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## Significance

- With an atmospheric life span of 7-12 years, the warming potential of methane (CH<sub>4</sub>) is 25 times greater than for carbon dioxide (CO<sub>2</sub>) over a 25 year period.
- More than 63% of atmospheric CH<sub>4</sub> (346 Tg Y<sup>-1</sup>) is derived from anthropogenic activities with landfills, oil and gas, and enteric fermentation being the main pollution sources.
- Methane oxidising organisms can fix CH<sub>4</sub>, providing a biological carbon sink by converting CH<sub>4</sub> to biomass carbon.
- This biomass can then be used for the production of value-adding renewable products (e.g. bioplastics, pigments, and lipids) providing added positive inputs into the techno-economics of CH<sub>4</sub> remediation.

## Background

- Four species of CH<sub>4</sub>-fixing yeasts were identified by Wolf *et al.*, (1980):
  - *Sporobolomyces roseus* and *S. gracilis*
  - *Rhodotorula rubra* and *R. glutinis*
- The ability of *R. glutinis* to utilise CH<sub>4</sub> as a sole carbon source was reported by Wolf *et al.* (1979, 1980).
- Reported CH<sub>4</sub>-oxidation rates were: 0.18 μmol CH<sub>4</sub>.mg dry weight.
- *Rhodotorula glutinis* is a unicellular, oleaginous, saprophytic yeast (Fig. 1) with a high lipid (>60%) and high carotenoid contents.
- As C18:1 was reported as the dominant fatty acid, *R. glutinis* fatty acids have potential for bioplastic production and the carotenoids (β-carotene, torulene and torularhodin) have high values on the antioxidant market (feed additives in aquaculture, natural food colourants, pharmaceuticals).
- Therefore, this study investigated the CH<sub>4</sub>-remediation potential of *R. glutinis* to evaluate benefits in a CH<sub>4</sub>-remediation context.

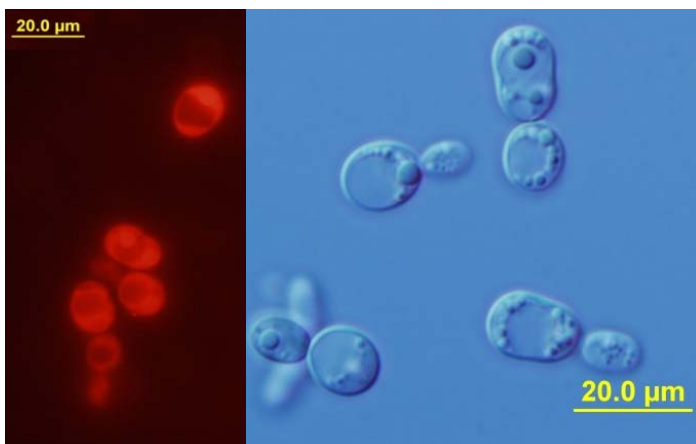


Fig 1. (Left) Fluorescence micrograph of *R. glutinis* stained with Nile-blue (Right) Differential interference micrograph of budding yeast cells (1,000x)

## Methods

- **Cultivation:** *Rhodotorula glutinis* cultures were grown in fed-batch reactors at 30 °C without shaking in basal medium (BM).
- **Methane treatments:** *R. glutinis* was purged with 20% CH<sub>4</sub> in air in batch cultures and 5% CH<sub>4</sub> in air in fermenter batch mode (Fig. 2). BM supplemented with CH<sub>4</sub> without *R. glutinis* served as control for non-biological CH<sub>4</sub> dissipation.
- **Carbon substrate utilisation:** *R. glutinis* was cultured in BM supplemented with acetate, glycerol, glycerol-diesel, or glycerol peptone.
- Growth was measured spectrophotometrically at 540 nm and via direct cell counts using a Neubauer haemocytometer.



Fig 2. Cultivation of *R. glutinis* in fermenter batch-mode, incubated with 5% CH<sub>4</sub> in air.

## Results

- Repetition of the exact same cultivation conditions used by Wolf *et al.* (1980) showed that *R. glutinis* did not fix CH<sub>4</sub>;
- Instead observed CH<sub>4</sub> declines were due to dissipation of CH<sub>4</sub> in the medium (non-organism control, Fig. 3).
- Best growth was achieved when cultivated in basal medium supplemented with glycerol-peptone (1%+0.001% w/w), while no growth was observed with CH<sub>4</sub> as the sole carbon source (Fig. 4).
- 22-day β-carotene productivities of *R. glutinis* on glycerol-peptone supplemented BM were 0.05 mg.g<sup>-1</sup>.dry weight.L<sup>-1</sup>.

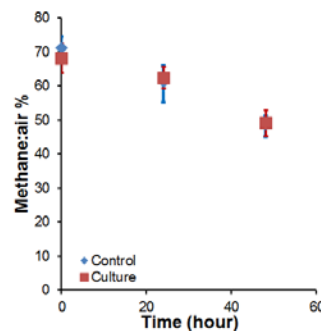


Fig 3. CH<sub>4</sub> dissolution in BM with (culture) and without *R. glutinis* (control)

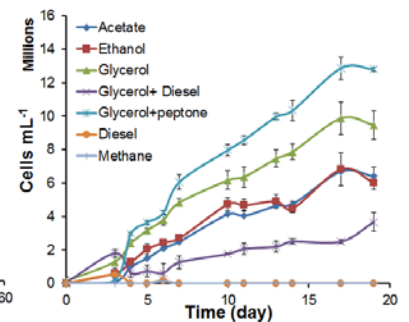


Fig 4. Growth of *R. glutinis* on different carbon sources

## Conclusions

- Our results show that *R. glutinis* is not a methane-oxidising yeast and refutes earlier reports by Wolf *et al.* (1979, 1980).
- *Rhodotorula glutinis* is not suitable for methane remediation.
- β-carotene productivities in glycerol-peptone-supplemented basal medium are not competitive with other natural sources.

## References

- Wolf, H. J., and R. S. Hanson. "Isolation and characterization of methane-utilizing yeasts." *Journal of General Microbiology* 114.1 (1979): 187-194.
- Wolf, H. J., M. A. R. C. I. A. Christiansen, and R. S. Hanson. "Ultrastructure of methanotrophic yeasts." *Journal of bacteriology* 141.3 (1980): 1340-1349.