## A sequencing-based approximation to phytoflagellate bacterivory at individual resolution



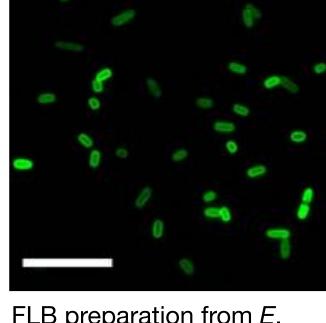
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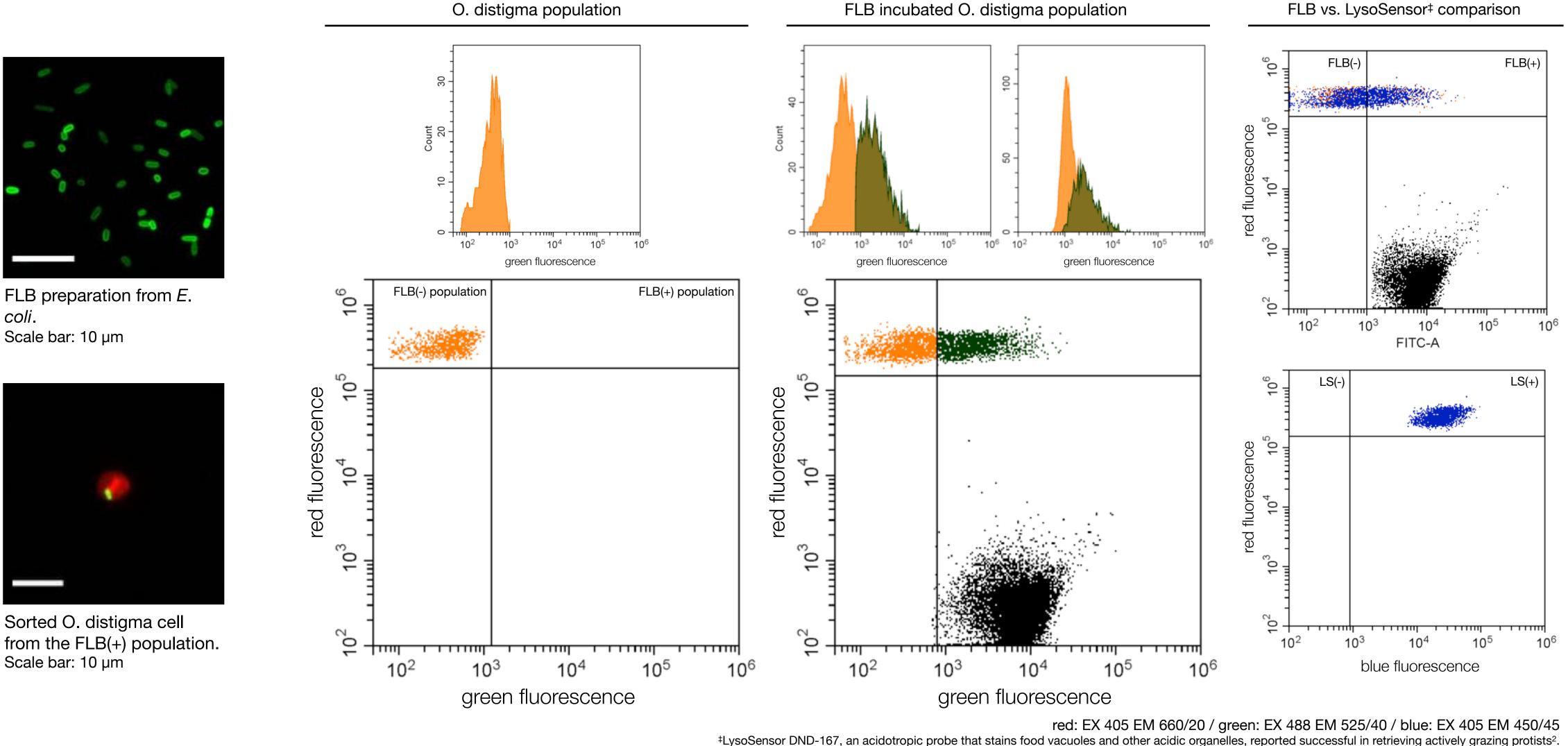
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cytometry tests with cultured Ochromonas distigma, used as a model mixotrophic phytoflagellate





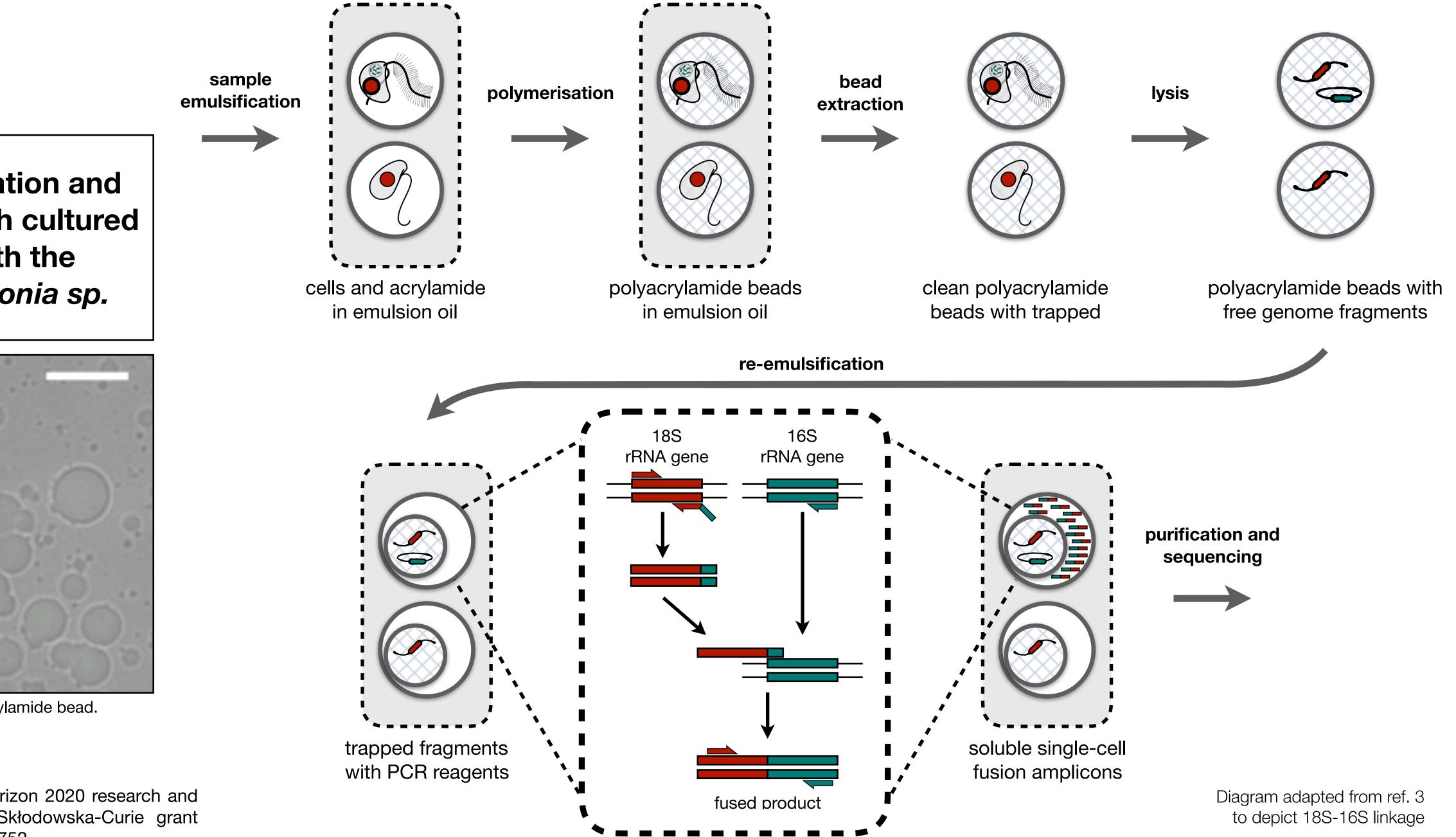
training network

**Mixotrophy** is largely unexplored but increasingly recognised as a rule rather than an exception within eukaryotic phytoplankters<sup>1</sup>. We are testing the efficiency, sensitivity and selectivity of employing **fluorescently labelled bacteria** (FLB) ingestion coupled to flow cytometry aiming to isolate mixotrophs via **fluorescence activated cell** sorting (FACS), taking advantage of the phagocytised FLB signal together with their natural chlorophyll autofluorescence.

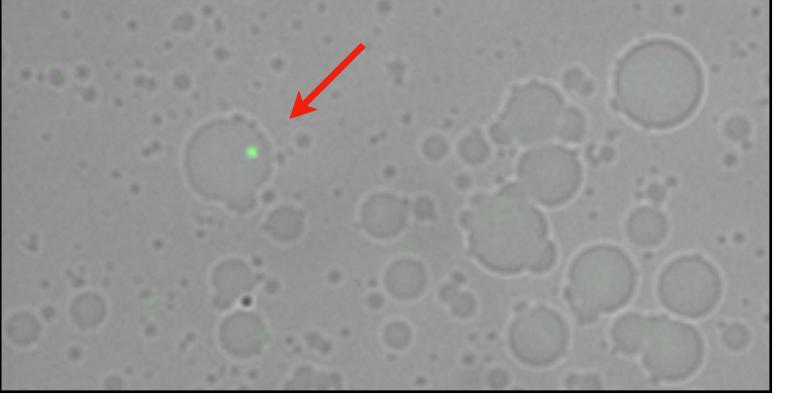
Who are they?

## Who do they prey upon?

Current methods to individually link prey to predator at microscopic level, such as CARD-FISH, require previous knowledge of the interacting species and thus lack in exploratory power. We are implementing emulsion, paired-isolation and concatenation PCR (epicPCR)<sup>3</sup> to uncover bacteria-protist interactions in aquatic environments via concatenation of taxonomic markers.



ongoing encapsulation and amplification tests with cultured O. distigma fed with the flavobacterium *Dokdonia* sp.



Encapsulated O. distigma cell within a polyacrylamide bead. Scale bar: 50 µm



## References

This work has been funded by the EU's Horizon 2020 research and innovation programme under the Marie Skłodowska-Curie grant agreement No. H2020-MSCA-ITN-2015-675752.

<sup>1</sup>Mitra, A. et al. Defining Planktonic Protist Functional Groups on Mechanisms for Energy and Nutrient Acquisition: Incorporation of Diverse Mixotrophic Strategies. Protist **167**, 106-120 (2016) <sup>2</sup>Carvalho, W. F. & Granéli. E. Acidotropic probes and flow cytometry: a powerful combination for detecting phagotrophy in mixotrophic and heterotrophic protists. Aquat Microb Ecol 44, 85-96 (2006). <sup>3</sup>Spencer, S. J., Tamminen, M. V. et al. *Massively parallel sequencing of single cells by epicPCR links functional genes with phylogenetic markers.* ISME J **10**, 427-436 (2016) The drawing of the flagellate with mastigonemes has been adapted from an original by Dennis Barthel (CC BY-SA 3.0).