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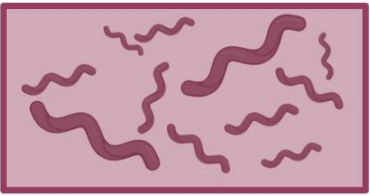
# Valorising organic waste for the production of high-value molecules by purple non-sulphur bacteria

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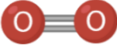
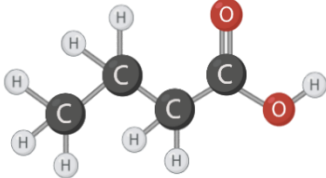





# Declaration of Interests

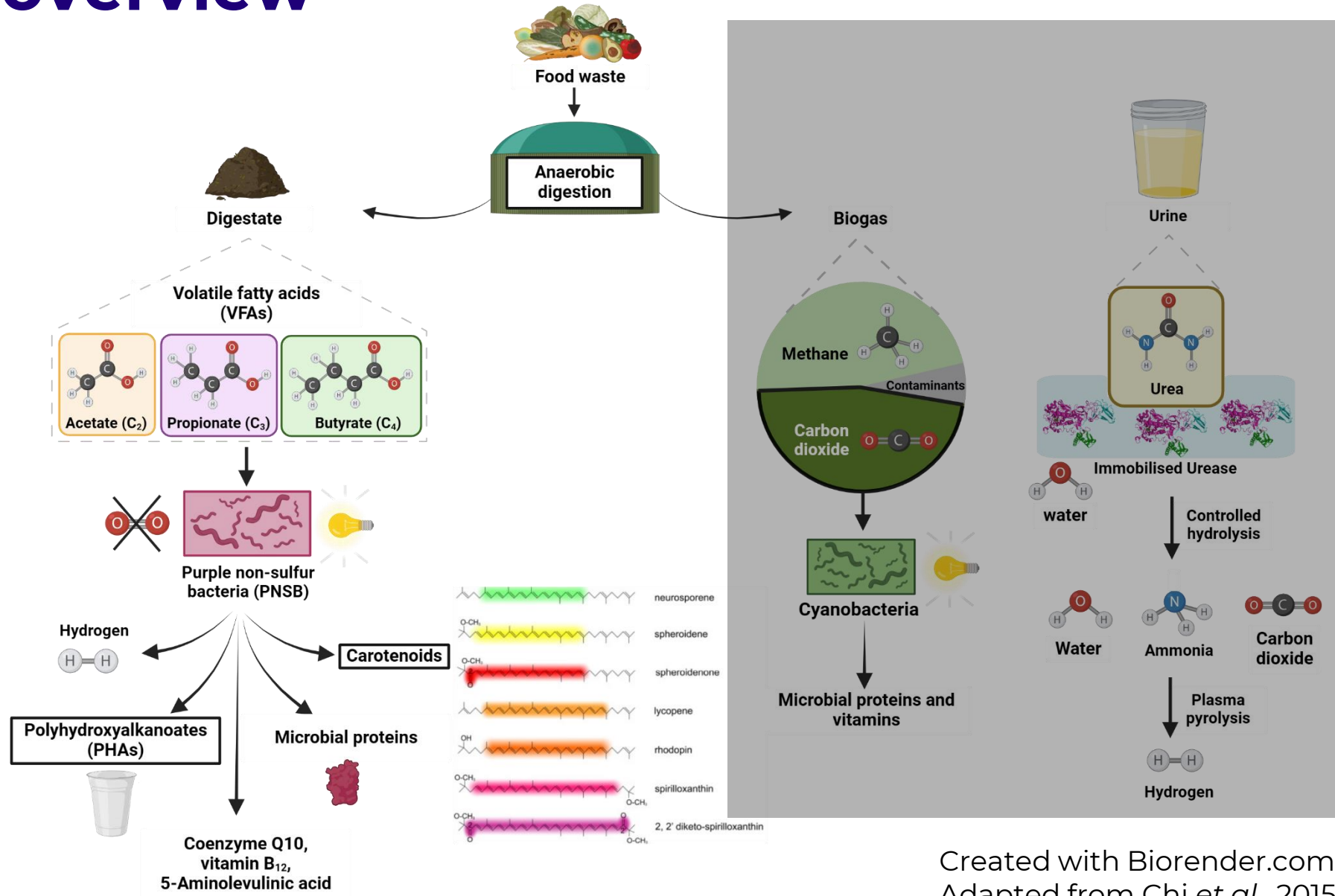
Affiliation / Financial interests	Organisation
Employment	Department of Proteomics and Microbiology, University of Mons
Grants / Research support	European fund for regional development (FEDER) 2021-2027 program under the WalBioPower project (DECARBOWAL portfolio) and the Bioprofiling project cofinanced by the EU and Wallonia
Scientific Advisory Board / Consultant	none
Government	none
Other	none



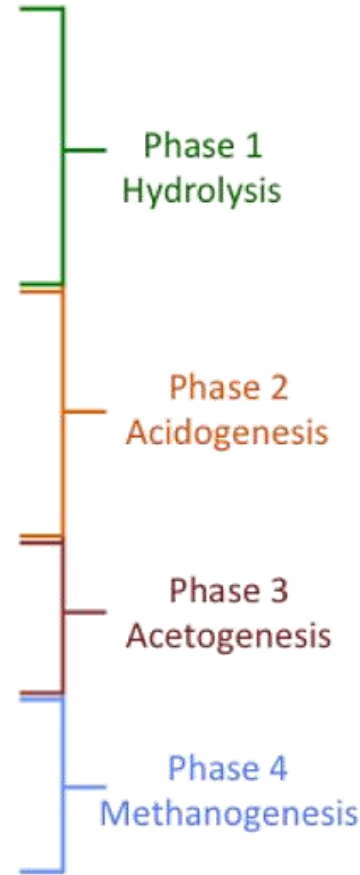
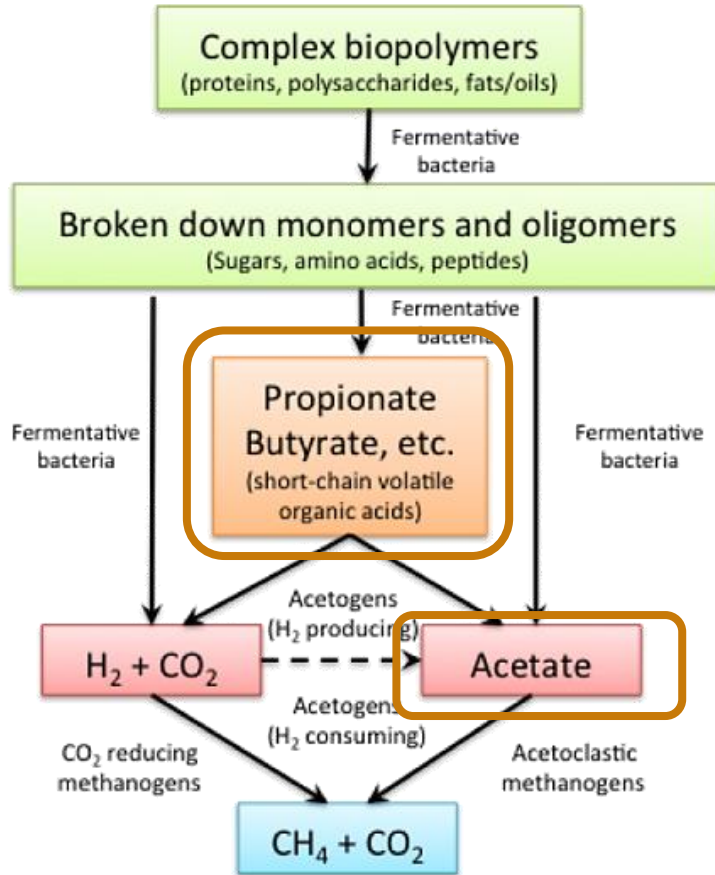
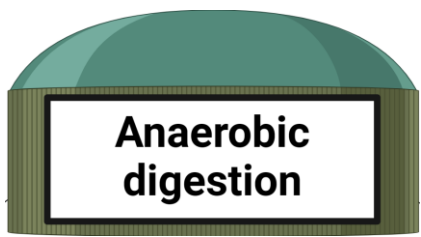
# Purple non-sulphur bacteria: Versatile metabolism

	 	 
C source: Organic compounds e- source: Organic compounds 	<b>Photoheterotrophy</b>	<b>Chemoheterotrophy</b>
C source:  e- source: Inorganic compounds  	<b>Photoautotrophy</b>	<b>Chemoautotrophy</b>

# Project overview



# Anaerobic digestion (AD)



- Waste streams
- Agriculture-based waste (manure and crops)
  - Wastewater
  - **Food** and municipal waste

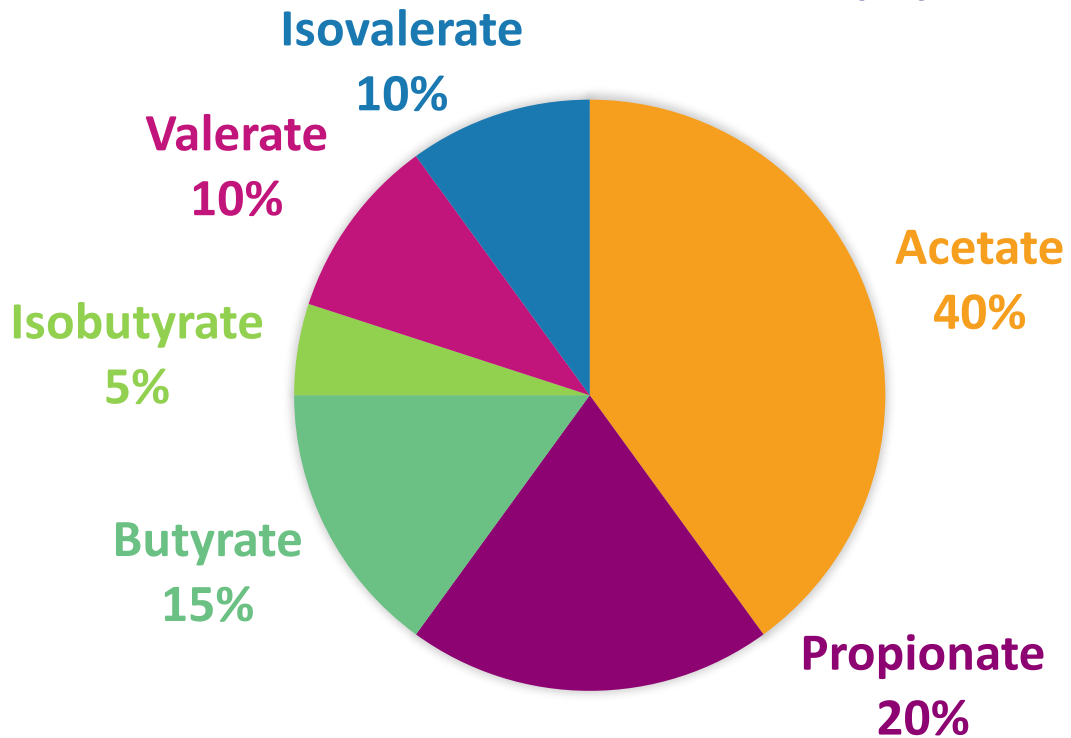
- Short-chain fatty acids (SCFAs) = Volatile fatty acids (VFAs)
- Acetate
  - Propionate
  - (Iso)butyrate
  - (Iso)valerate



Purple non-sulphur bacteria (PNSB)

# Culturing PNSB in Synthetic Digestate

## SYNTHETIC DIGESTATE (%)

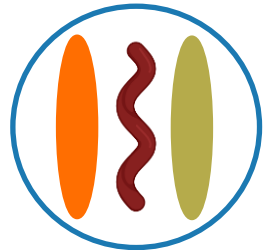


*Rhodospirillum rubrum*



Co-culture crs:

- *Rhodobacter capsulatus*
- *Rhodospirillum rubrum*
- *Cereibacter sphaeroides*

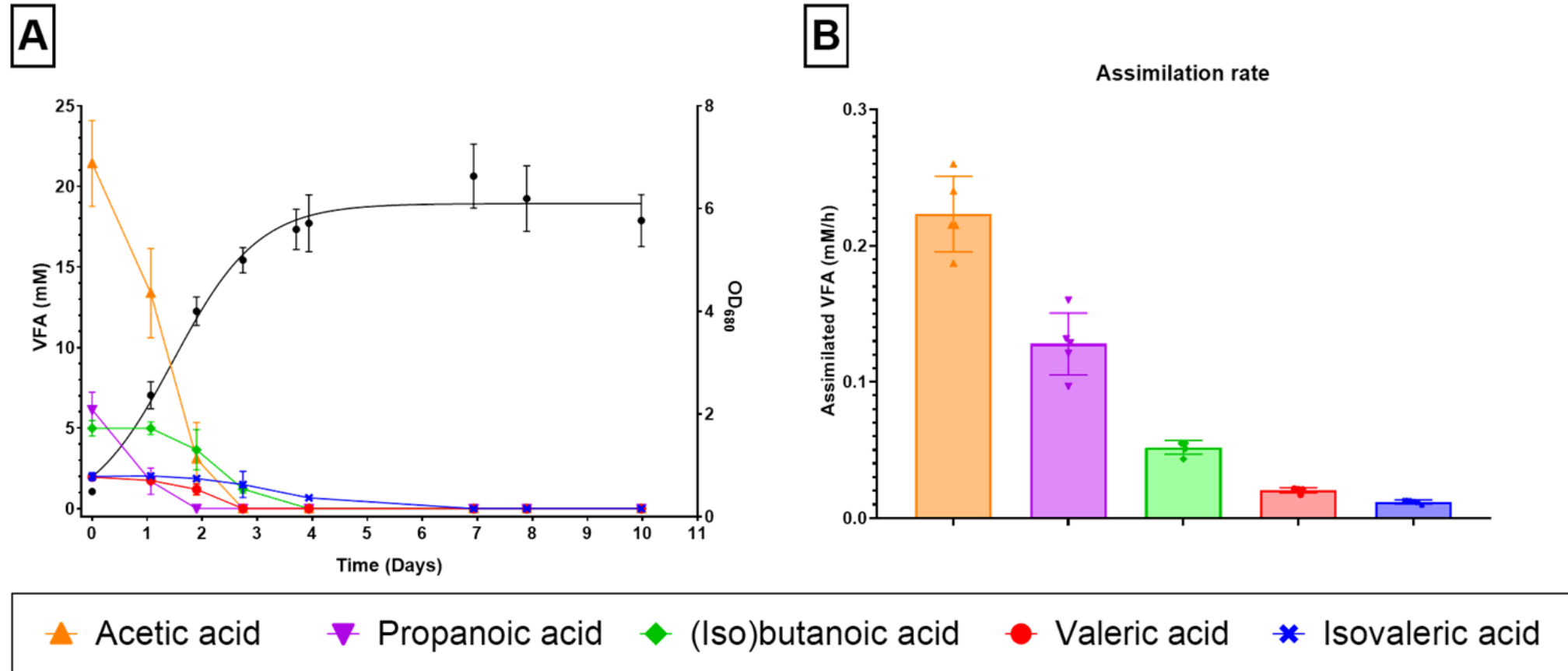


120 mM C equivalents



# Culturing PNSB in Synthetic Digestate

- LC-MS analysis of VFA assimilation





# Bacterial strain proportions

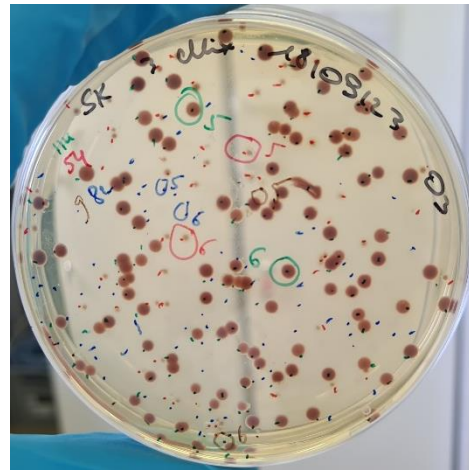
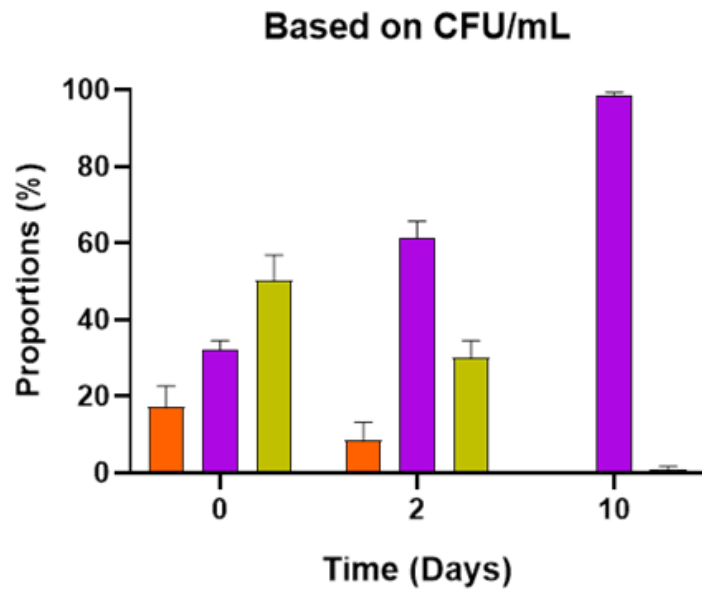
- $OD_{680} = 0.133$  of each strain  $\Rightarrow$  Achieve Start  $OD_{680} = 0.4$



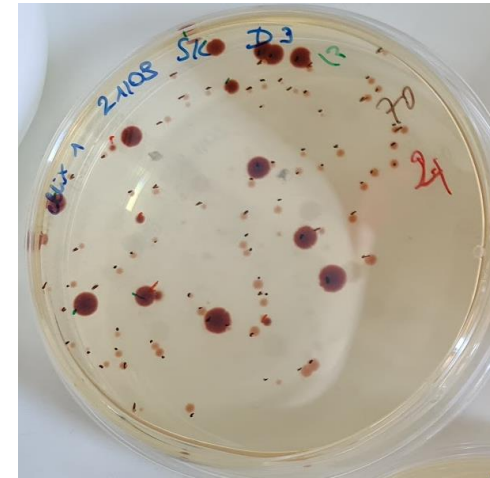


# Bacterial strain proportions

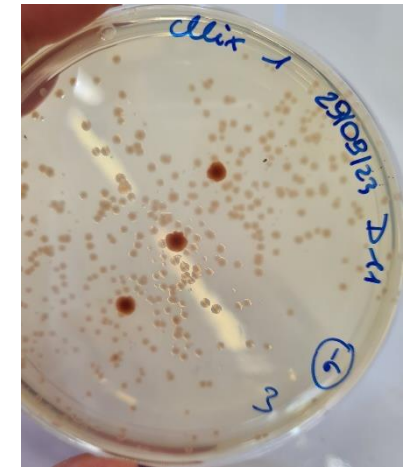
- $OD_{680} = 0.133$  of each strain  $\Rightarrow$  Achieve Start  $OD_{680} = 0.4$



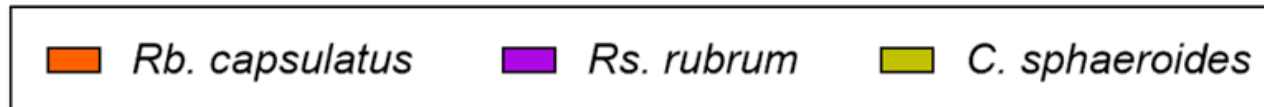
Day 0



Day 2



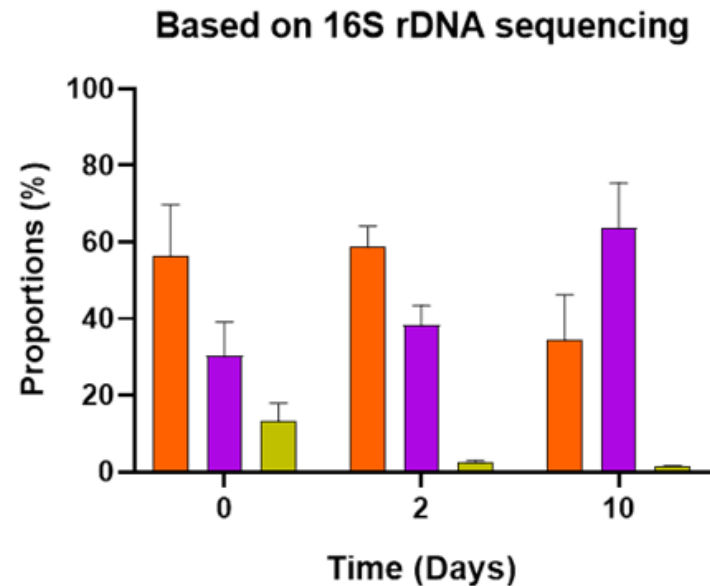
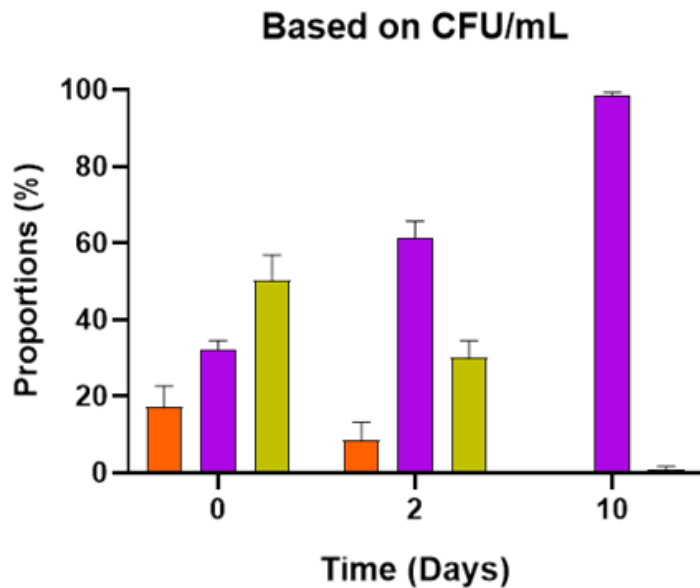
Day 10



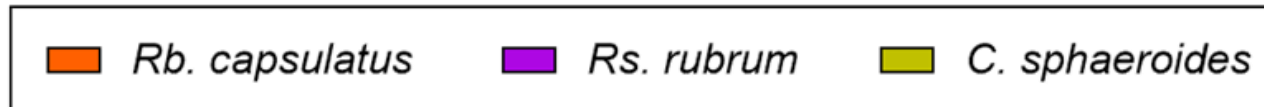


# Bacterial strain proportions

- $OD_{680} = 0.133$  of each strain  $\Rightarrow$  Achieve Start  $OD_{680} = 0.4$



Oxford Nanopore GridION  
+ 16S rDNA primers from  
Oxford Nanopore

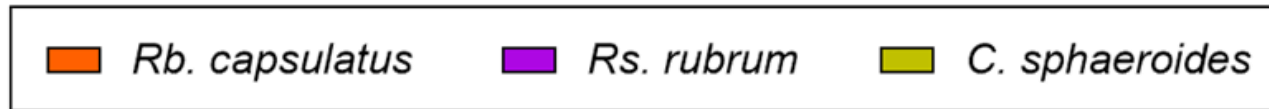
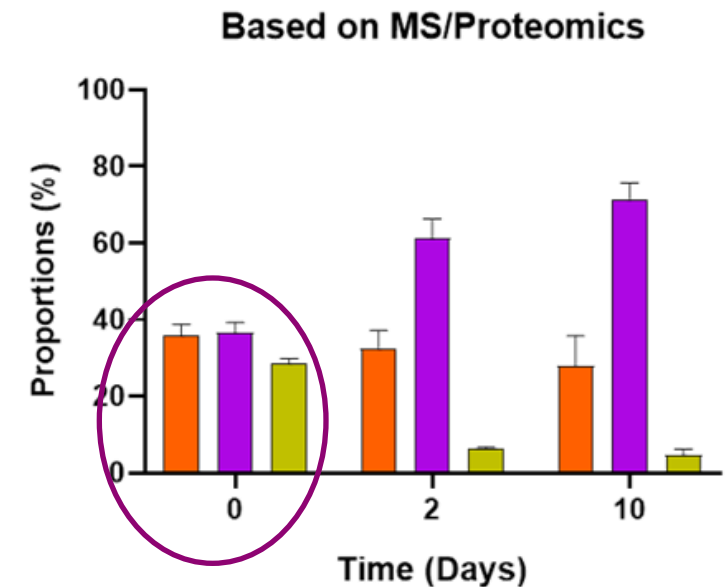
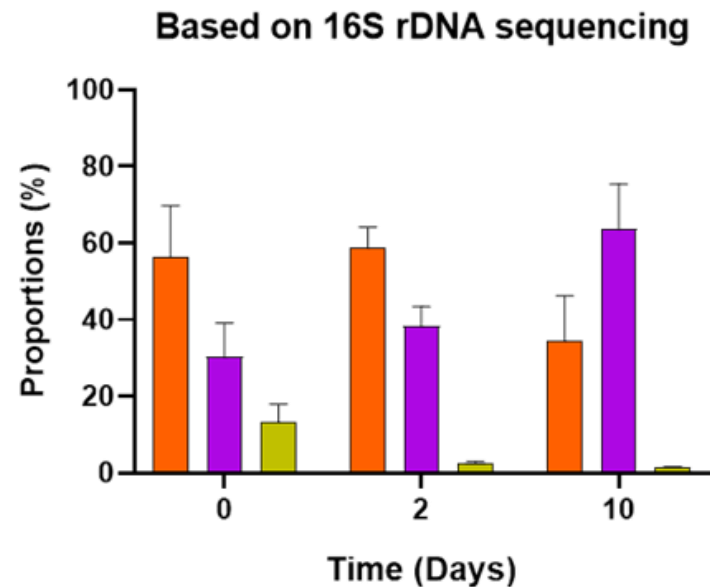
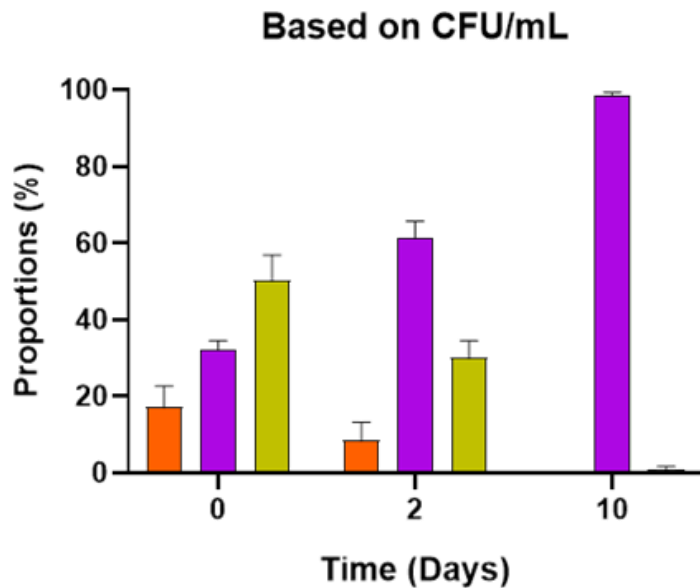


Primers have a higher affinity for *Rb. capsulatus*



# Bacterial strain proportions

- $OD_{680} = 0.133$  of each strain  $\Rightarrow$  Achieve Start  $OD_{680} = 0.4$



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

# Ammonium in Food-waste Digestates

Standard MELiSSA Medium:  
 35 mM NH<sub>4</sub>Cl  
 → 0.5 g/L N  
 → 0.635 g/L NH<sub>4</sub><sup>+</sup>

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 <sub>4</sub> <sup>+</sup> -N (g/kg)	C/N ratio	References
		96.4	8.9	10.8	91.0	43.5	9.6	2.4	2.3	7.2	20.9	
		85.3	8.1	5.2	63.6	32.1	4.6	1.0	1.3	4.0	11.3	
		69.8	7.3	0.8	38.5	12.8	1.1	0.1	0.4	1.5	3.1	
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 Drogg (2013)
Food waste	Germany	-	7.6-8.1	-	-	-	3-6.8	-	-	1.5-5.6	-	Fuchs and Drogg (2013)
Food 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

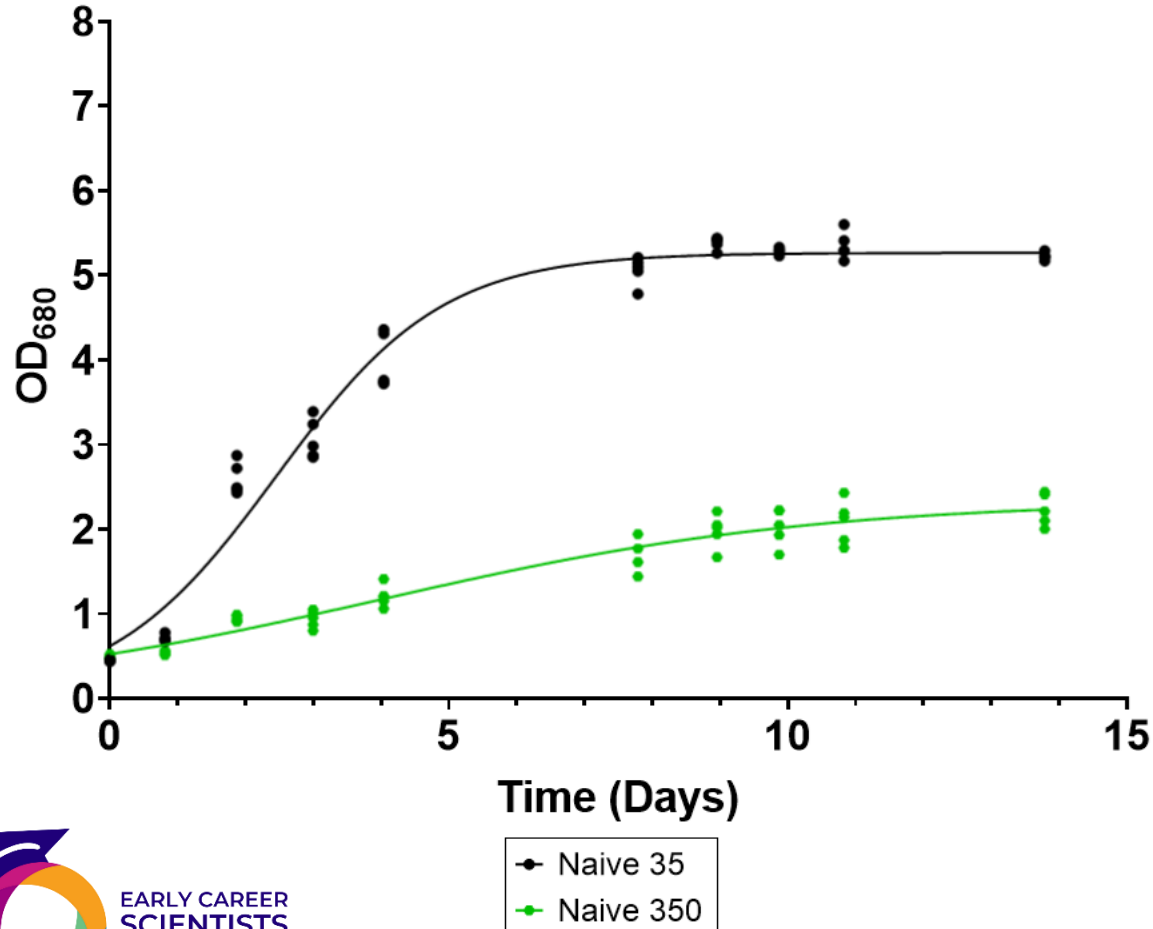
~350 mM NH<sub>4</sub>Cl  
 → 5 g/L N  
 → 6.35 g/L NH<sub>4</sub><sup>+</sup>





# Effects of High Ammonium Levels

## Co-culture crs



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

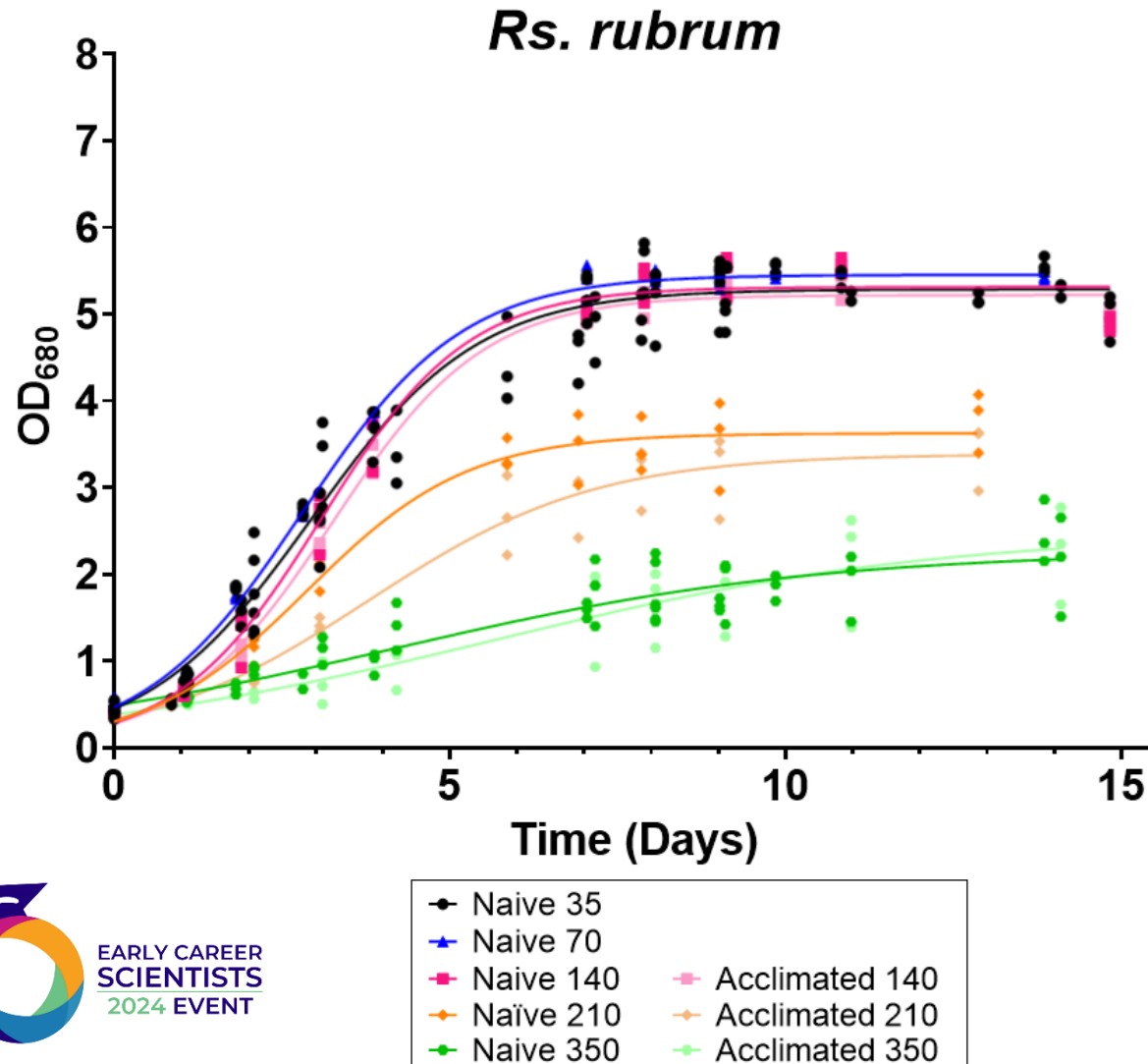
⇒ *Why do the bacteria suffer in high-ammonium medium?*

⇒ *How could they adapt to these stringent conditions?*

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



# Effects of High Ammonium Levels



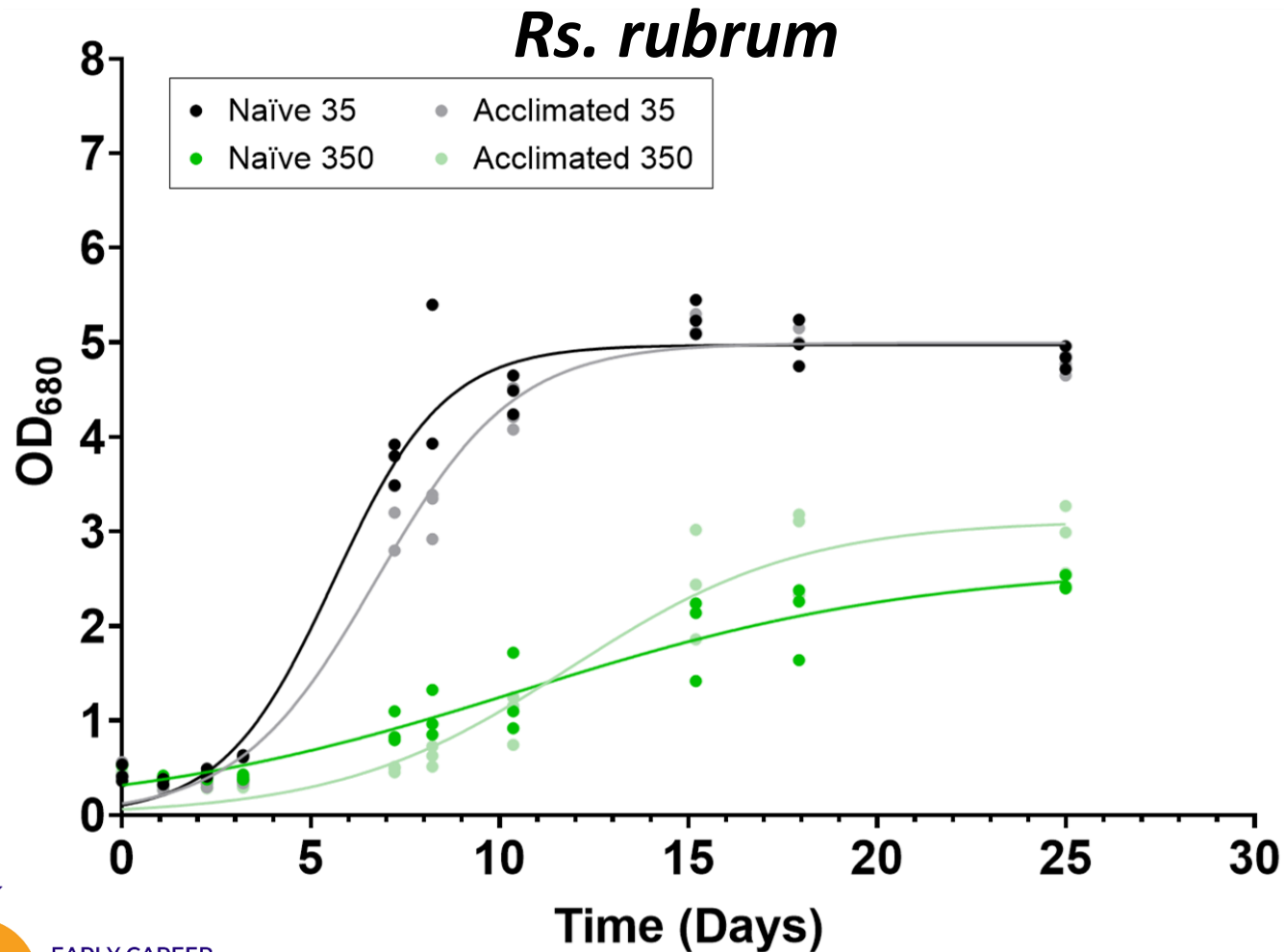
⇒ Growth is impaired at 210 mM and 350 mM NH<sub>4</sub>Cl and the cultures did not reach the same OD<sub>680</sub> as in lower NH<sub>4</sub>Cl medium

⇒ Acclimatation did not take place

⇒ *Organic acid contents and proteomic analyses will follow*



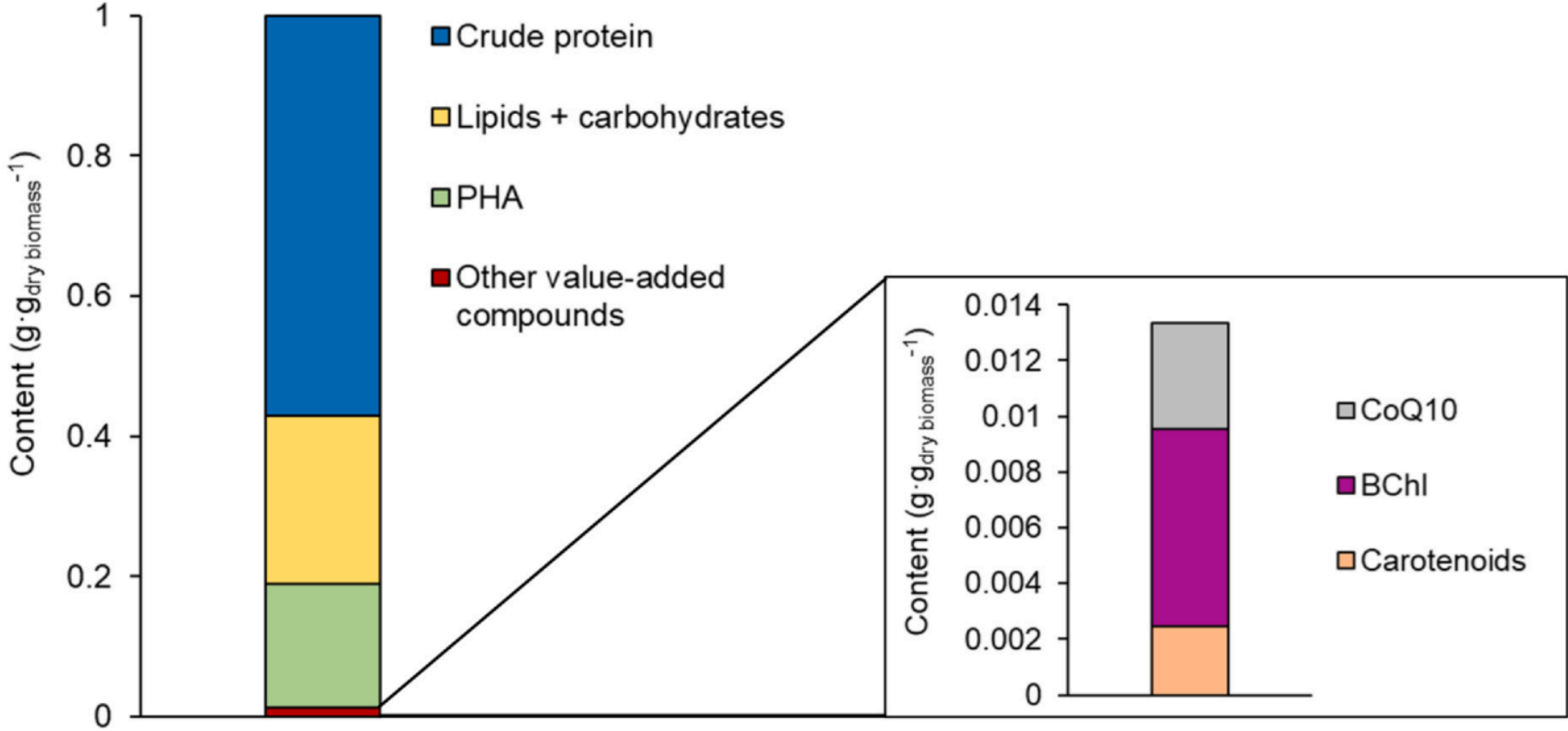
# Acclimatation to High Ammonium Levels



⇒ Is the acclimatation taking place at the moment?

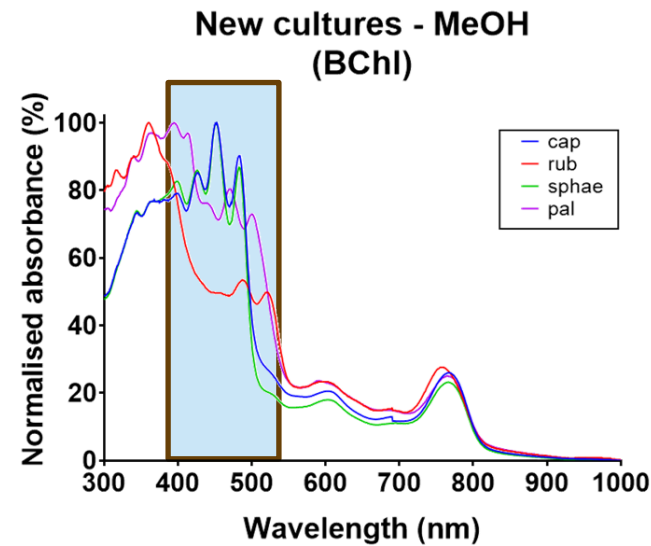
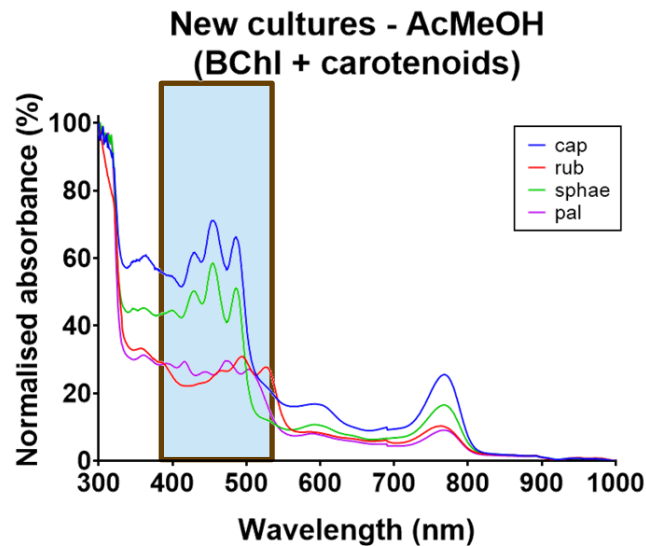
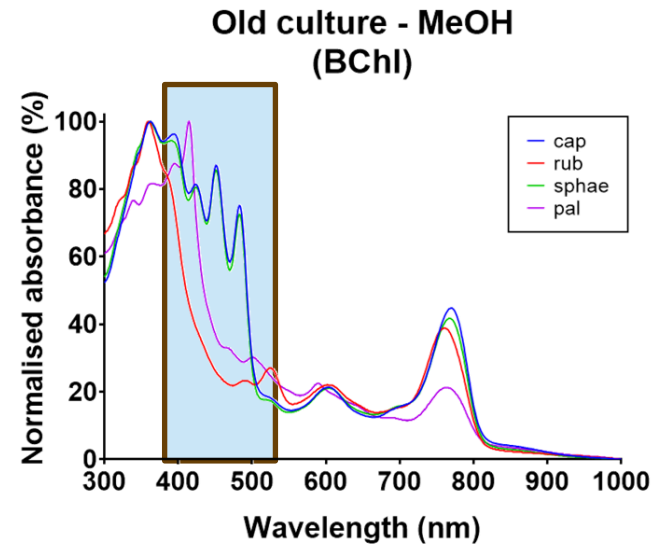
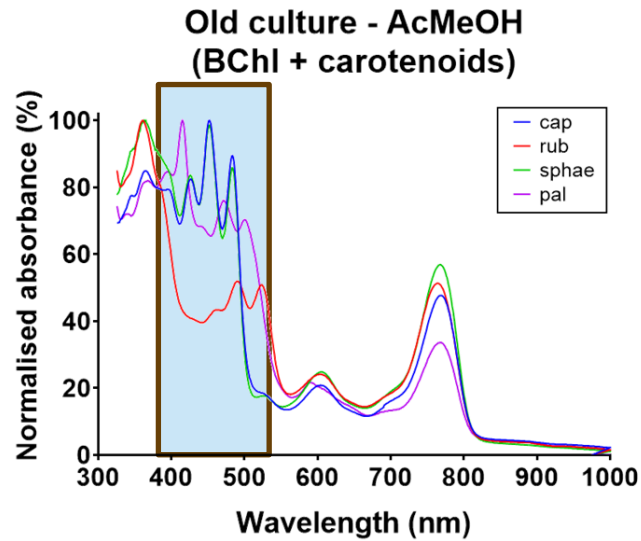
⇒ *Organic acid contents and proteomic analyses will follow*

# Contents of PNSB

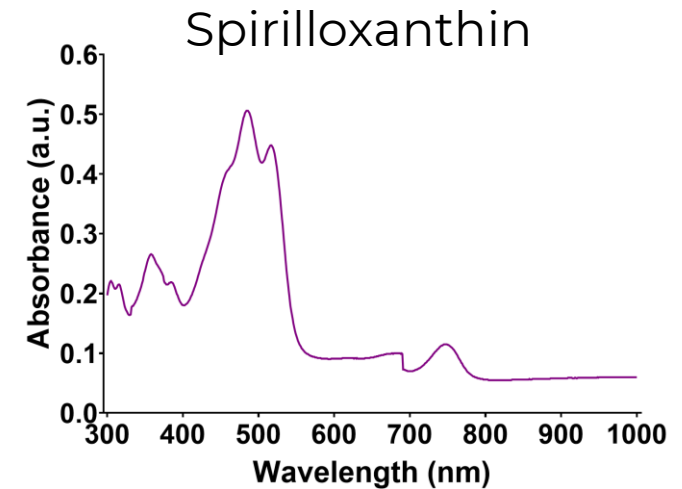




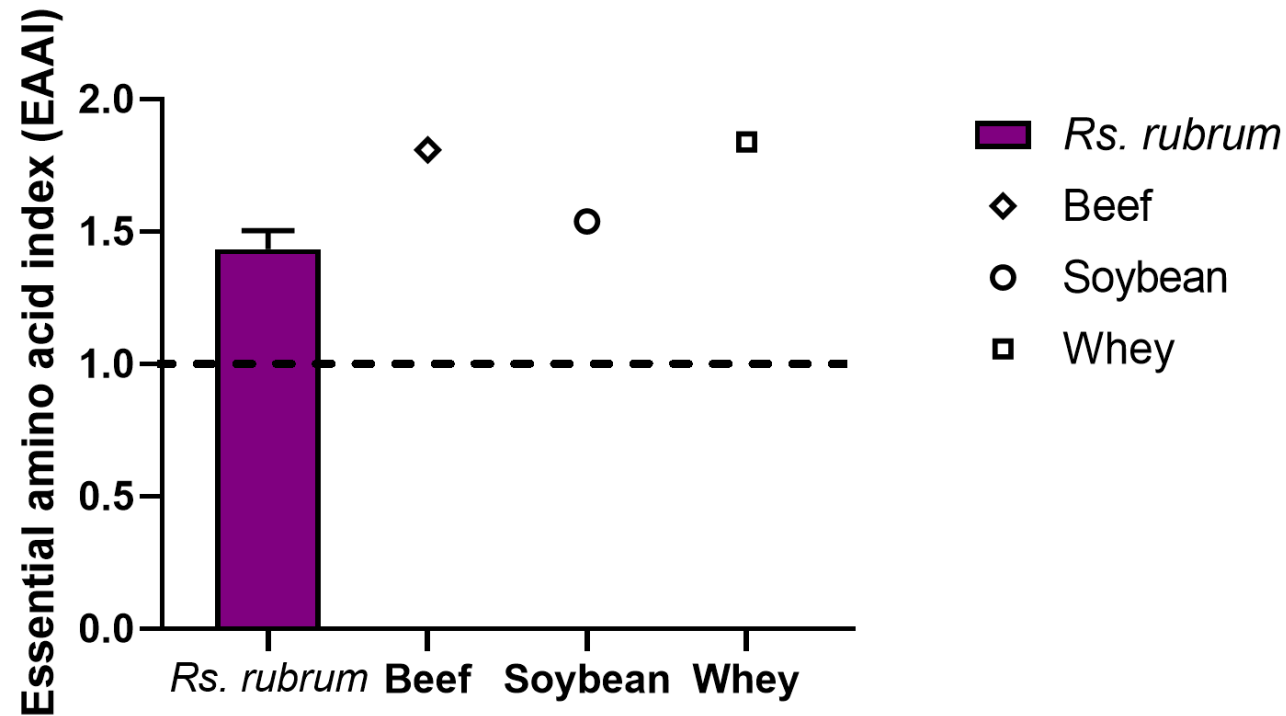
# Carotenoid extractions



*Currently testing  
the antioxidant effects  
of carotenoids  
on walnut oil  
(UCLouvain)*



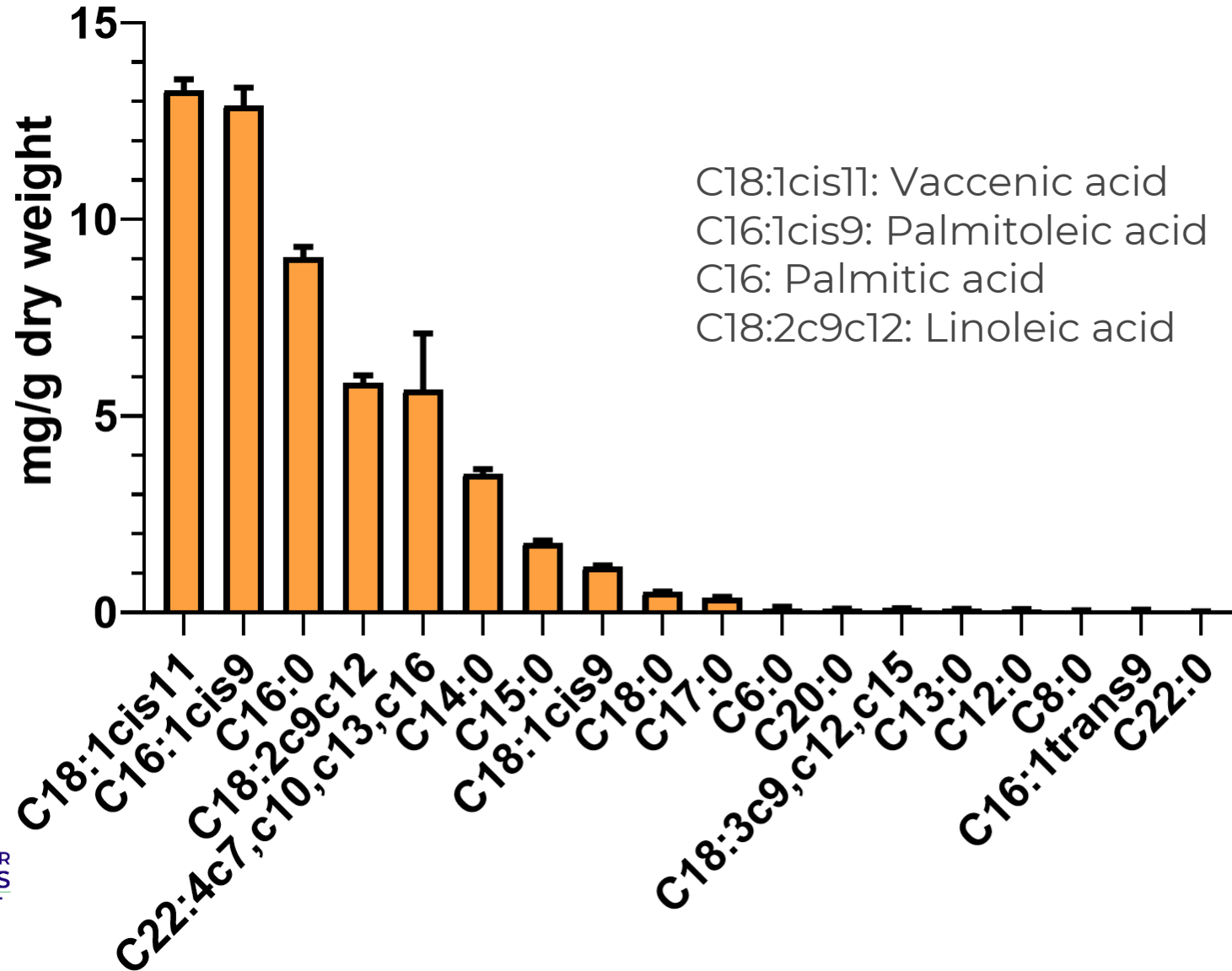
# Protein-rich part of *Rhodospirillum rubrum*



Conversion of molasses into protein-rich *Rhodospirillum rubrum* biomass

Essential amino acid contents are similar to other protein sources

# Fatty acids in *Rhodospirillum rubrum*



Many *cis* fatty acids

# Take-home messages

- **Co-cultures** of *Rhodospirillum rubrum*, *Rhodobacter capsulatus* and *Cereibacter sphaeroides* assimilate **acetic and propionic acid** first, followed by **(iso)butyric acid and valeric acid** and lastly, **isovaleric acid**
- **Mass spectrometry** is a great alternative method to determine **bacterial proportions** in mixed cultures
- ***Rhodospirillum rubrum*** is able to survive to **high ammonium** levels of 140 mM  $\text{NH}_4\text{Cl}$  (= **2.54 g/L  $\text{NH}_4^+$** ) without reduction in growth
- ***Rhodospirillum rubrum*** is able to produce **microbial proteins (containing essential amino acids)** and **cis fatty acids**, which **could have health benefits**

# Acknowledgements



This research is supported by the European fund for regional development (FEDER) 2021-2027 program under the WalBioPower project (DECARBOWAL portfolio) and the Bioprofiling project cofinanced by the EU and Wallonia



# Thank you!



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