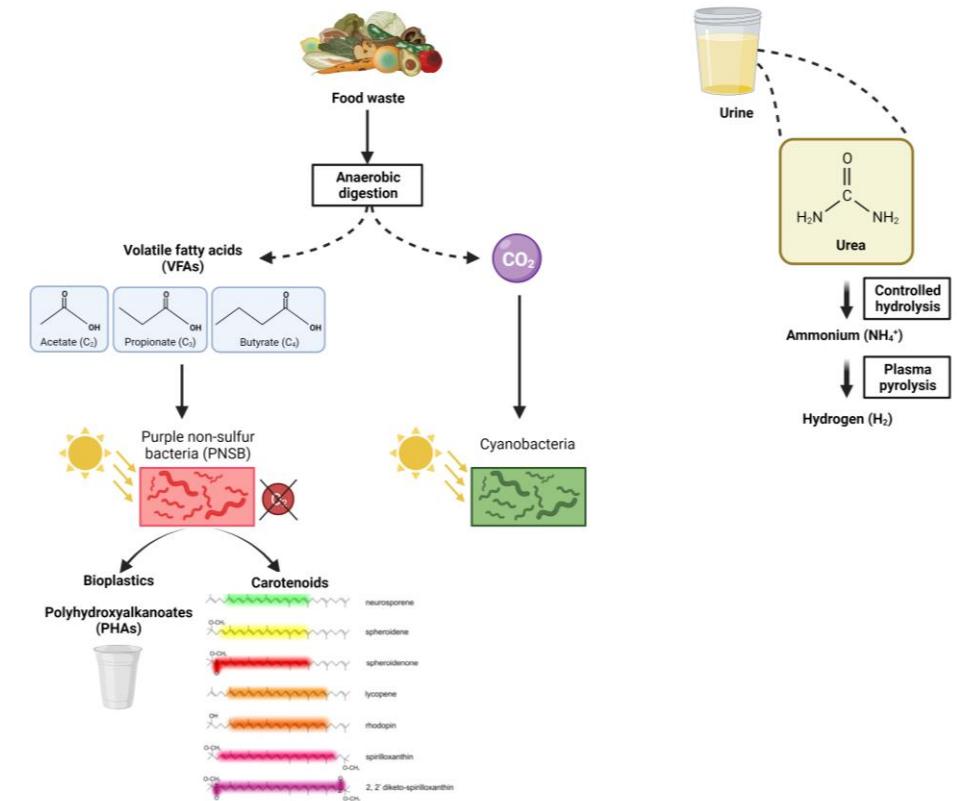


WalBioPower: Valorisation of organic waste for the production of clean energy

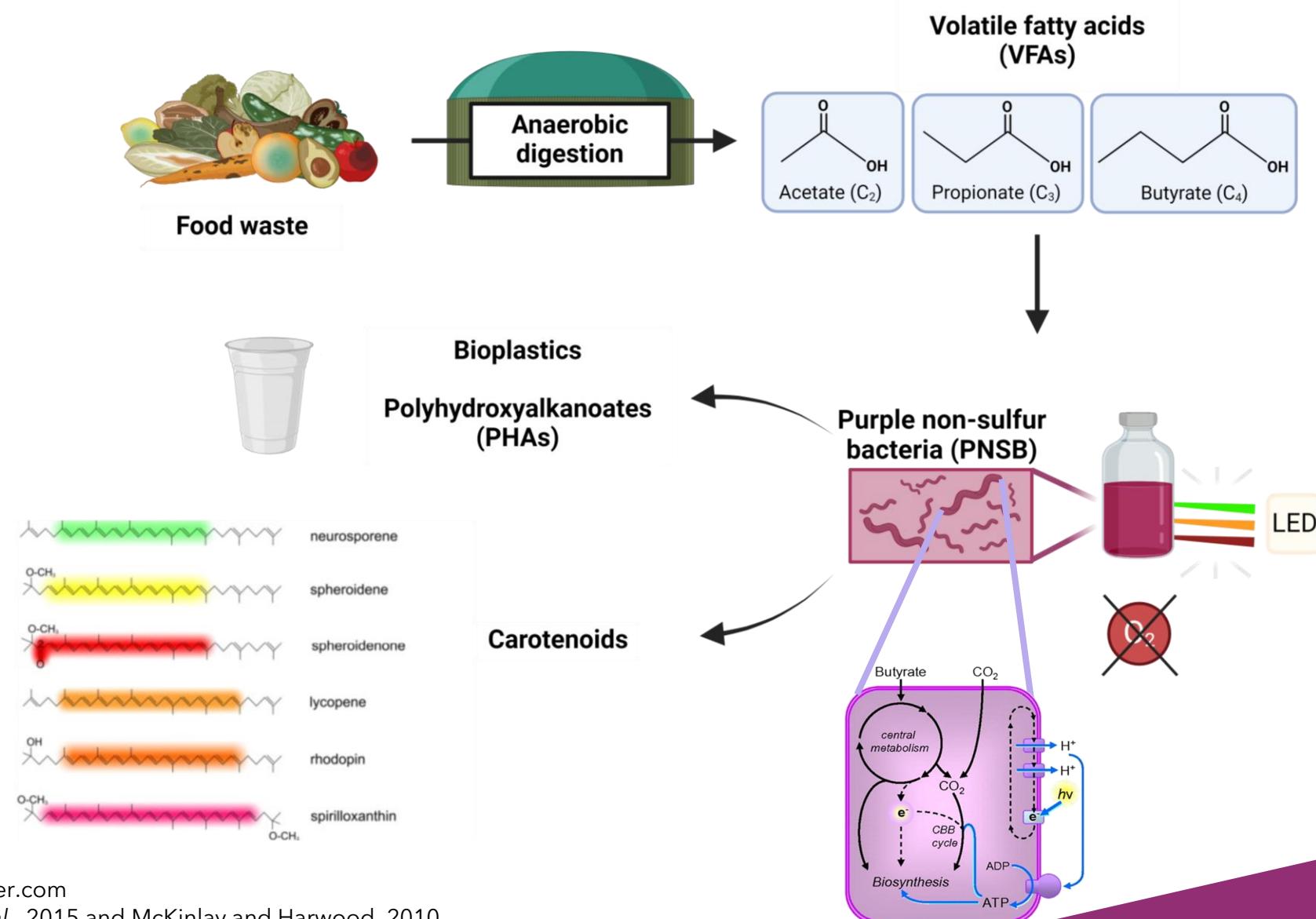
Simone Krings, Ruddy Wattiez, Baptiste Leroy

18/04/2024

DEPARTMENT OF PROTEOMICS
AND MICROBIOLOGY



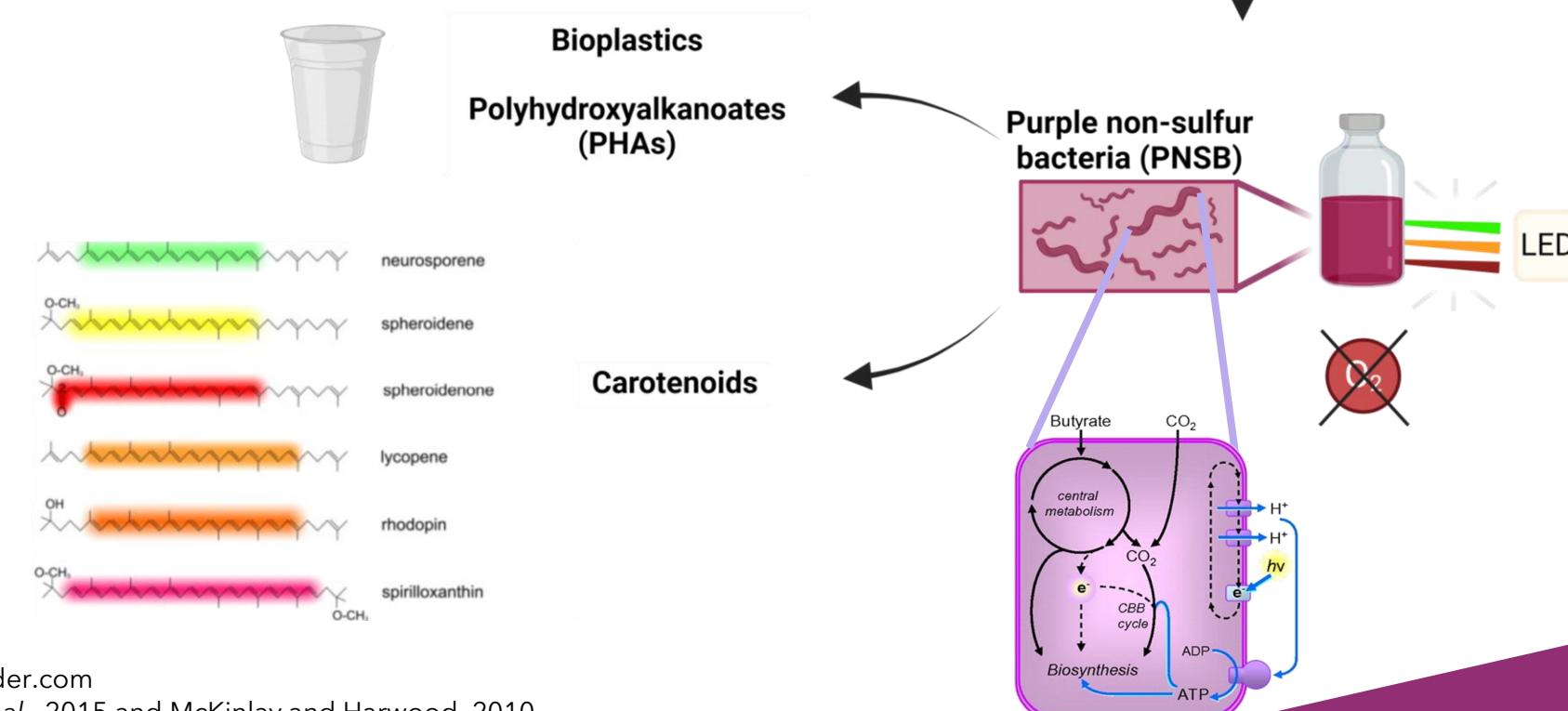
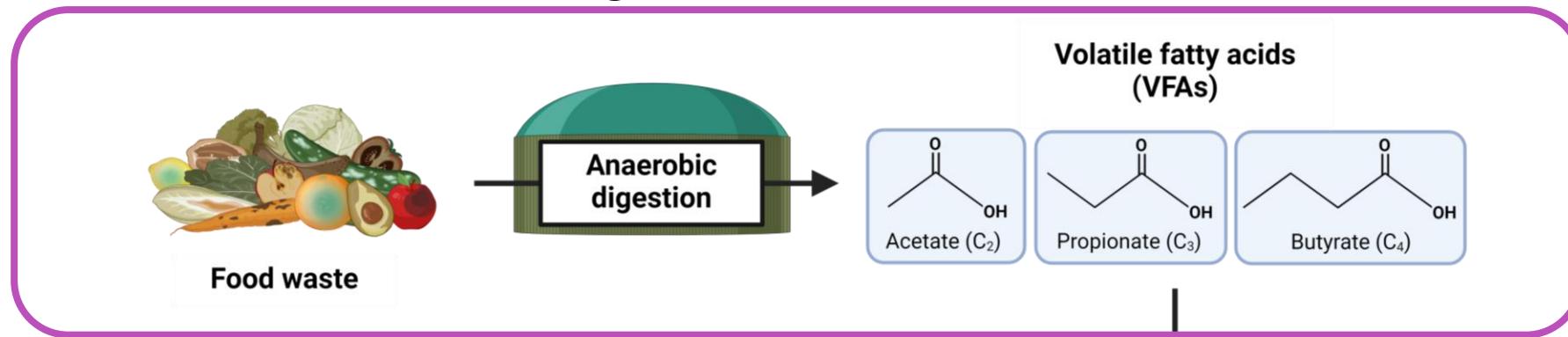
Project overview



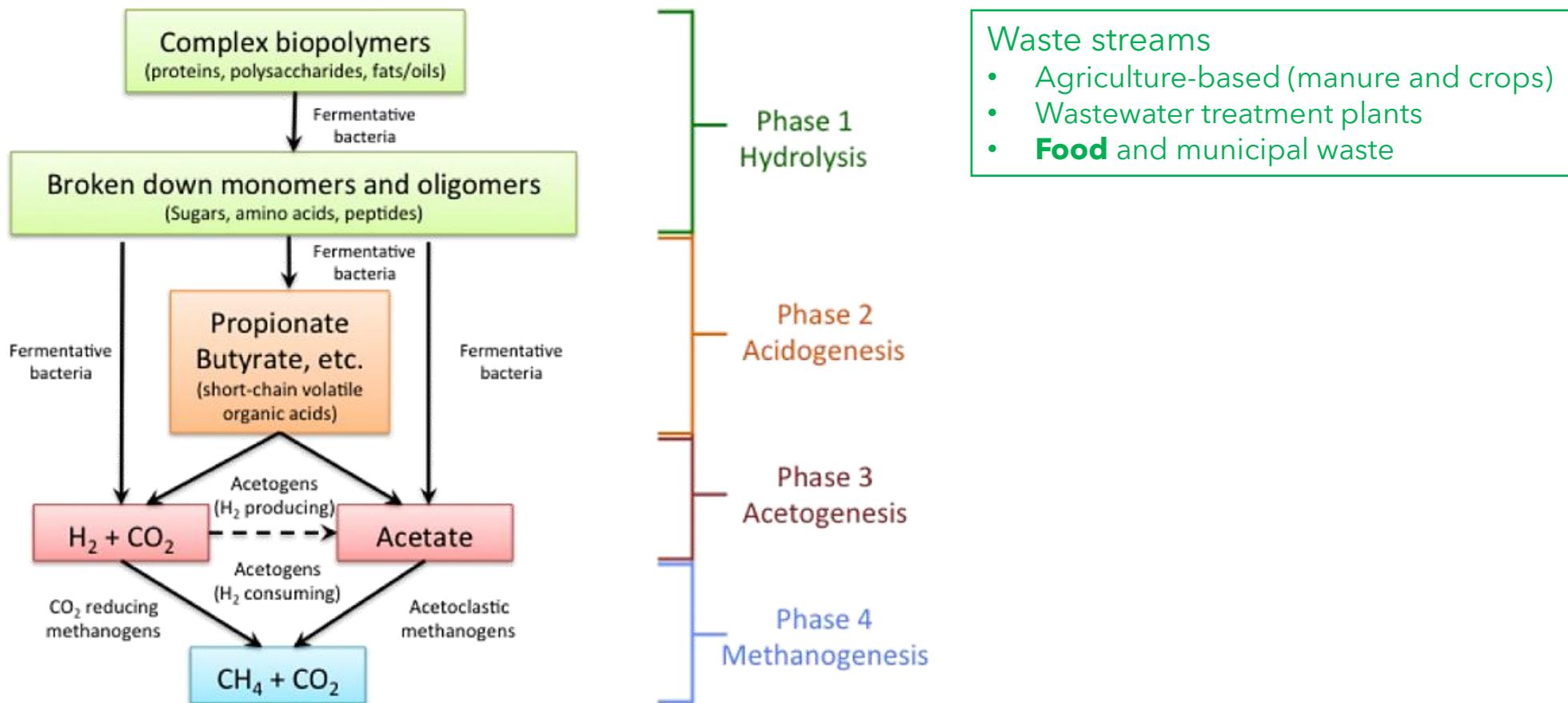
Created with Biorender.com

Adapted from Chi et al., 2015 and McKinlay and Harwood, 2010

Project overview

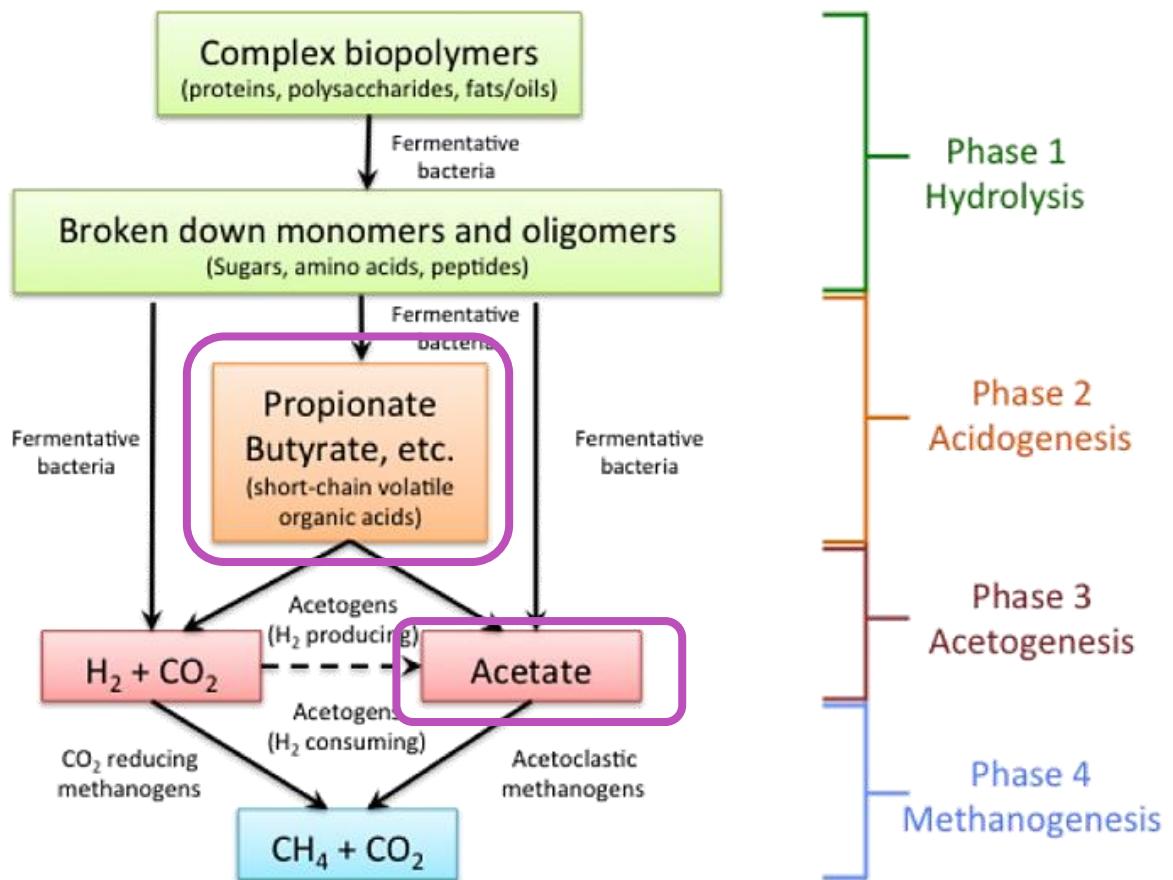


Anaerobic digestion (AD)



Credit: BEEMS Module B7 - Anaerobic Digestion

Anaerobic digestion (AD)



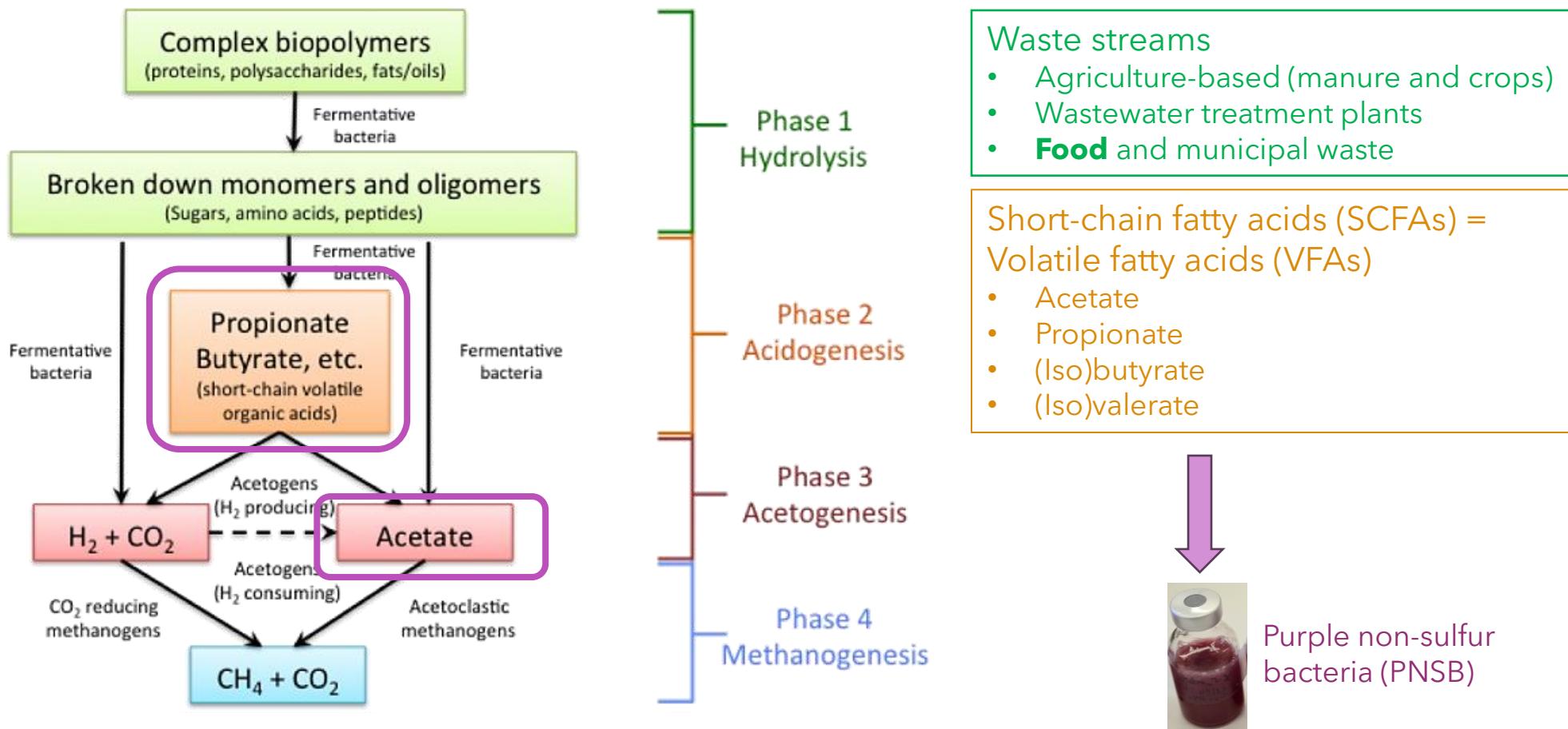
Waste streams

- Agriculture-based (manure and crops)
- Wastewater treatment plants
- **Food** and municipal waste

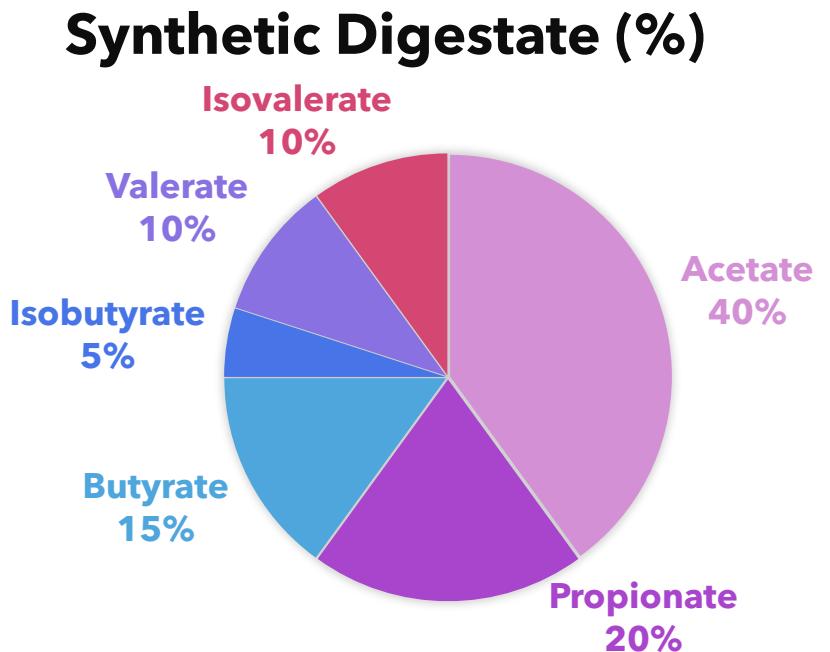
Short-chain fatty acids (SCFAs) =
Volatile fatty acids (VFAs)

- Acetate
- Propionate
- (Iso)butyrate
- (Iso)valerate

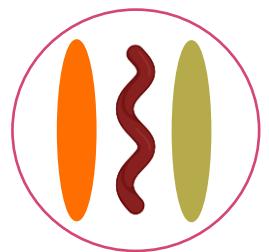
Anaerobic digestion (AD)



Culturing PNSB in Synthetic Digestate



- *Rhodospirillum rubrum* 
- Co-culture crs: *Rhodobacter capsulatus*, *Rhodospirillum rubrum* and *Cereibacter sphaeroides*

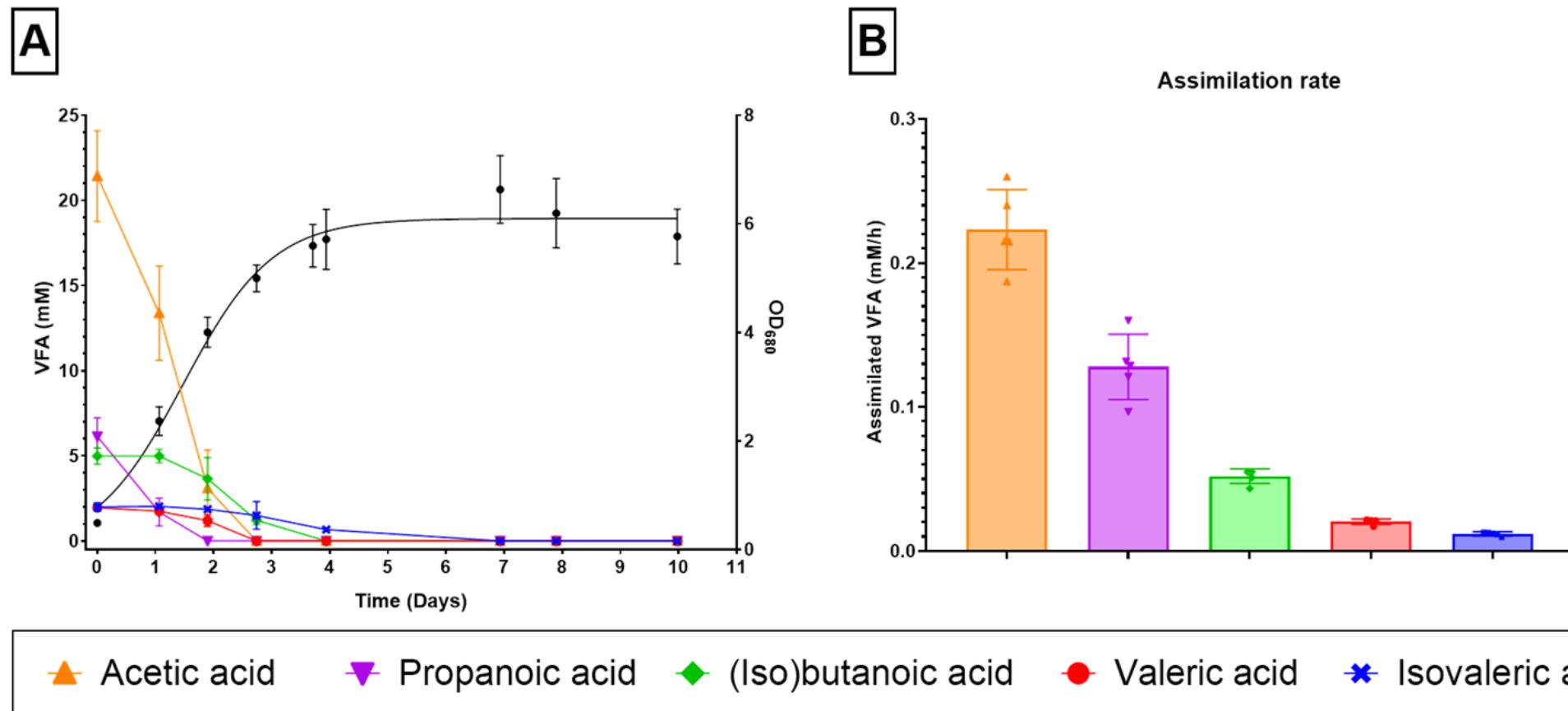


- Culture in SMN broth
- Preculture in MELiSSA medium + VFA (Acetate, Propionate and Butyrate (1:1:1) + 50 mM NaHCO_3 + Thiamine and Niacin
- Experiment in MELiSSA with synthetic digestate



Culturing PNSB in Synthetic Digestate

- LC-MS analysis of VFA assimilation





Bacterial strain proportions

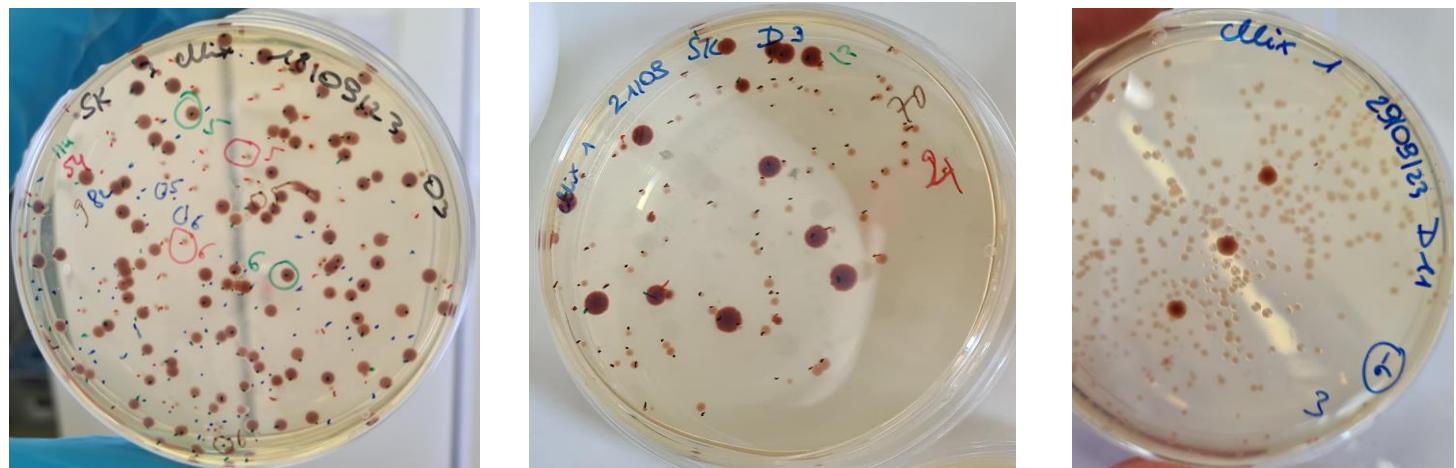
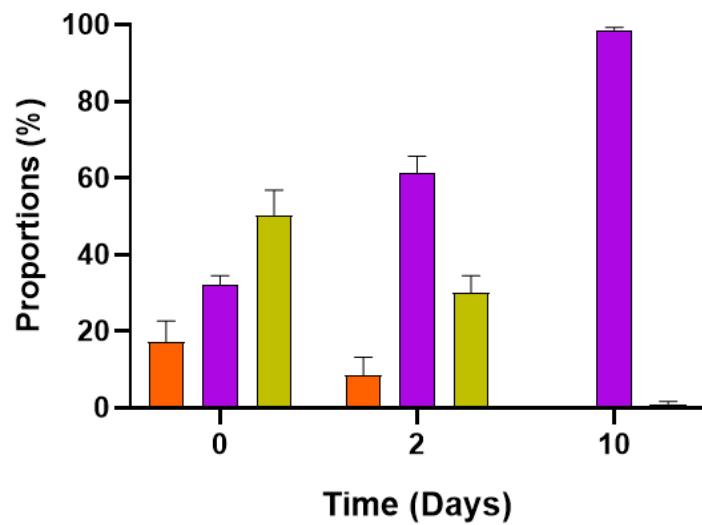
- $OD_{680} = 0.133$ of each strain => Achieve Start $OD_{680} = 0.5$



Bacterial strain proportions

- $OD_{680} = 0.133$ of each strain => Achieve Start $OD_{680} = 0.5$

Based on CFU/mL



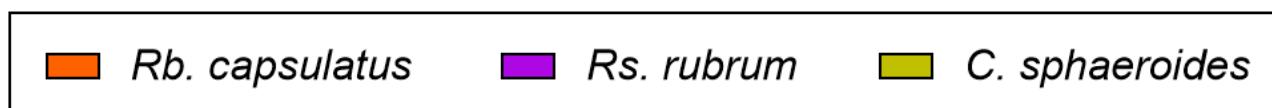
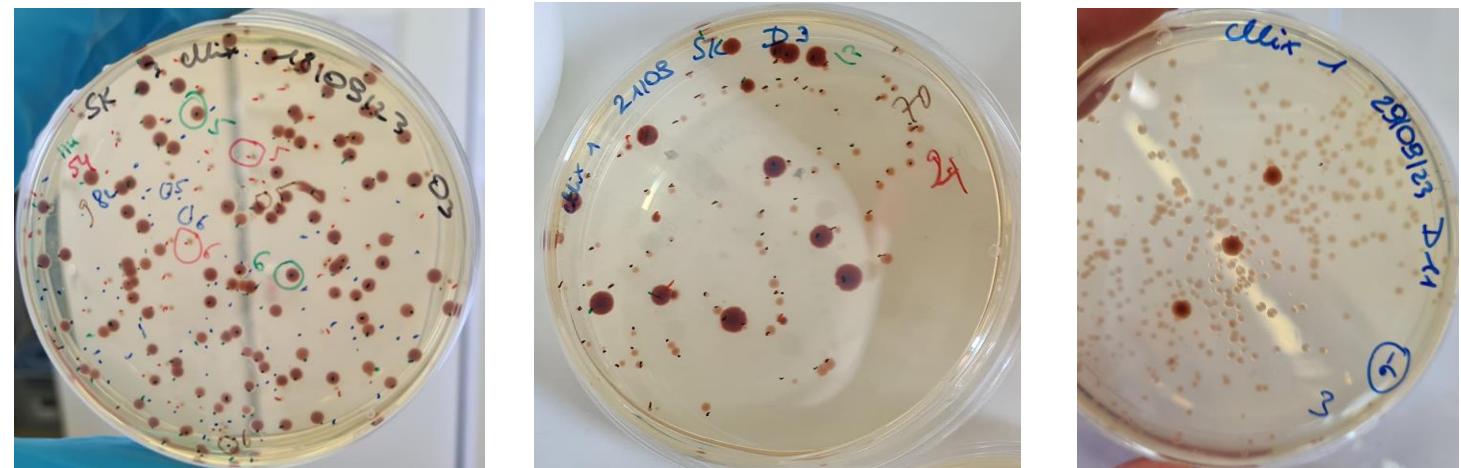
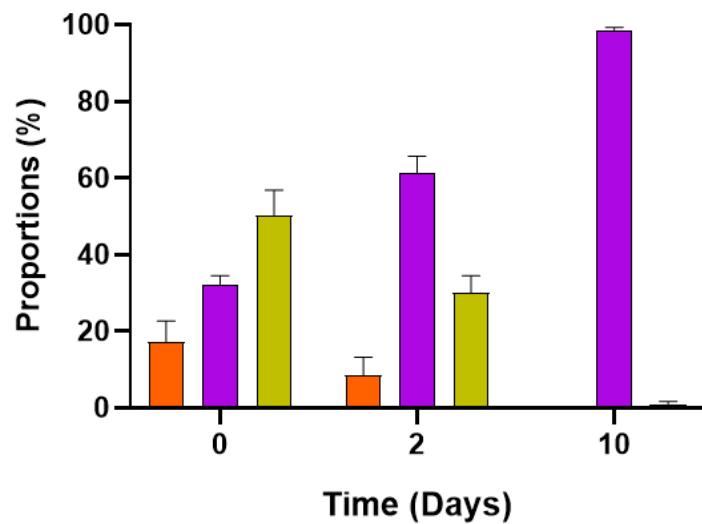
■ *Rb. capsulatus* ■ *Rs. rubrum* ■ *C. sphaeroides*



Bacterial strain proportions

- $OD_{680} = 0.133$ of each strain => Achieve Start $OD_{680} = 0.5$

Based on CFU/mL

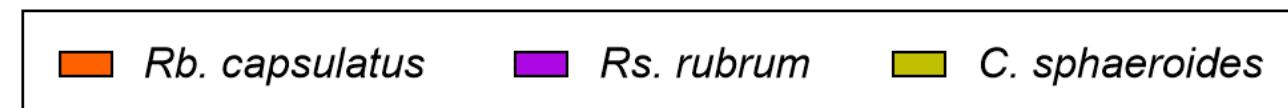
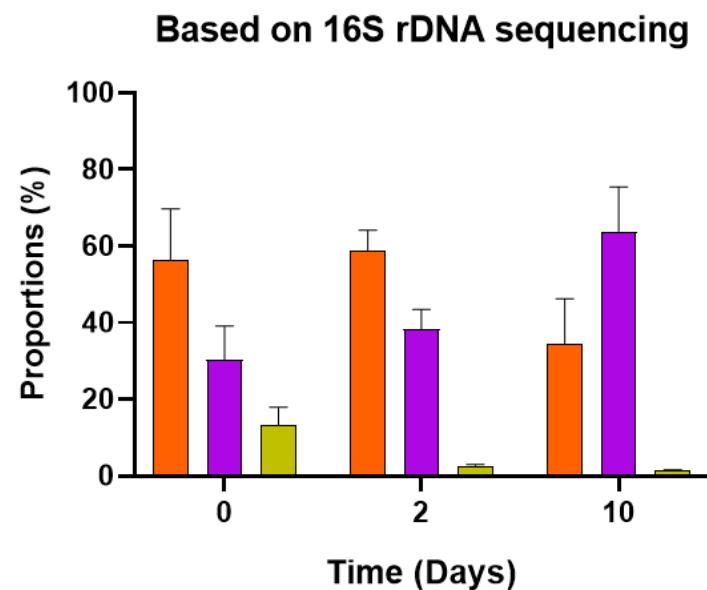
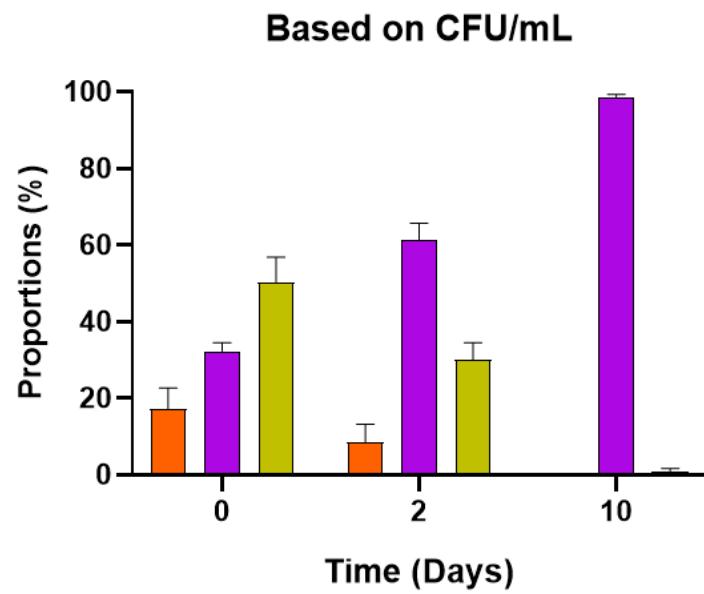


Similarity of colonies on agar plates



Bacterial strain proportions

- $OD_{680} = 0.133$ of each strain => Achieve Start $OD_{680} = 0.5$

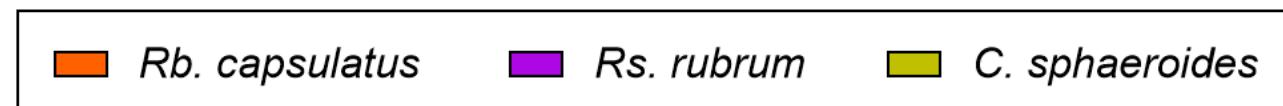
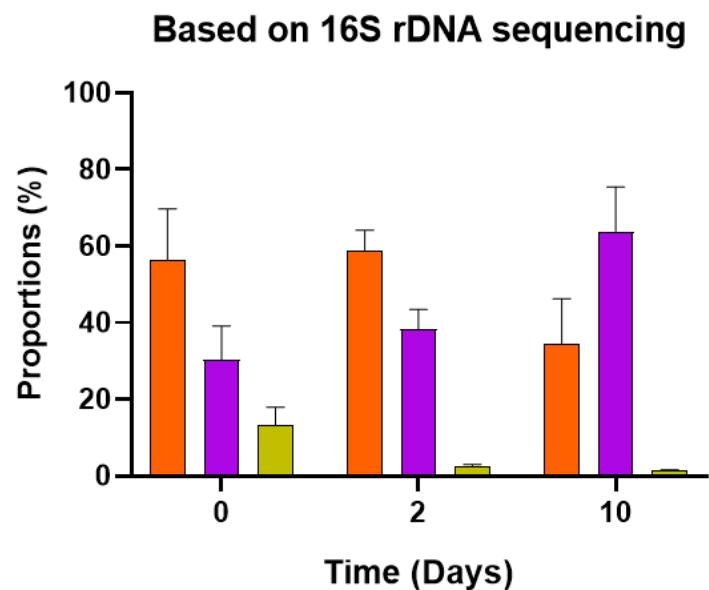
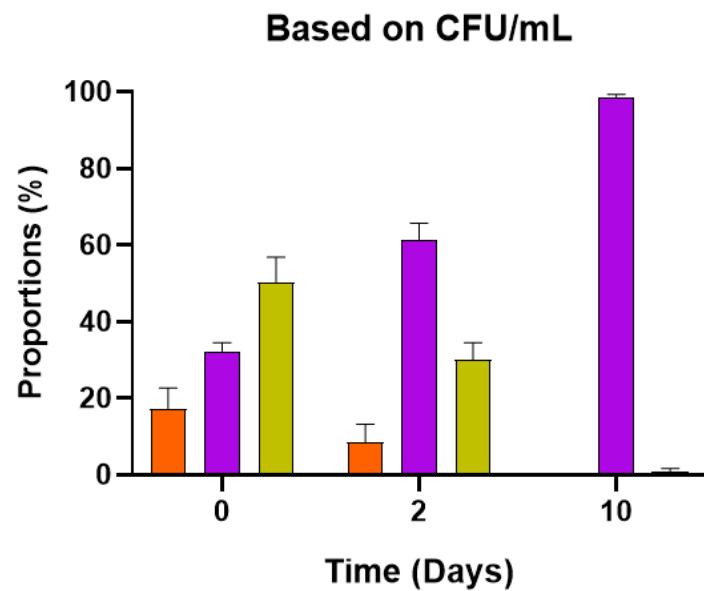


Oxford Nanopore GridION
+ 16S rDNA primers from Oxford
Nanopore



Bacterial strain proportions

- $OD_{680} = 0.133$ of each strain => Achieve Start $OD_{680} = 0.5$



Oxford Nanopore GridION
+ 16S rDNA primers from Oxford
Nanopore

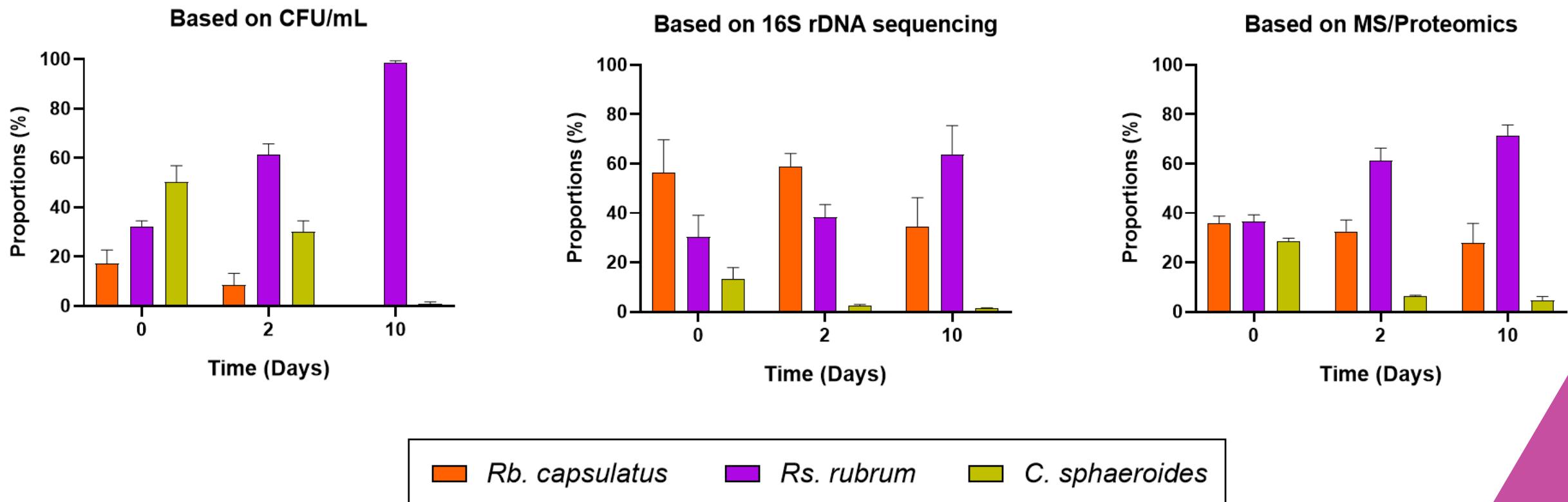


Primers have a higher affinity
for *Rb. capsulatus*



Bacterial strain proportions

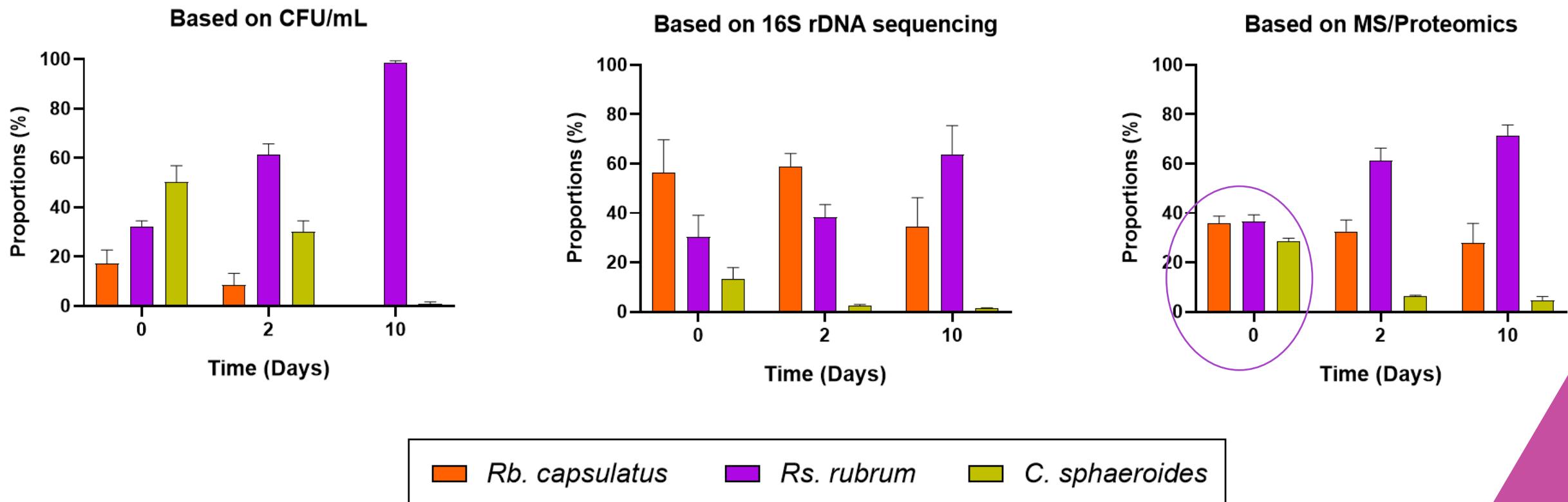
- $OD_{680} = 0.133$ of each strain => Achieve Start $OD_{680} = 0.5$





Bacterial strain proportions

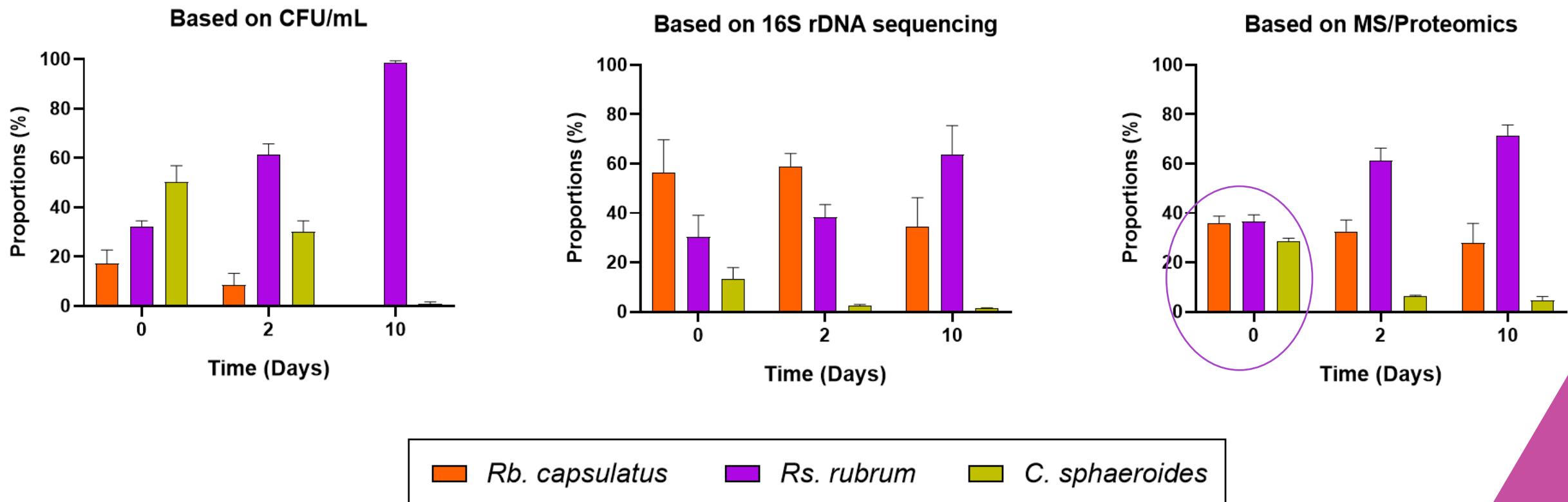
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Bacterial strain proportions

- $OD_{680} = 0.133$ of each strain => Achieve Start $OD_{680} = 0.5$



MS/Proteomics could be a great alternative to the classical methods!

Ammonium in Food-waste Digestates

Table 1

Food waste digestate characteristics (MC: moisture content; EC: electrical conductivity; OM: organic matter; TOC: total organic carbon; TN: total nitrogen; TP: total phosphorous; TK: total potassium).

Waste type	Country	MC (%)	pH	EC (dS/m)	OM (%)	TOC (%)	TN (%)	TP (%)	TK (%)	NH ₄ ⁺ -N (g/kg)	C/N ratio	References
OFMSW	Spain	71.8	8.5	6.8	45.3	—	3.2	0.75	—	—	—	García-Albacete et al. (2014)
Source selected biowaste	Spain	75.6	8.31	—	63	34.1	4.3	—	—	—	11.9	Cerda et al. (2019)
OFMSW	Spain	76	8.3	—	63	34	4.3	—	—	—	11.8	Rodríguez et al. (2019)
Food waste	Ireland	80	—	—	64.7	—	—	—	—	—	—	Chen et al. (2019)
OFMSW	Italy	92.1	7.69	—	65	35.7	—	—	—	—	—	Beggio et al. (2019)
Food waste	UK	92.5	8.05–8.24	—	—	—	3.3–4.4	0.56–0.76	1.05–1.55	2.9–3.6	—	Sánchez-Rodríguez et al. (2018)
Food waste	China	—	—	—	66.1	43.5	1.92	—	—	—	—	Liu et al. (2020)
Food waste	UK	93.9–95.6	8.1–8.8	—	—	—	5.4–8	—	—	3.9–6.3	—	Nicholson et al. (2017)
Food waste	China	96.4	8.6	—	91	—	4.8	0.53	—	2.54	—	Wang et al. (2020)
Residual household waste	France	69.8	8.2	—	38.5	—	—	—	—	4.18	—	Zeng et al. (2016)
Organic household waste	Germany	—	—	—	85.1	34.3	1.9	—	—	—	—	Cao et al. (2019)
Food waste	USA	—	—	2.5	—	—	6.96	1.2	1.9	3.4	—	Barzee et al. (2019)
Food waste	Italy	—	8.4–8.9	—	50.1–58.8	30.2–31.3	1.6–3.5	0.5–0.9	0.4–0.5	2.3–2.4	8.9–18.9	Grigatti et al. (2020)
Food waste	Poland	—	8.1	—	63	—	6.6	1.6	1.63	—	—	Czekala et al. (2020)
Segregated biodegradable waste	Italy	—	7.7	—	64	31.4	4.9	—	—	3.48	—	Peng et al. (2018)
Food waste	Singapore	—	7.81	—	—	—	—	0.5	—	4.7	—	Cheong et al. (2020)
Food waste	Austria	—	7.8	—	—	39.7	1.4	—	—	5.1	—	Franke-Whittle et al. (2014)
OFMSW	Canada	—	8.5	10.8	48	24.5	1.6	1.5	1.5	6.19	—	Arab and McCartney (2017)
Food waste	Finland	—	7.6–8.3	—	50.2–63.7	25.9–26.9	7.8–8.7	0.1–0.3	1.9–3.2	3.2–4.5	3.1–3.3	Tampio et al. (2016)
OFMSW	Canada	—	8.5	0.8	—	—	—	1	—	5	—	McLachlan et al. (2004)
Food waste	Australia	—	8.02	—	—	42.1	5.8	1.97	0.62	1.5	—	Opatokun et al. (2016)
Food waste	Italy	—	7.97	—	80.4	—	3.97	0.9	2.3	—	—	Peng and Pivato (2019)
OFMSW	Italy	—	8.8	—	68–71	12.8–22.7	1.09	1.49	0.78	—	12.1–20.9	Peng and Pivato (2019)
Food waste	Germany	—	7.3–8.3	—	—	4.2–6.7	—	—	—	5.1–7.2	—	Fuchs and Drosig (2013)
Bio waste	Germany	—	7.6–8.1	—	—	—	3–6.8	—	—	1.5–5.6	—	Fuchs and Drosig (2013)
Fruit waste	Australia	94.8	—	—	73.1	43.5	9.6	0.8	1	—	—	Serrano et al. (2020)
Household organic waste	Switzerland	—	—	—	64.7	33.1	4.7	2.45	0.63	—	—	Loes et al. (2018)

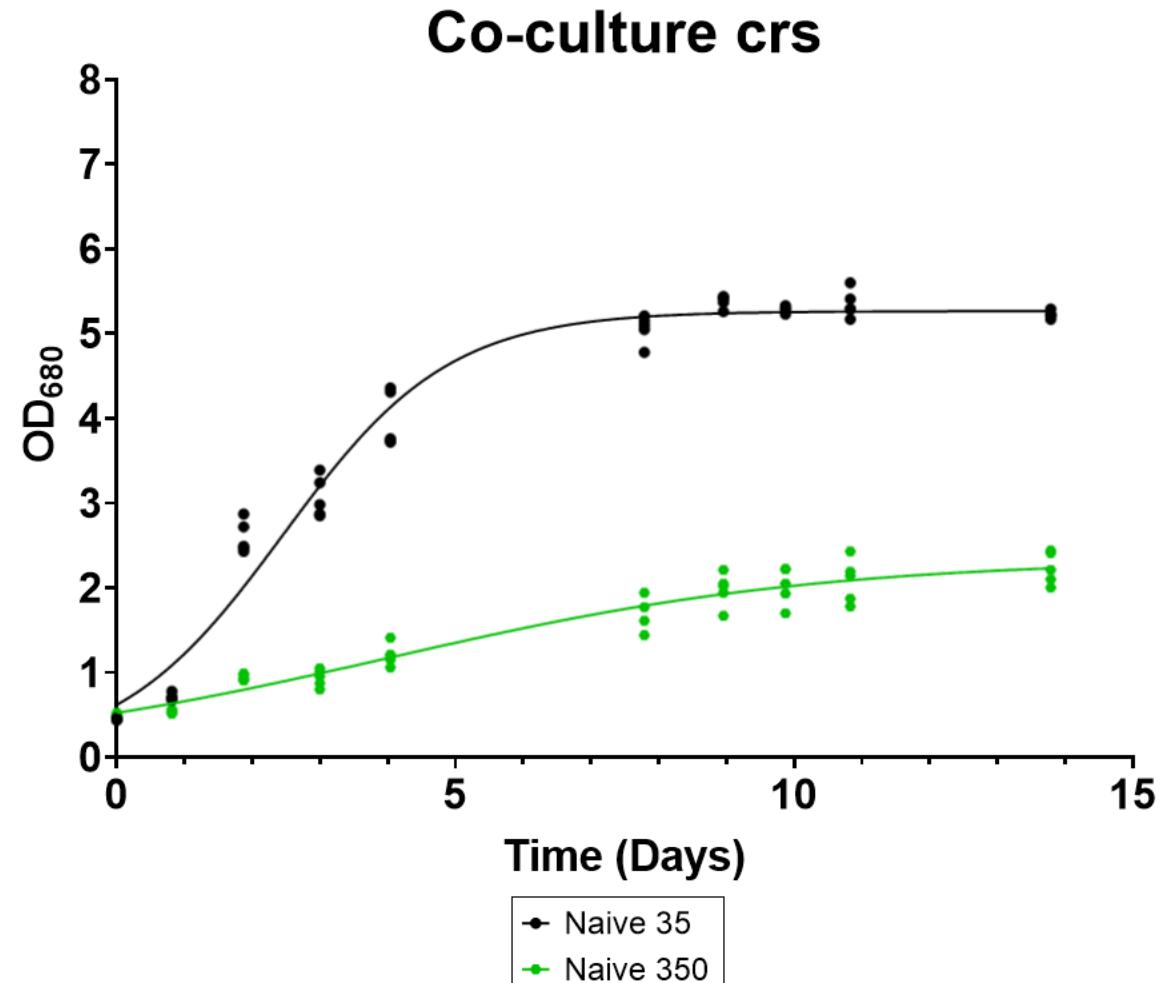
10 x

Standard MELiSSA
Medium:
35 mM NH₄Cl
→ 0.5 g/L N
→ 0.635 g/L NH₄⁺

~350 mM NH₄Cl
→ 5 g/L N
→ 6.35 g/L NH₄⁺



Effects of High Ammonium Levels



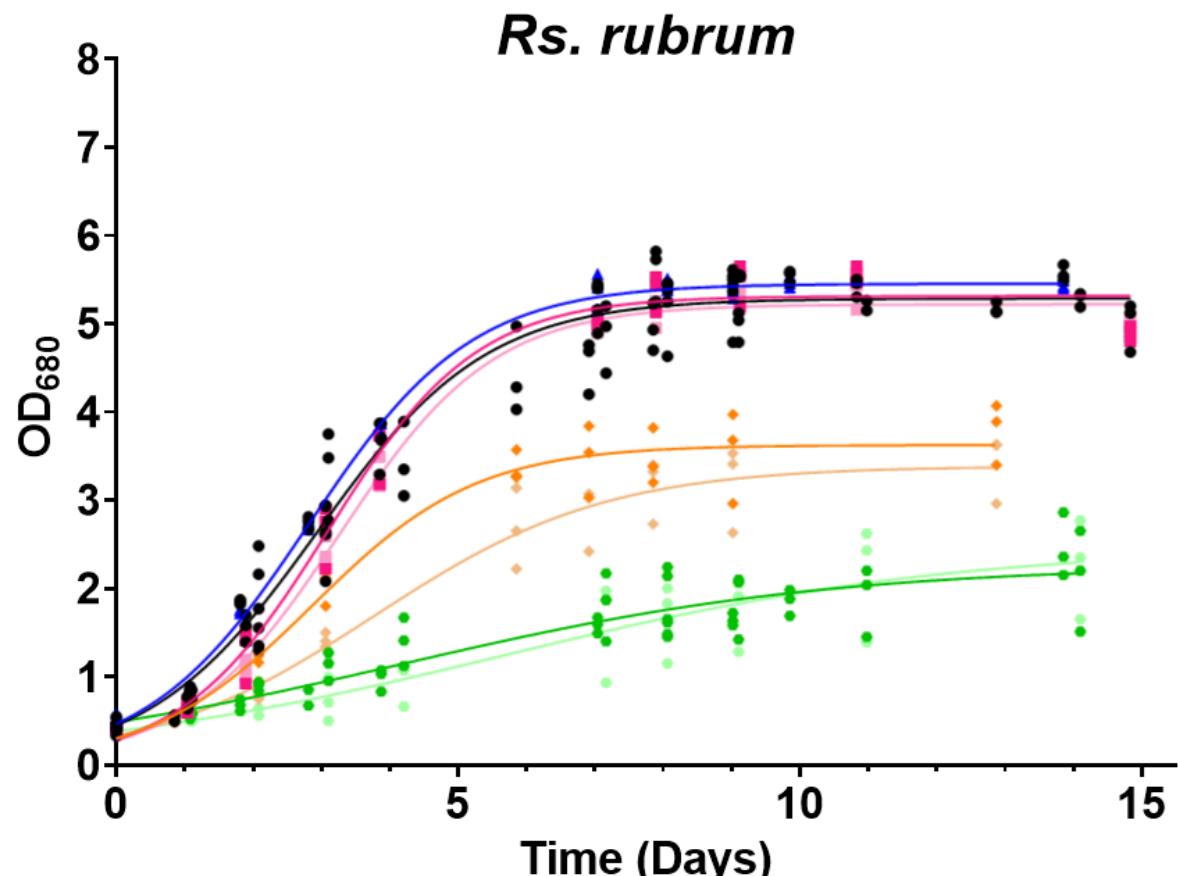
Reduced growth of co-cultures of *Rb. capsulatus*, *Rs. rubrum*, *C. sphaeroides* in high-ammonium medium

- ⇒ Why do the bacteria suffer in high-ammonium medium?
- ⇒ What is the minimal inhibitory concentration?
- ⇒ How can they adapt to these stringent conditions?

⇒ Organic acid contents, proteomic analysis and bacterial strain proportions will follow



Effects of High Ammonium Levels



⇒ Growth is impaired at 210 mM and 350 mM NH₄Cl and the cultures did not reach the same OD₆₈₀ as in lower NH₄Cl medium

⇒ Acclimation did not take place

⇒ Organic acid contents and proteomic analyses will follow

Acknowledgements



**Thank you for
your attention!**



This research is supported by the European fund for regional development (FEDER) through the WalBiopower project (DECARBOWAL portfolio) and Bioprofiling project cofinanced by EU and Wallonia.

