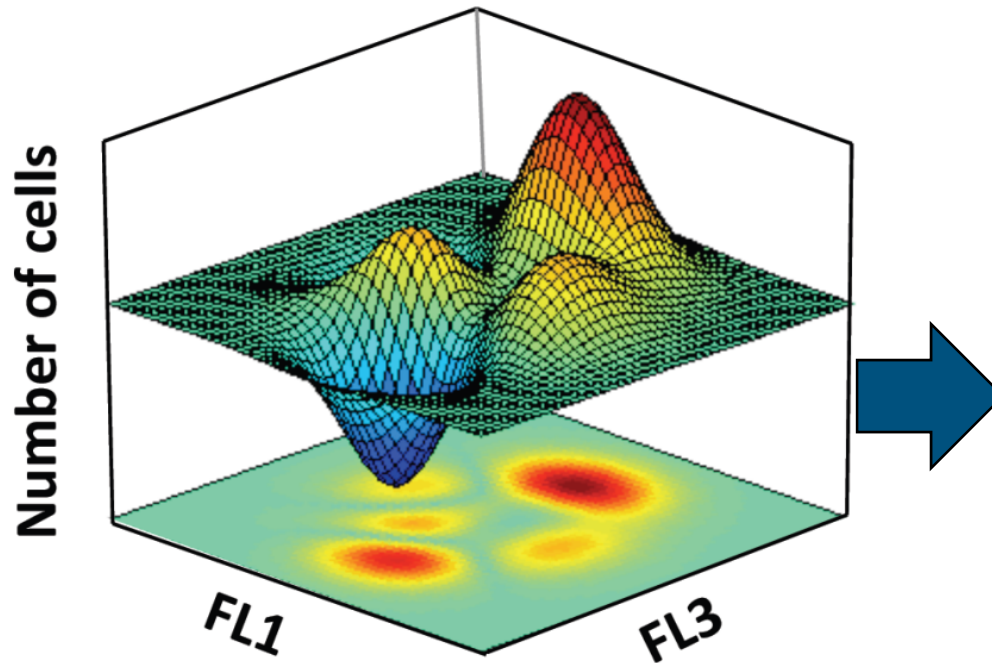
Morphology ~ Scattering  
DNA content ~ Fluorescence  
...



“The community landscape represents a phenotypic state of the whole microbial community.”



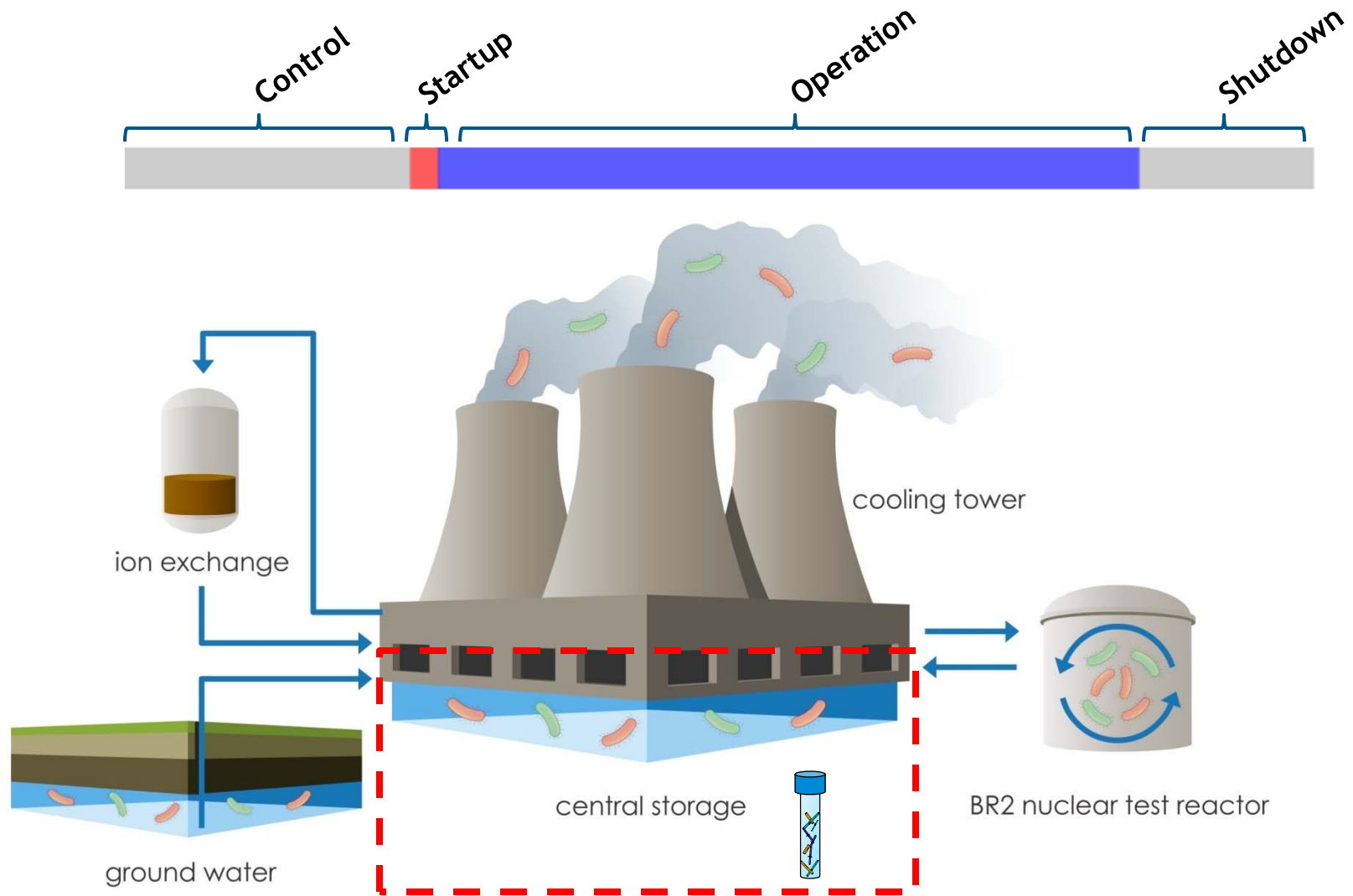
# Rationale



25	90	63	22	16	2	0	0	0	0
35	127	278	214	73	25	8	3	1	0
241	897	2532	3205	1736	588	155	52	9	3
581	2058	6762	11946	8168	3268	995	301	103	26
57	220	780	1596	1526	713	279	89	33	13
0	4	8	15	15	14	3	2	1	0

Hill Order (q)	Diversity metric ( $D_q$ )
0	$D_0 = S$
1	$D_1 = e^{-\sum_{i=1}^S p_i \ln(p_i)}$
2	$D_2 = \frac{1}{\sum_{i=1}^S (p_i)^2}$

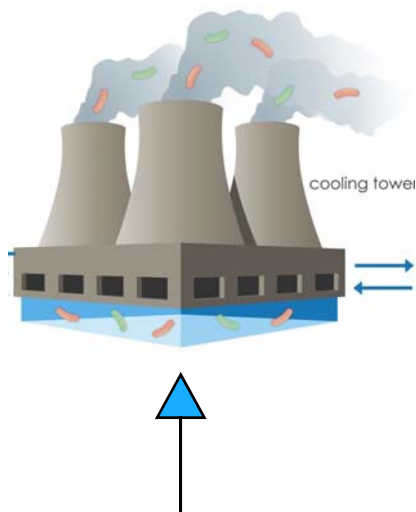
# Oligotrophic ecosystem model



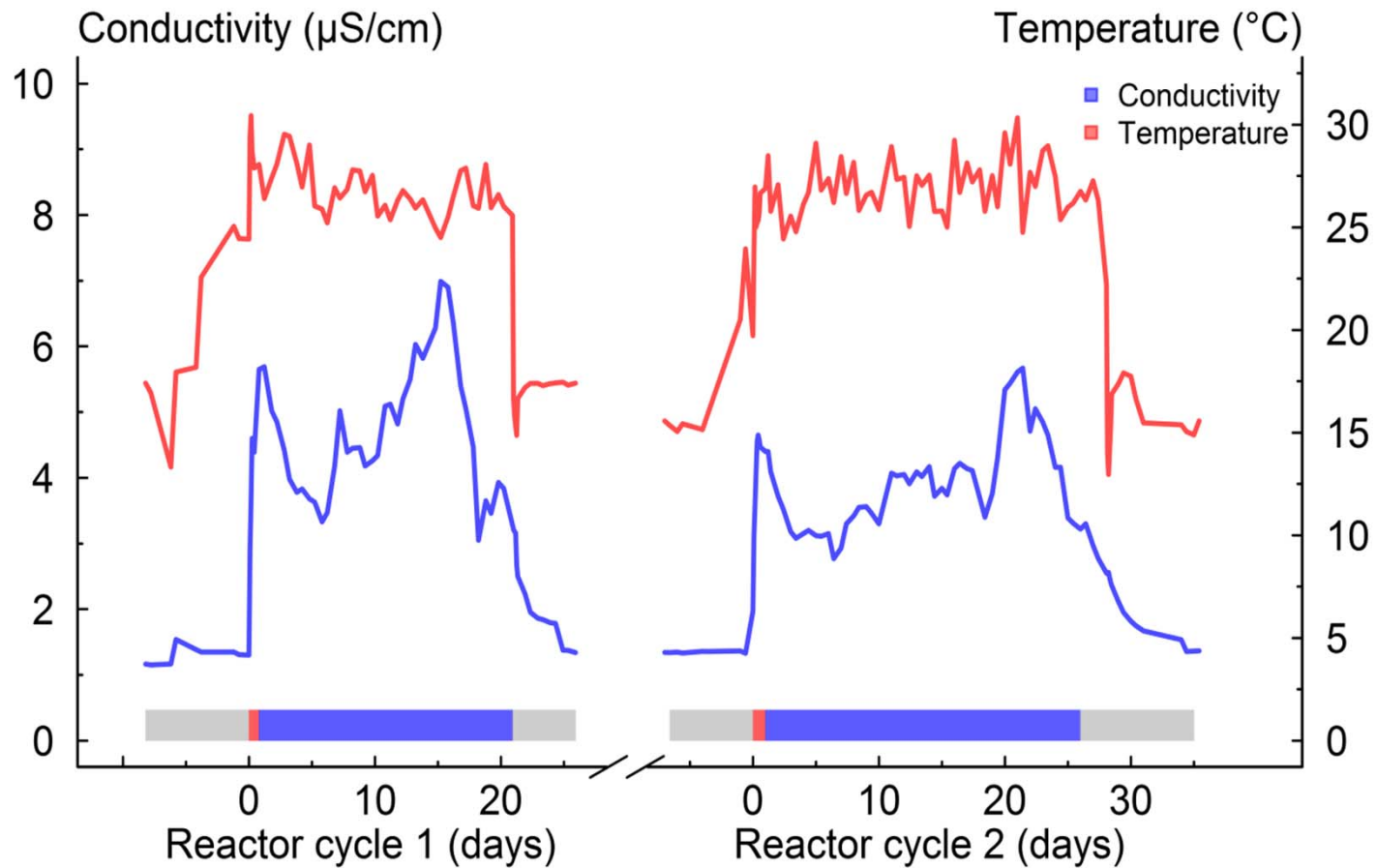


## Both a dynamic and stable system

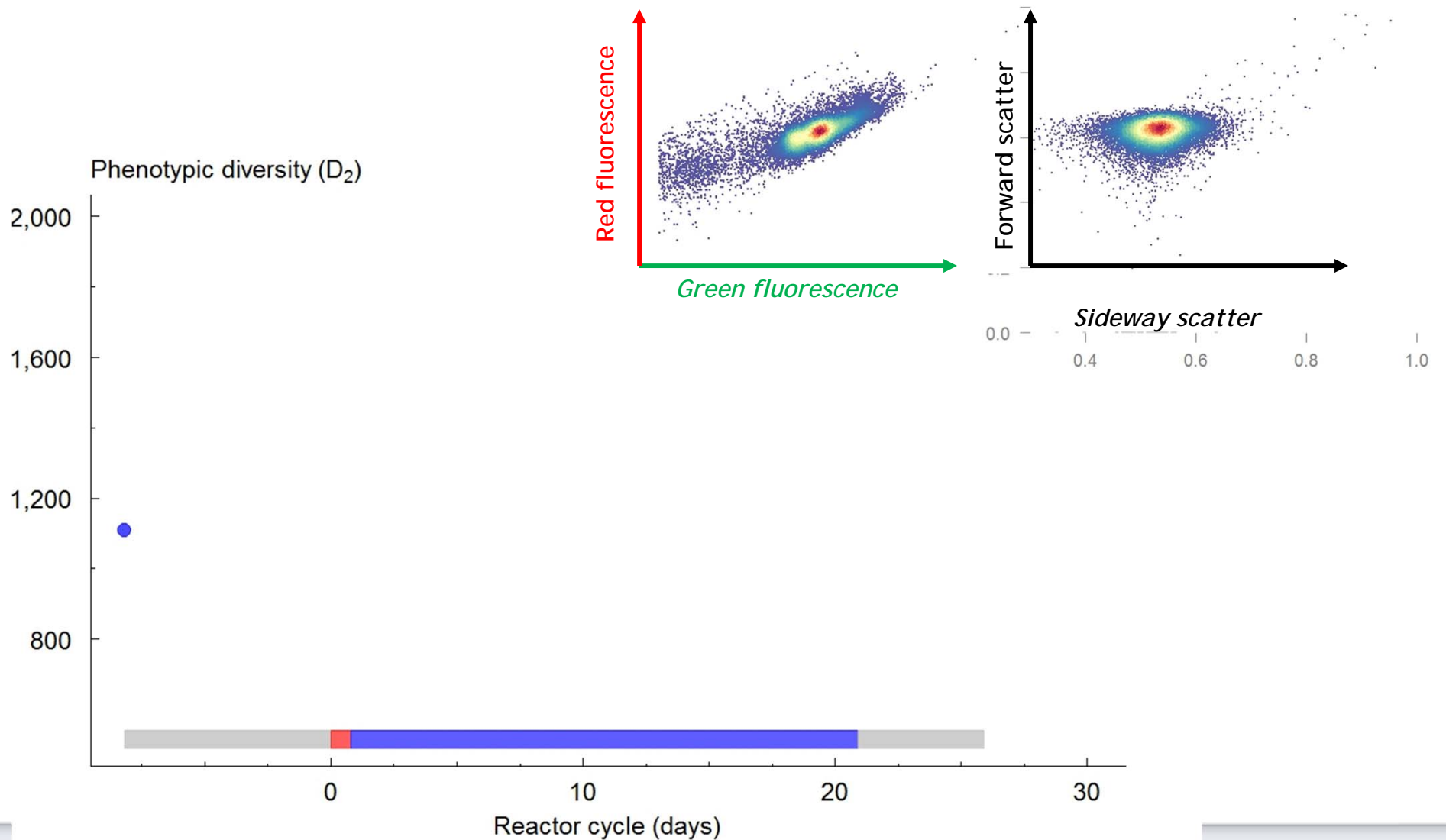
- $60 \text{ m}^3 \text{ h}^{-1} \rightarrow 4,000 \text{ m}^3 \text{ h}^{-1}$
- $15 \text{ }^\circ\text{C} \rightarrow 30 \text{ }^\circ\text{C}$
- $1 \text{ } \mu\text{S cm}^{-1} \rightarrow 7 \text{ } \mu\text{S cm}^{-1}$



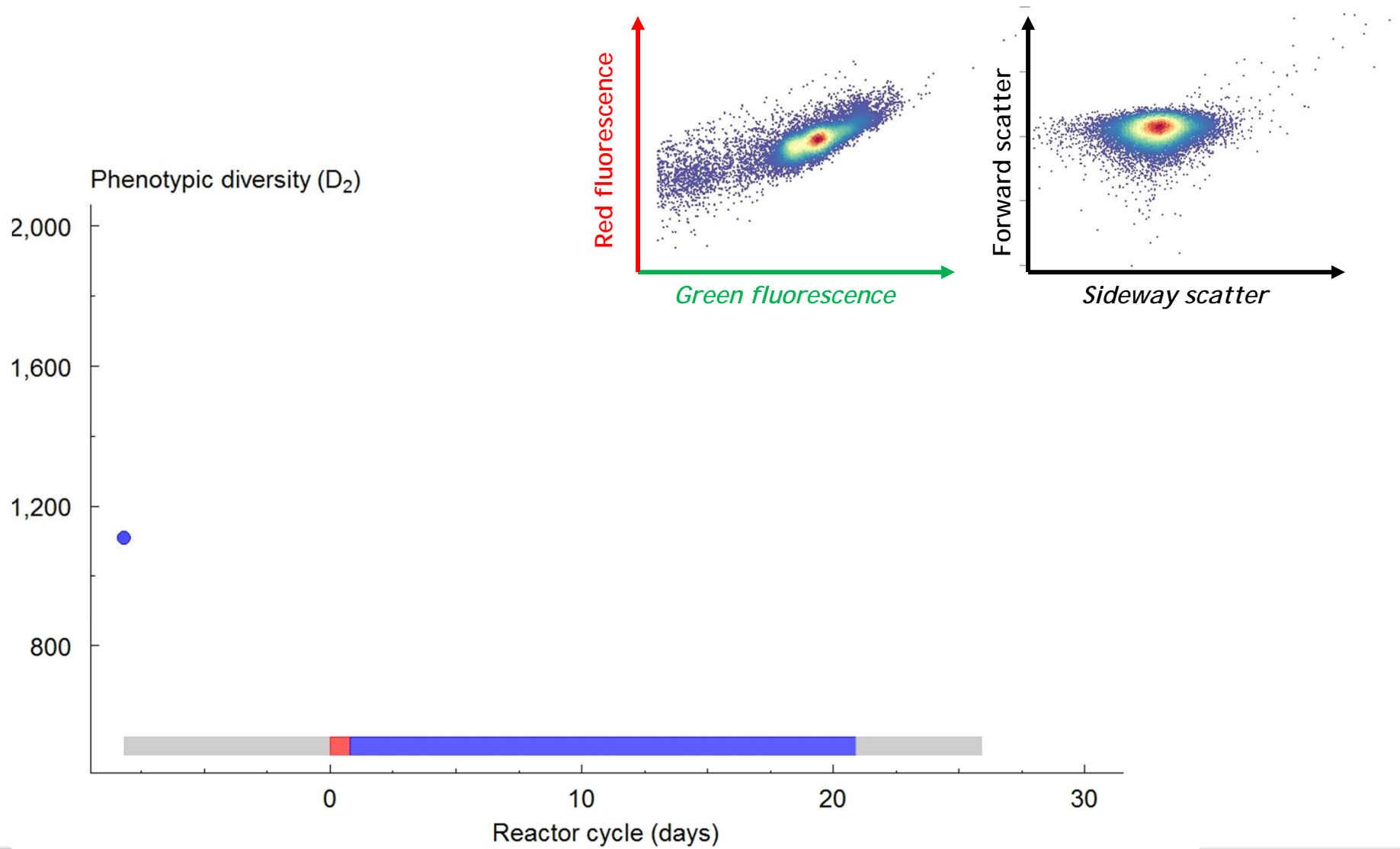
# Both a dynamic and stable system



# Monitoring the biodiversity

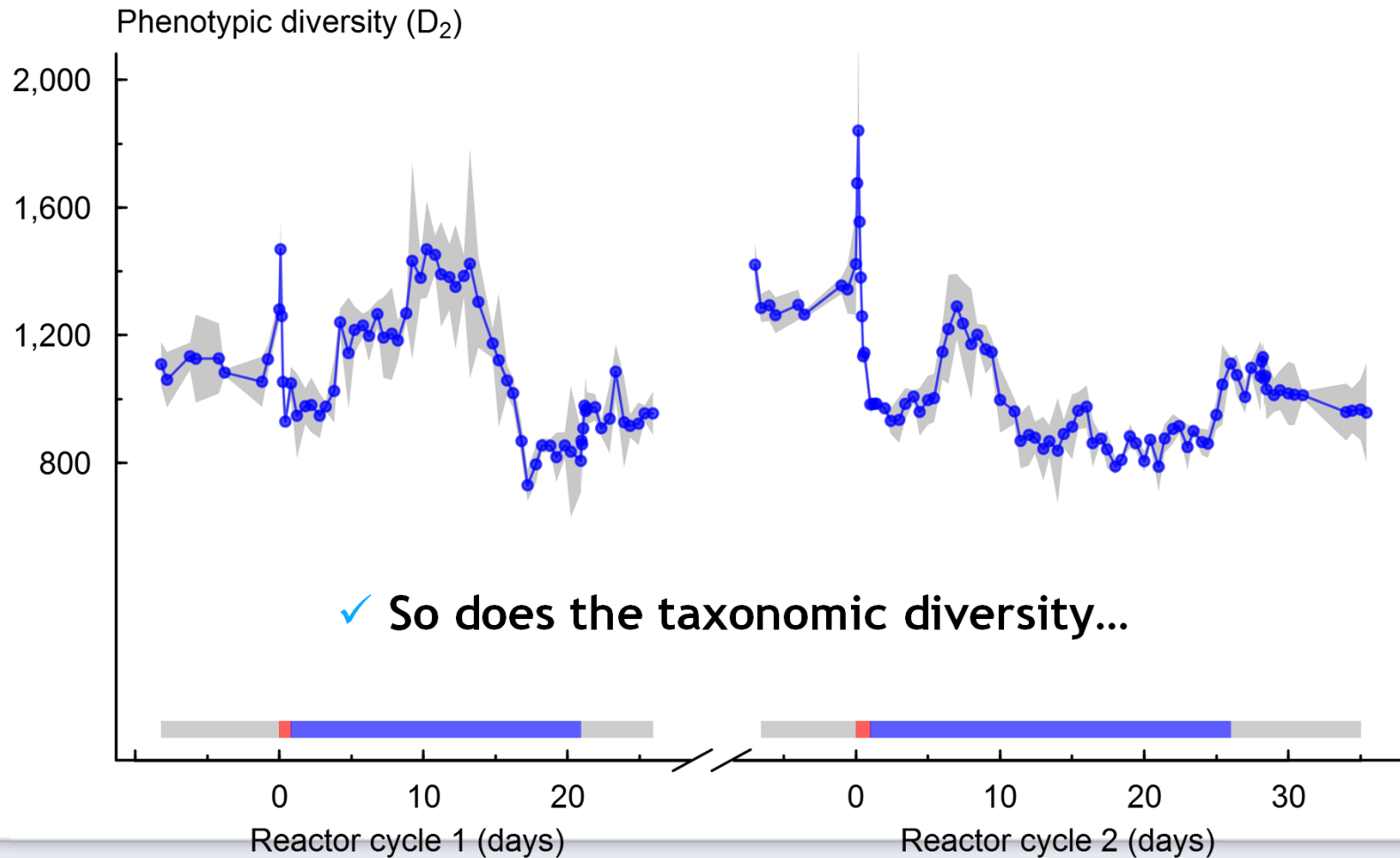


# Monitoring the biodiversity

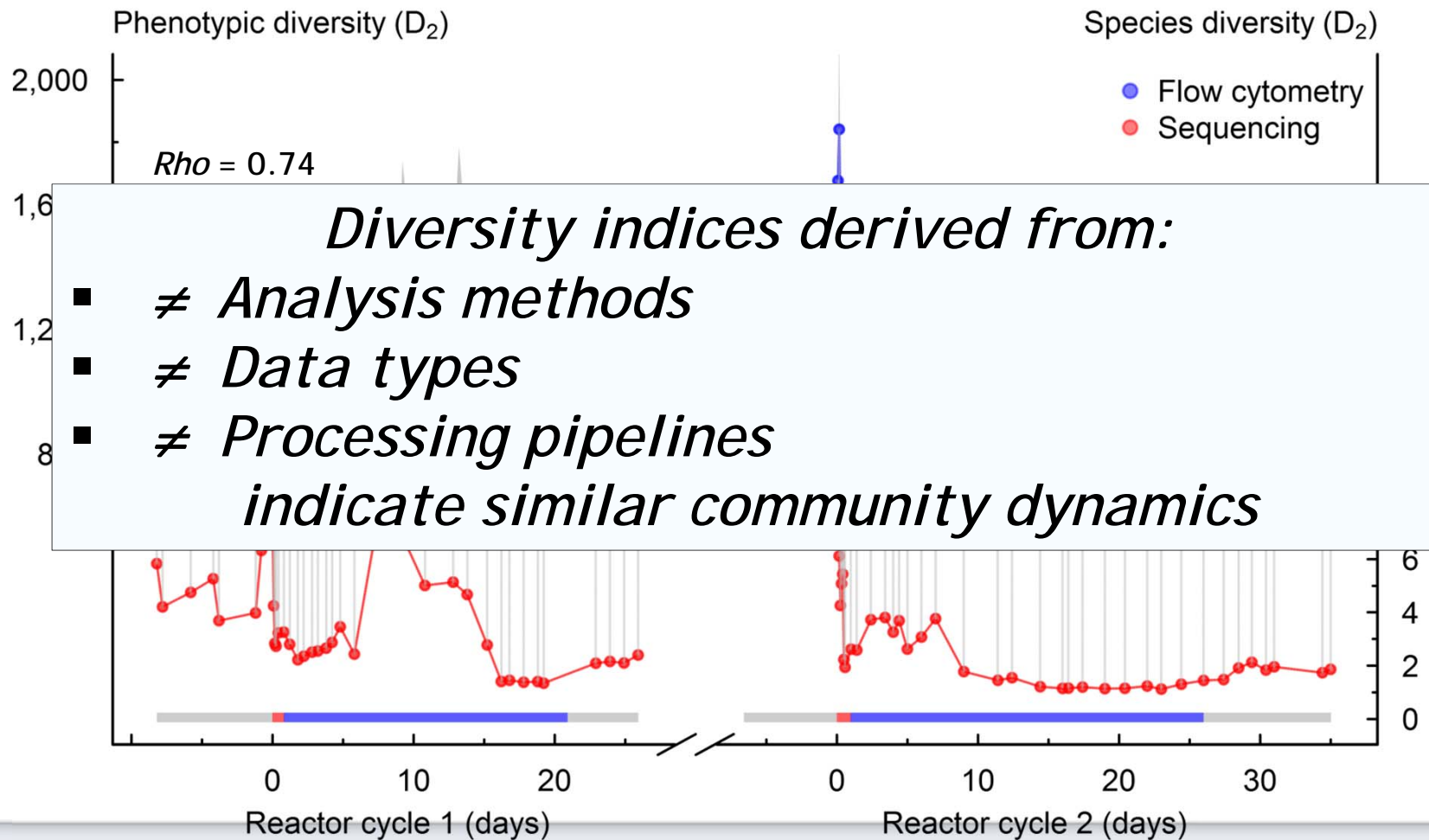


# Monitoring the biodiversity

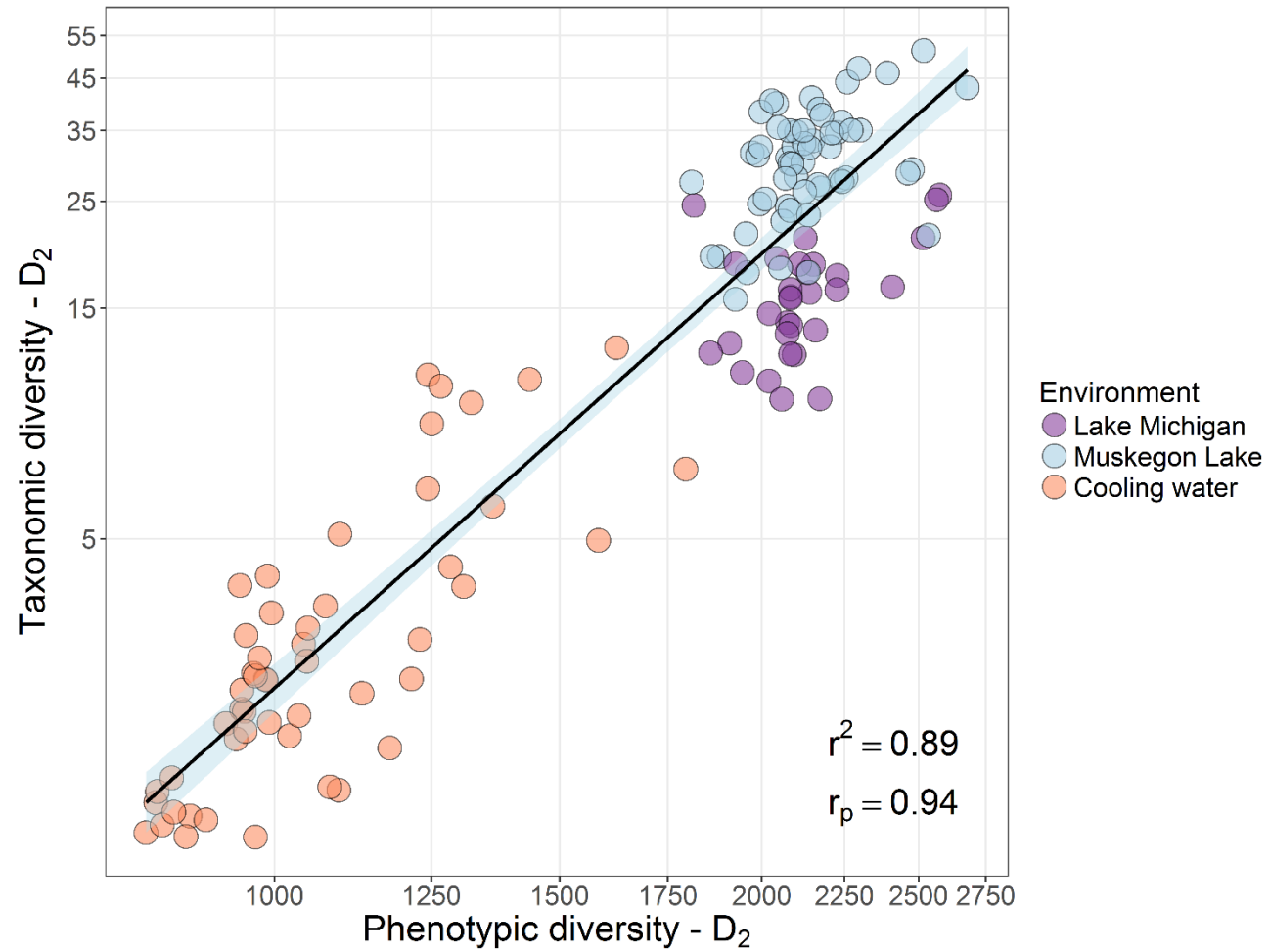
✓ Phenotypic diversity indicates dynamic changes



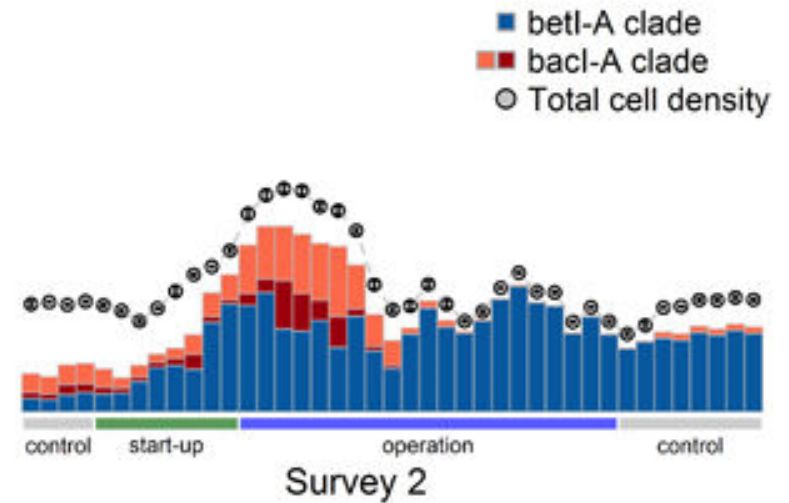
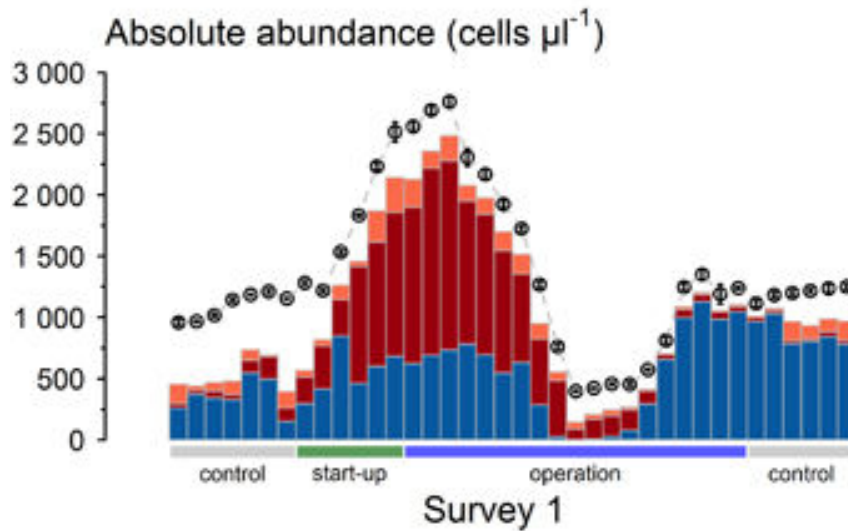
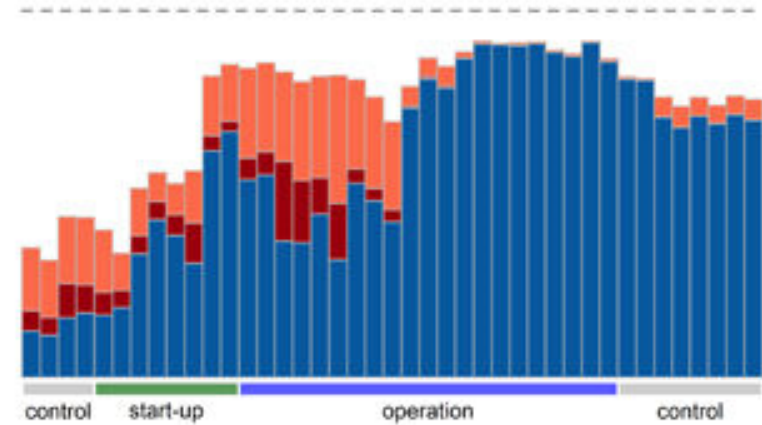
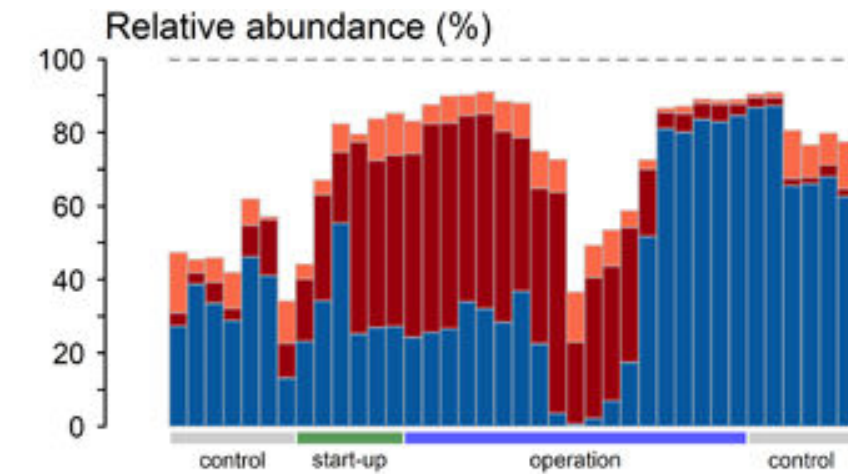
# Monitoring the biodiversity



# Other Ecosystems



# Absolute quantification of OTUs

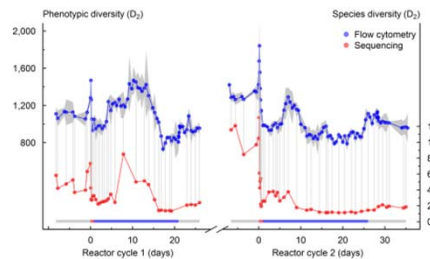




- ✓ In-situ real-time monitoring of biodiversity



- ✓ Intelligent experimental design



*When do I sample? And how frequently?*

- ✓ Complementary tool to sequencing platform

[https://github.com/rprops/Phenoflow\\_package](https://github.com/rprops/Phenoflow_package)

# Acknowledgements

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