

Cold-adapted carboxylic ester hydrolases from two Antarctic *Psychrobacter* strains: genomic analyses and *in-vitro* studies



PNRA19_00073 (TENORE)
PNRA18_00232 (AMICI)



Ca' Foscari
University
of Venice



Institute of
Biomolecular
Chemistry

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Sampling campaigns in Antarctica

Environmental samples

Genomic approach

Study of bacterial communities

Culturomic approach

In-lab isolation of extremophilic bacteria

Taxonomical characterization and study of biodiversity

Study of biomolecules

Polymers

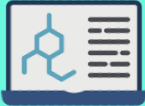
e.g. exo- and capsular-polysaccharides, polyglutamic acid

Enzymes

e.g. carboxylic ester hydrolases (EC 3.1.1), glycoside hydrolases (EC 3.2.1)

Genome sequencing and annotation

OVERVIEW OF THE PhD PROJECT

 **In-plate screening**  **In-silico screening**

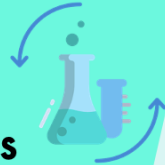
Recovery and purification

In-vitro assay



Definition of the biotechnological potential

Set-up to monitor the degradation of plastics



Optimization of the production

Taxonomical characterization and study of biodiversity

Study of biomolecules

Polymers

e.g. exo- and capsular-polysaccharides, polyglutamic acid

Enzymes

e.g. **carboxylic ester hydrolases (EC 3.1.1)**, glycoside hydrolases (EC 3.2.1)

Genome sequencing and annotation

ENVIRONMENTAL SAMPLING IN ANTARCTICA

XXXIII Italian Antarctic expedition (2017/2018)

XXXIV Italian Antarctic expedition (2018/2019)

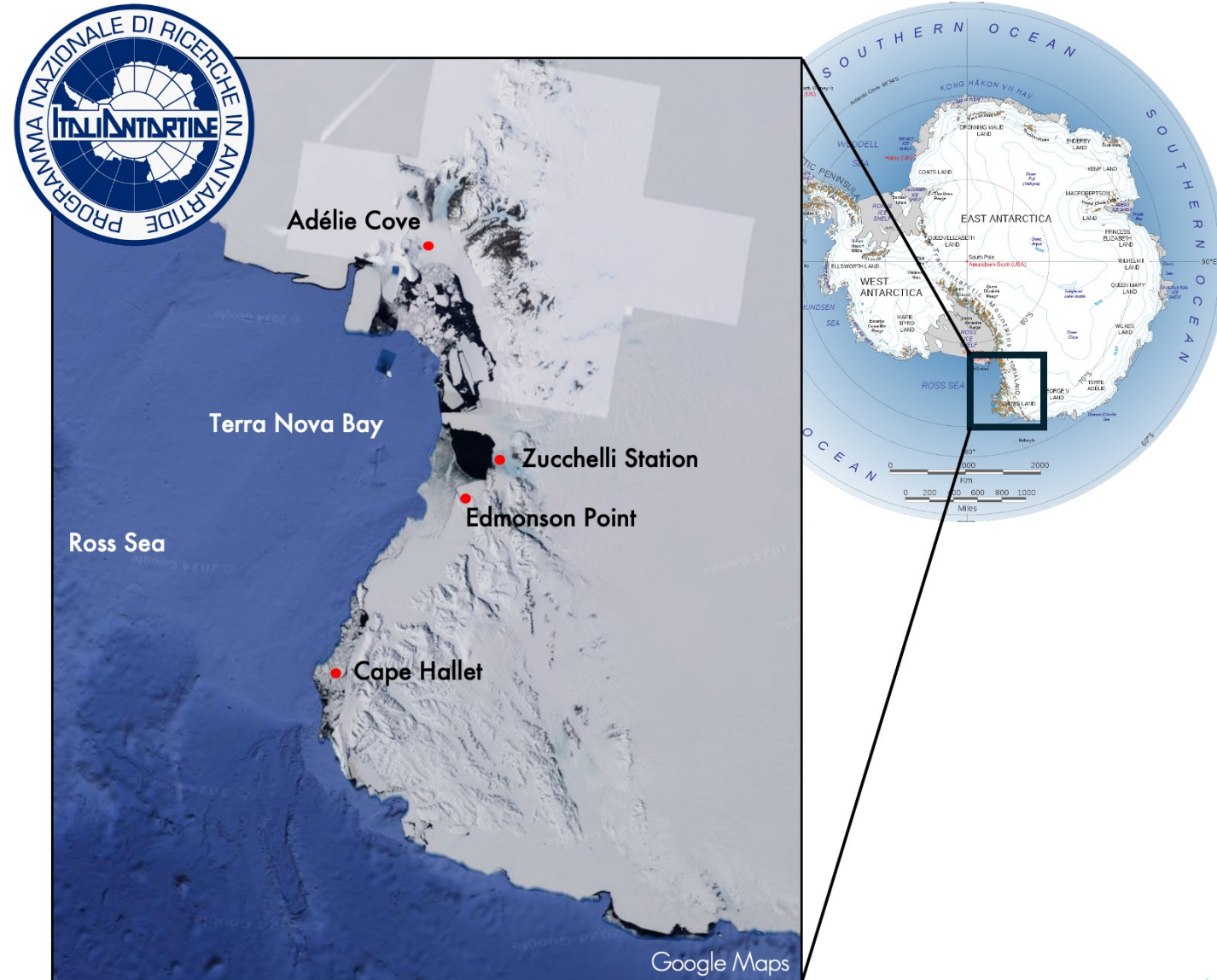
Supported by Research Project PNRA16_00274

- Terra Nova Bay, in proximity of the Mario Zucchelli Station and the Adélie Cove Bay (ASPA 161)
Marine sediment samples from 20 and 50 m depth
- Edmonson Point (ASPА 165)
Sand and sediment samples
- Cape Hallet (ASPА 106)
Sand and sediment samples

XXXIX Italian Antarctic expedition (2023/2024)

Supported by Research Project PNRA19_00073

- Ross Sea polynya with Laura Bassi Research Vessel
Sea water samples



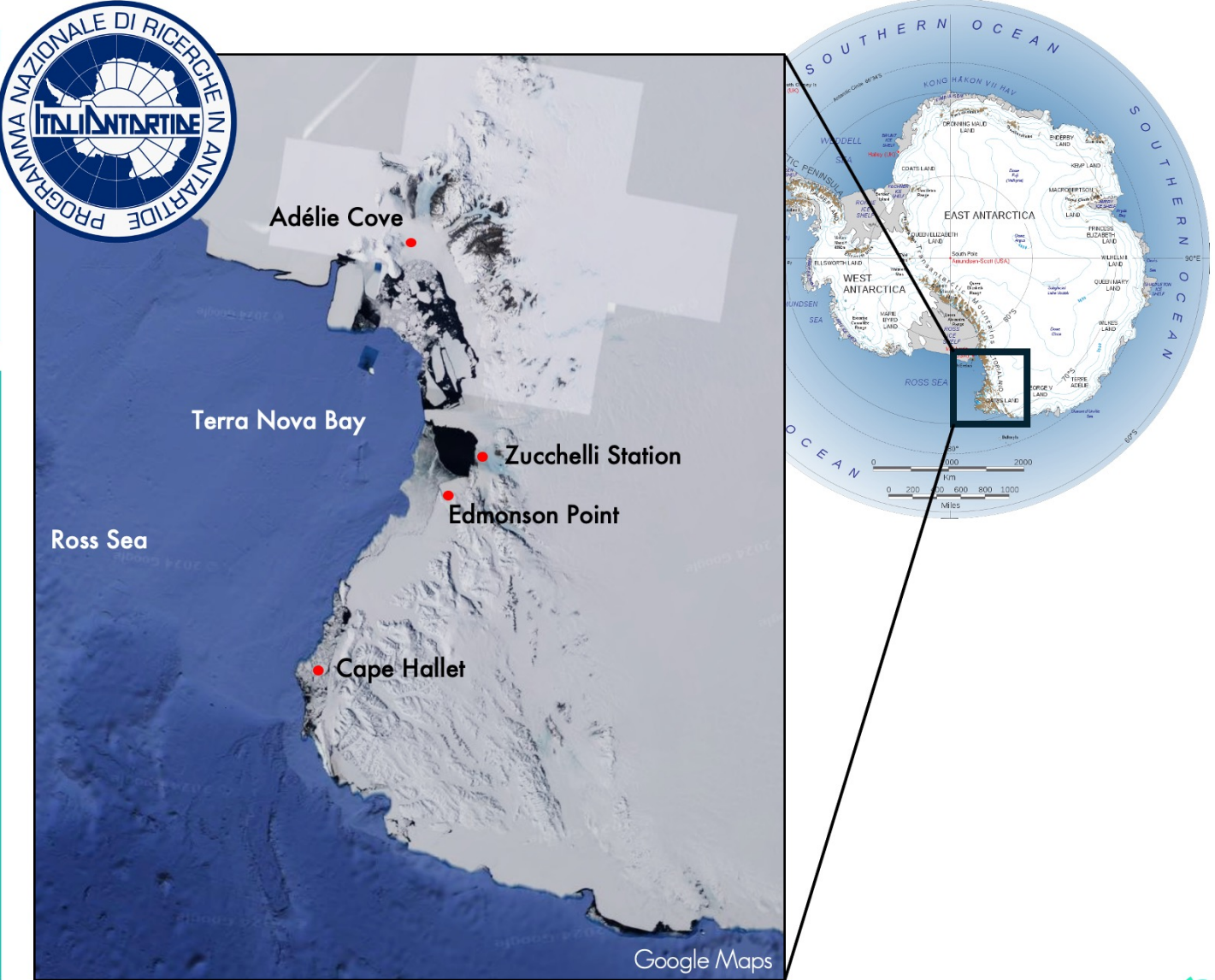
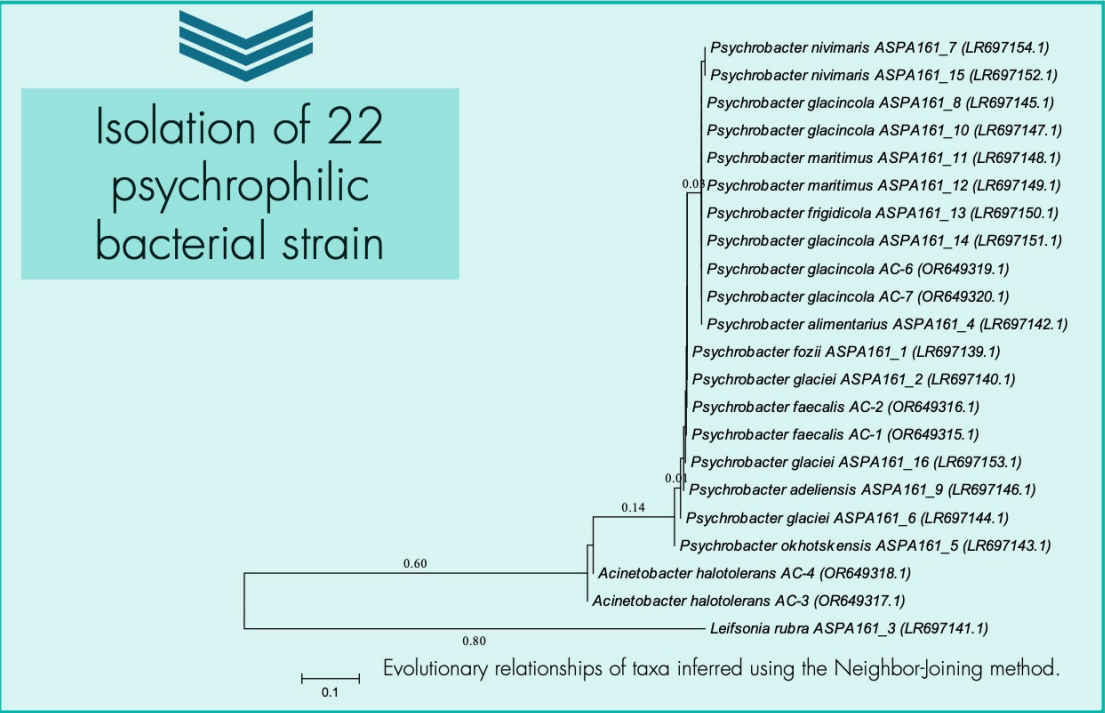
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STUDY OF EXTREMOPHILIC BACTERIA: ICB CULTURE COLLECTION

Culturomic approach

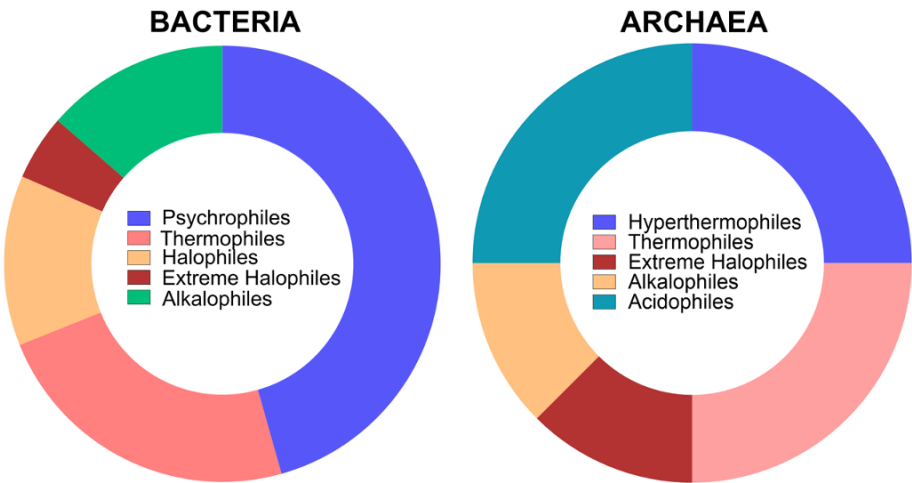
CE-ICB Extremophiles Collection

Institute of Biomolecular Chemistry – National Research Council (Napoli, IT)
Collection, characterization, distribution and preservation of microbial strains

SUS-MIRRI.IT project, funded by NextGeneration EU 2022



More information @ our Posters



Psychrobacter sp. strain ASPA161_6

Psychrobacter sp. strain ASPA161_9

Temperature	Medium	pH	Salinity
10-30 °C, optimum 20 °C	Tryptic Soy Broth	6.0-8.5	0.5-5%
10-30 °C, optimum 20 °C	Tryptic Soy Broth	6.0-9.0	0.5-5%

STUDY OF EXTREMOPHILIC BACTERIA: ICB CULTURE COLLECTION

Culturomic approach

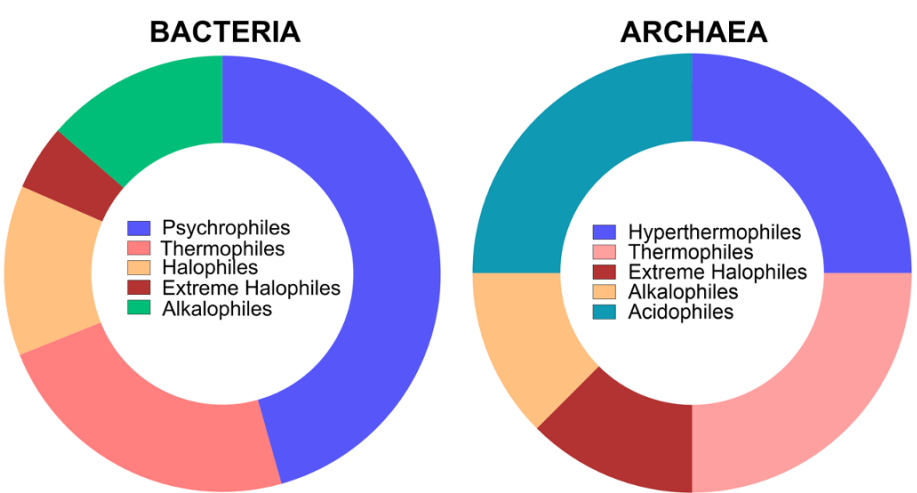
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Genome sequencing: dDDH and ANI

Psychrobacter sp. strain ASPA161_6

Accession number JBEFNJ0000000000

Genome compared	Access. number	dDDH [%confidence interval]	ANiB [%aligned nucleotide]	ANIm [%aligned nucleotide]
<i>Psychrobacter glaciei</i> KCTC 42280	GCA_014652895	67.7 [64.8-70.6]	95.72 [82.56]	96.37 [84.12]
<i>Psychrobacter okhotskensis</i> MD17	GCF_904846405	30.2 [27.8-32.7]	85.06 [66.87]	87.97 [60.20]
<i>Psychrobacter fjordensis</i> BSw21516B	GCF_904845995	25.7 [23.4-28.2]	81.01 [59.89]	86.97 [33.89]
<i>Psychrobacter fozii</i> CECT 5889	GCA_003217155.1	25.5 [23.2-28.0]	81.09 [60.76]	86.82 [35.29]

Psychrobacter sp. strain ASPA161_9

Accession number JBEFNI0000000000

<i>Psychrobacter okhotskensis</i> MD17	GCF_904846405	28.8 [22.5-27.3]	79.69 [55.73]	86.29 [26.66]
<i>Psychrobacter fulvigenes</i> KC40	GCF_904846155	24.3 [22.0-26.7]	79.91 [57.12]	85.63 [29.12]
<i>Psychrobacter maritimus</i> Pi2-20	GCF_904846295	24.2 [21.9-26.7]	79.44 [54.22]	85.99 [24.65]
<i>Psychrobacter luti</i> CECT 5885	GCF_014192115	23.9 [21.6-26.4]	79.18 [52.68]	85.74 [23.20]

Genome annotation
Search for **cold-adaptation and stress responses** genes

Gene	COG	Product	EC number
cspA_1	COG1278	Cold shock protein	/
cspA_2	COG1278	Cold shock protein	/
cspV	COG1278	Cold shock protein	/
deaD	COG0513	ATP-dependent RNA helicase	3.6.4.13
desA3	COG3239	NADPH-dependent stearyl-CoA 9-desaturase	1.14.19
katE	COG0753	Catalase C	1.11.1.6
katG	COG0336	Catalase-peroxidase	1.11.1.21
sodA	COG0605	Superoxide dismutase [Mn/Fe]	1.15.1.1
sodB	COG0605	Superoxide dismutase [Fe]	1.15.1.1
msrAB	COG0225	Peptide methionine sulfoxide reductase	1.8.4.12

Genome annotation

Search for carboxylic ester hydrolases: **lipase (EC 3.1.1.3)** and **esterase (EC 3.1.1.1)**

***Psychrobacter* sp.
strain ASPA161_6**

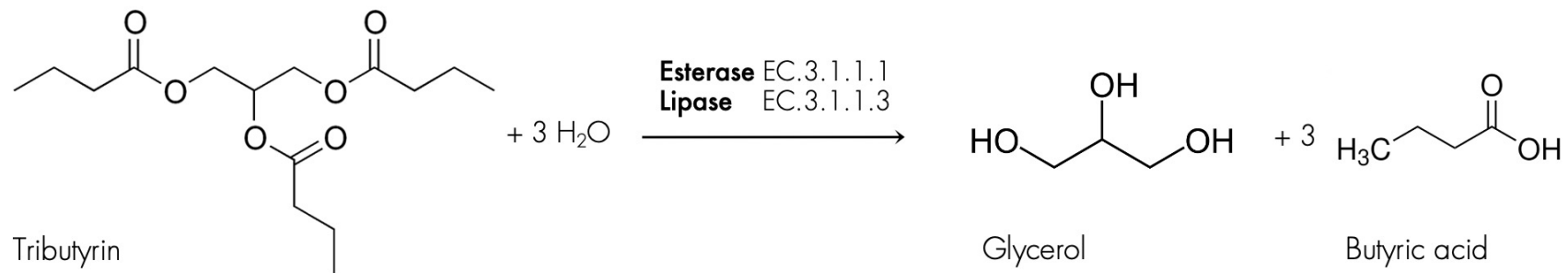
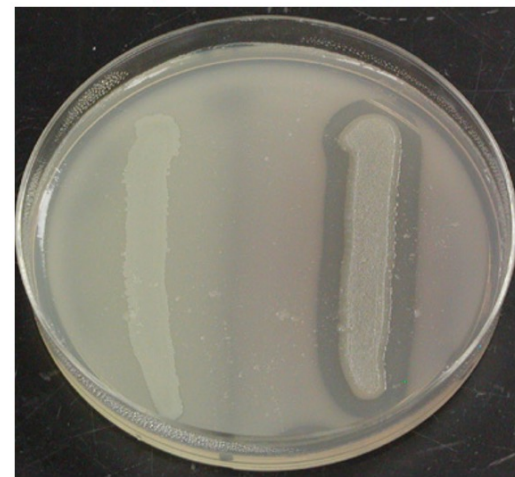
Gene	Length (bp)	COG	Product	EC number
estB	672	COG0400	Carboxylesterase 2	3.1.1.1
lip1	987	COG1075	Lipase 1	3.1.1.3
lip2	1302	COG1075	Lipase 2	3.1.1.3
lip3	945	COG1075	Lipase 3	3.1.1.3
lip	1077	COG1075	Triacylglycerol lipase	3.1.1.3

***Psychrobacter* sp.
strain ASPA161_9**

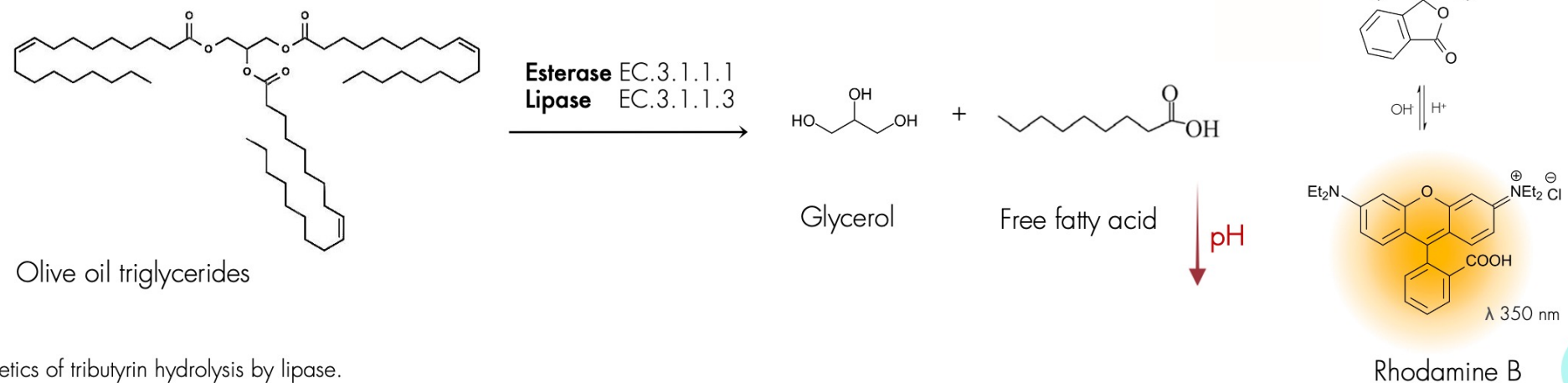
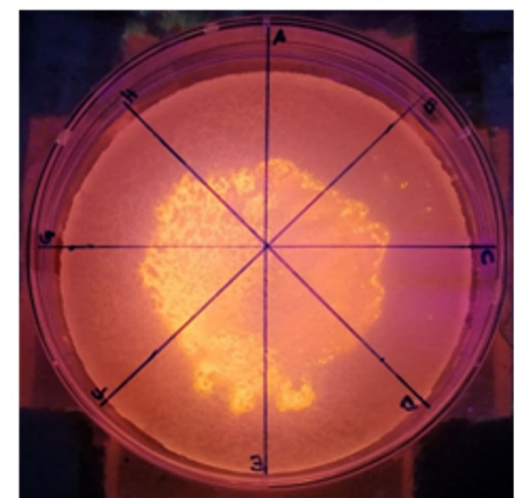
estB	672	COG0400	Carboxylesterase 2	3.1.1.1
lip2	1548	COG1075	Lipase 2	3.1.1.3
lip3	948	COG1075	Lipase 3	3.1.1.3
lip	1053	COG1075	Triacylglycerol lipase	3.1.1.3

IN-PLATE SCREENING FOR POLYESTER HYDROLASES

Tributyrin 1% (v/v) agar plate

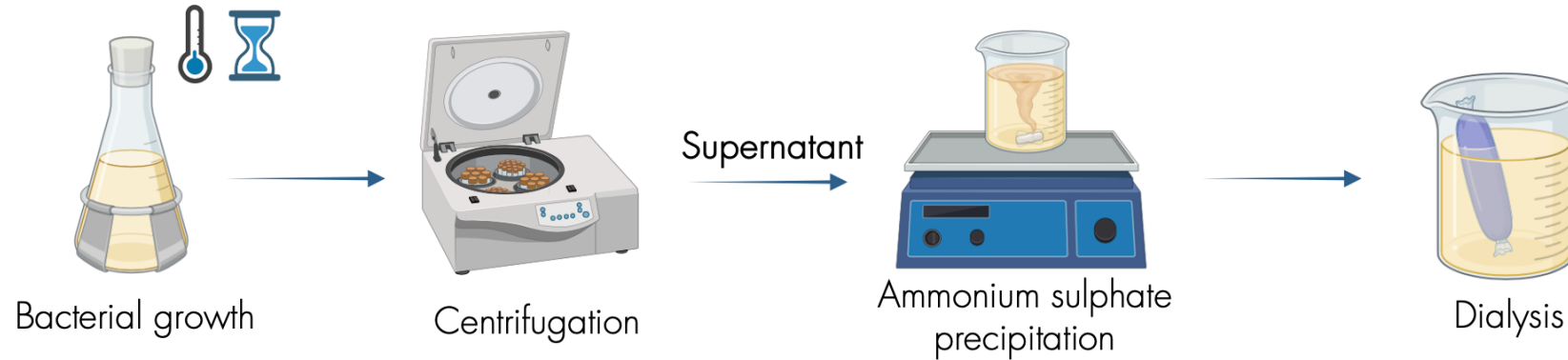


Olive oil 2.5% (v/v) and rhodamine B agar plate



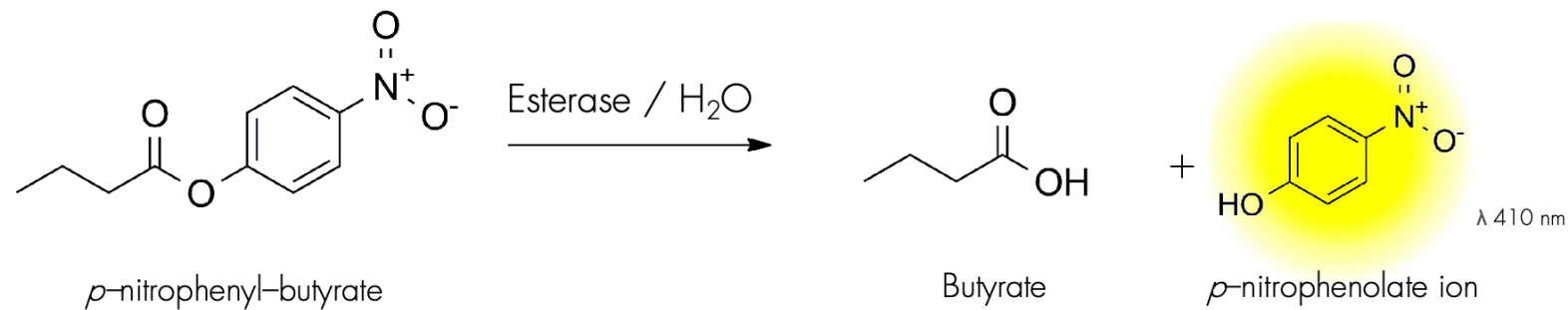
IN-VITRO ASSAY: COLORIMETRIC-SPECTROPHOTOMETRIC ASSAYS

Recovery of the extracellular enzymatic activities in the partially purified supernatant



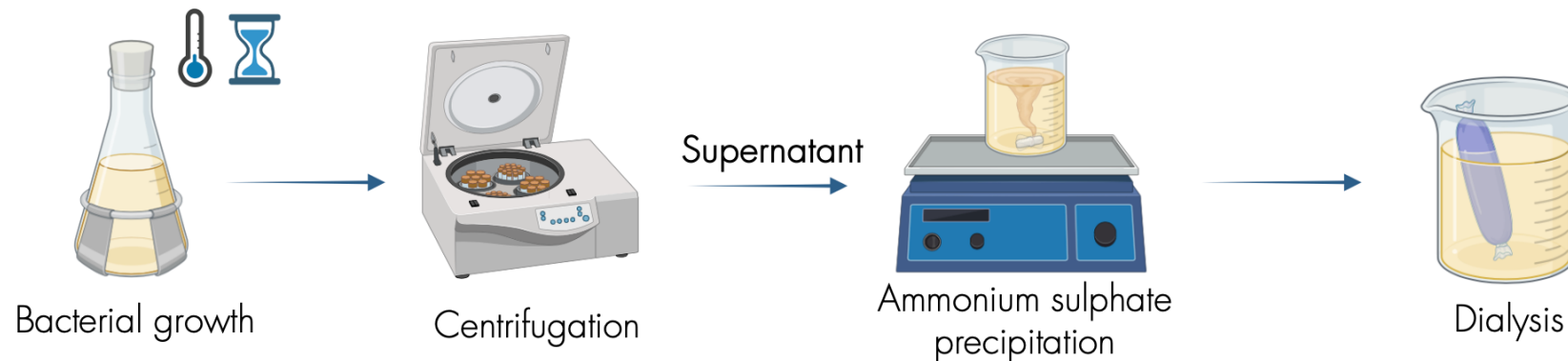
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Colorimetric-spectrophotometric assays with chromogenic substrates: p-NP esters



IN-VITRO ASSAY: COLORIMETRIC-SPECTROPHOTOMETRIC ASSAYS

Recovery of the extracellular enzymatic activities in the partially purified supernatant



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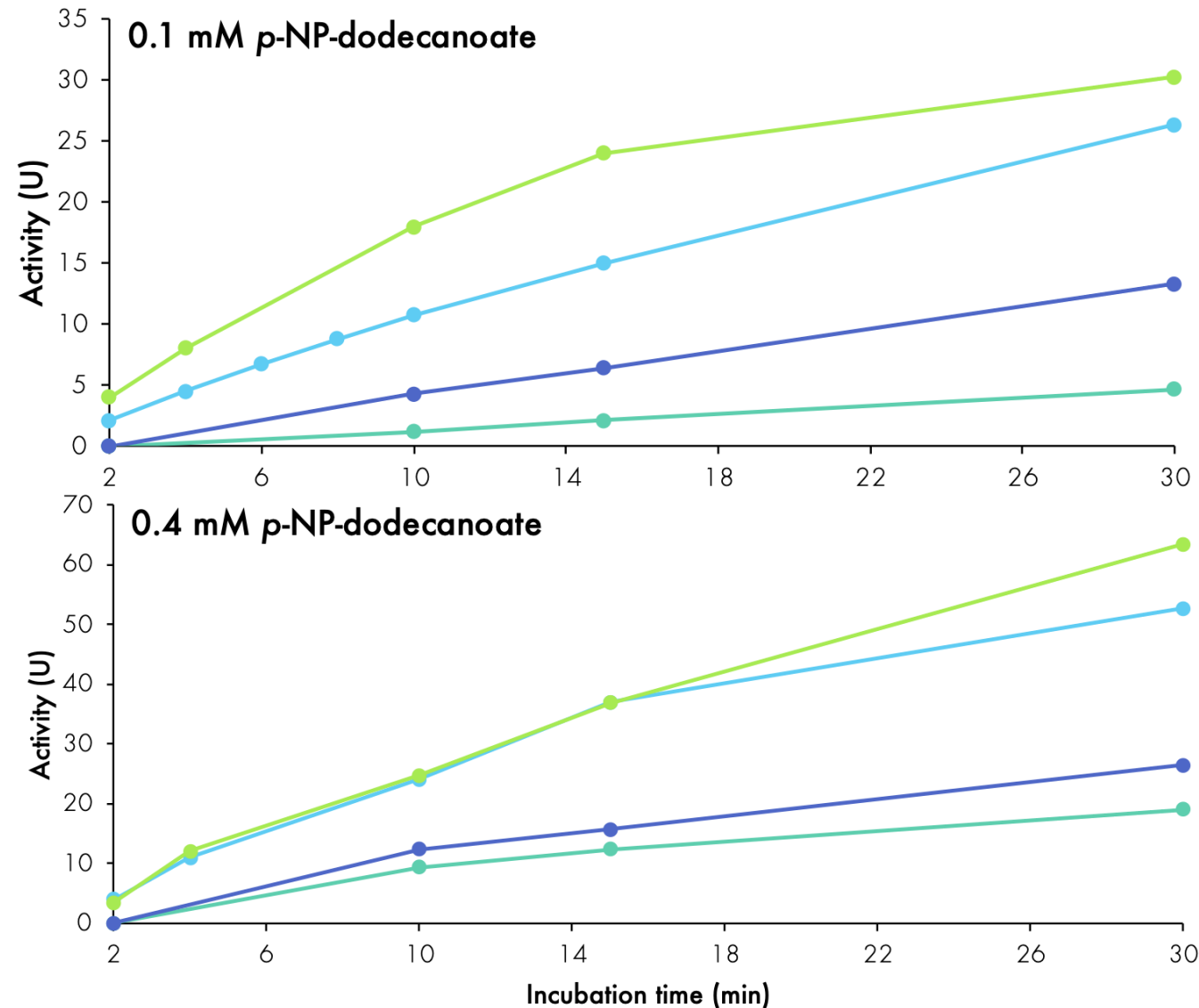
Colorimetric-spectrophotometric assays with chromogenic substrates: p-NP esters

Enzymatic activity	pNP esters	ASPA161_6	ASPA161_9
Esterase EC.3.1.1.1	<i>p</i> -NP-acetate (C ₂)	+	+
	<i>p</i> -NP-butyrate (C ₄)	+	+
Lipase EC.3.1.1.3	<i>p</i> -NP-decanoate (C ₁₀)	+	+
	<i>p</i> -NP-dodecanoate (C ₁₂)	+	+

IN-VITRO ASSAY: COLORIMETRIC-SPECTROPHOTOMETRIC ASSAYS

Hydrolysis of ester bonds (static incubation, different temperatures and incubation times)

EXAMPLE: *Psychrobacter* sp. strain ASPA161_9

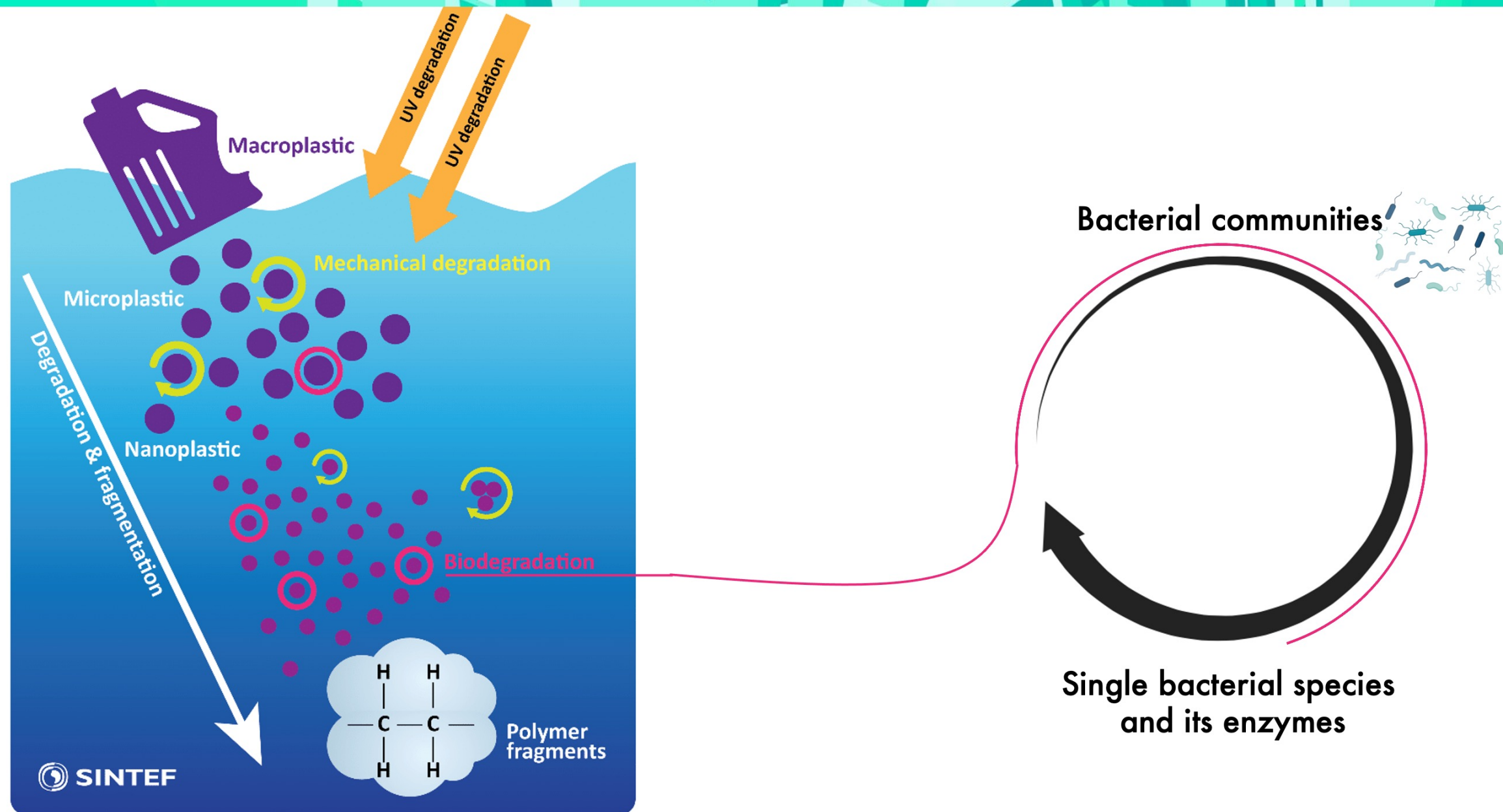


**Similar results with both strains
and the different substrates**

- 35 °C
- 25 °C
- 15 °C
- 4 °C

1U: activity releasing 0.001 μ mol of *p*-nitrophenol under the assay conditions

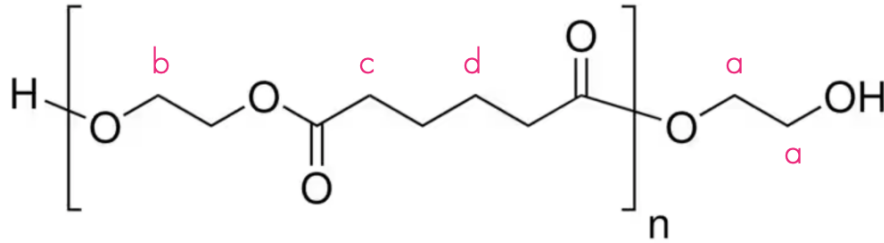
INSIGHTS INTO PLASTIC POLLUTION AND MICROBIAL PLASTIC TRANSFORMATION



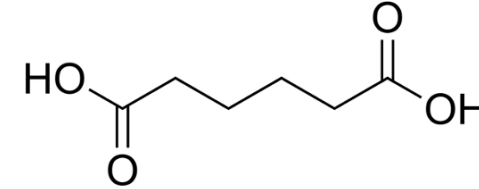
SET-UP: CONTROL OF THE DEGRADATION OF PLASTICS



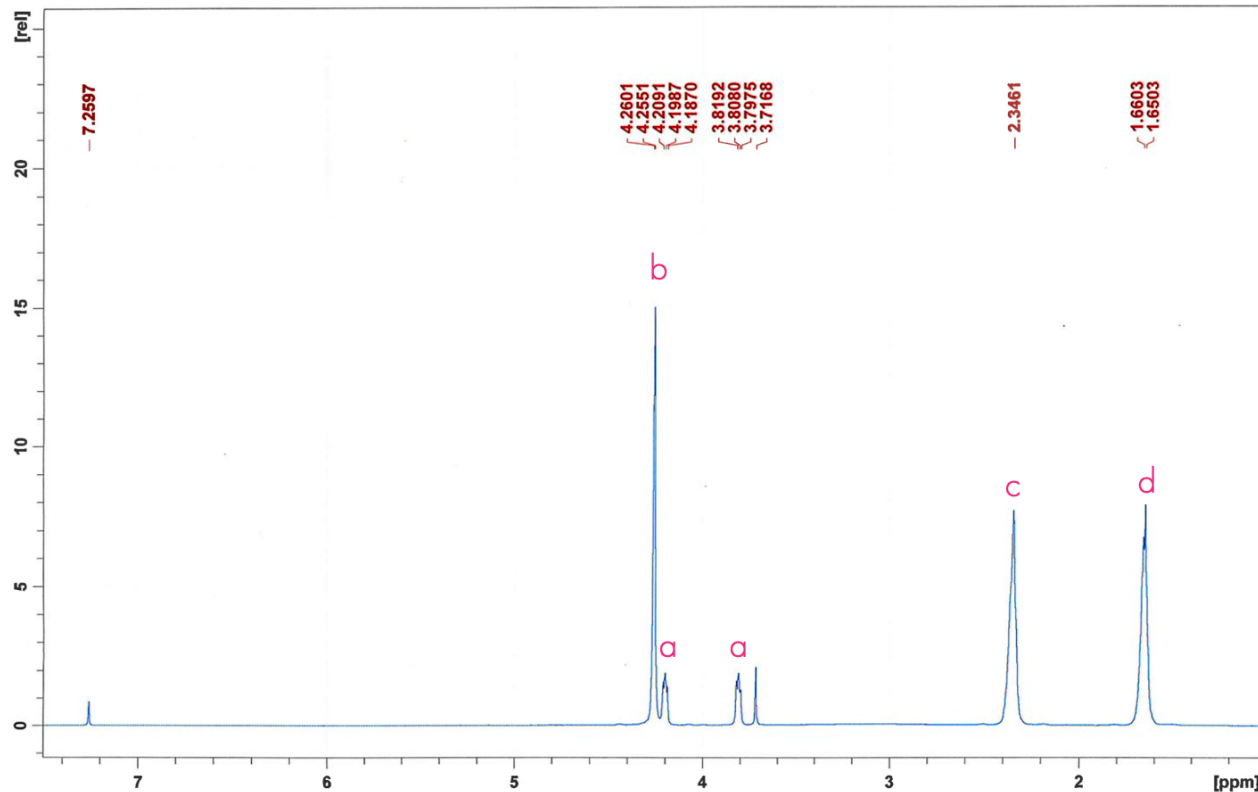
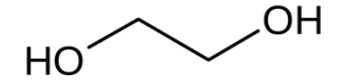
Poly(ethylene adipate) PEA



Adipic acid



Ethylene glycol



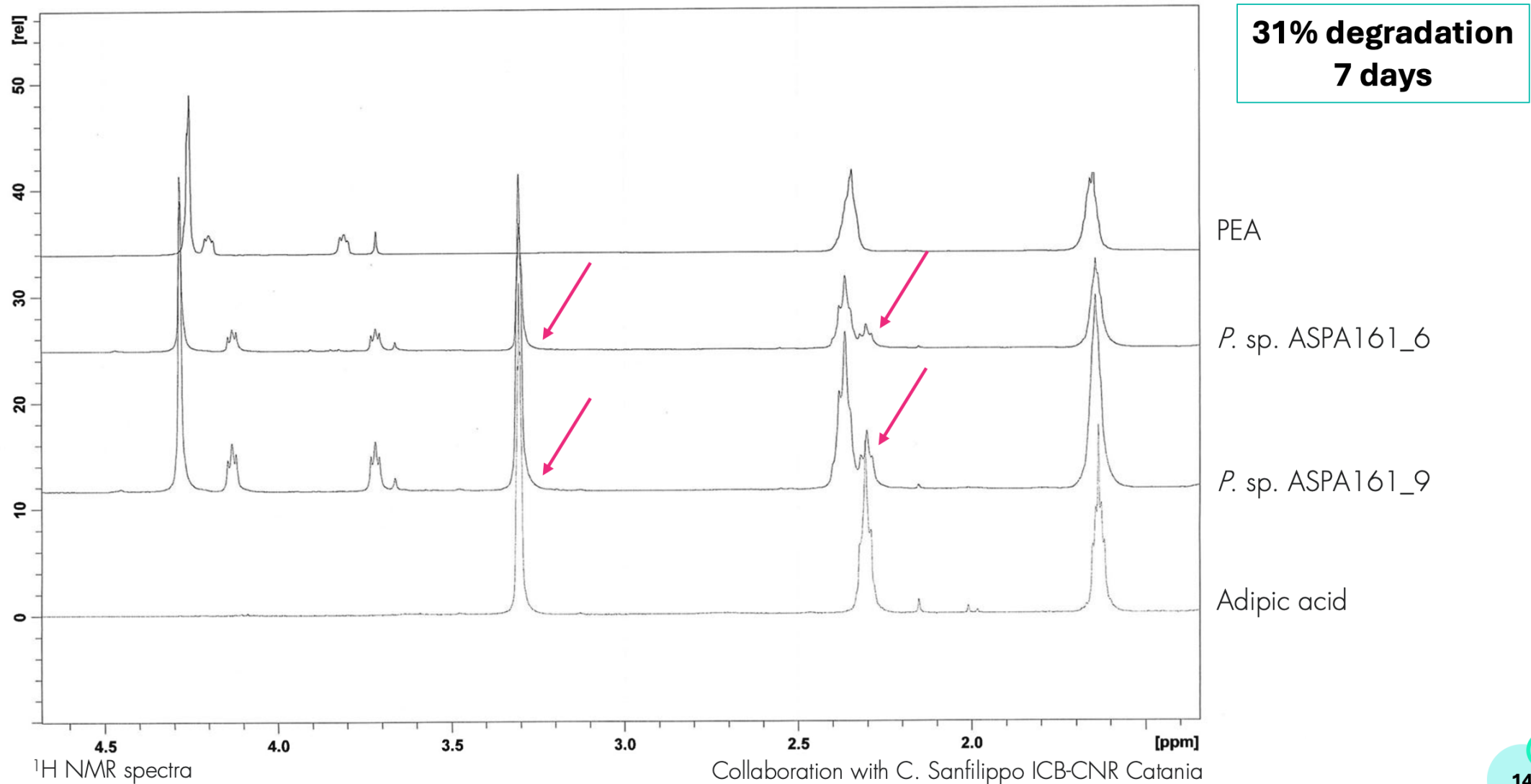
^1H NMR spectrum of PEA

Collaboration with C. Sanfilippo ICB-CNR Catania

Biodegradation time > 30 days (PEA film, 37 °C)

SET-UP: CONTROL OF THE DEGRADATION OF PLASTICS

PEA degradation by partially purified extracellular enzymes from the supernatant



SET-UP: CONTROL OF THE DEGRADATION OF PLASTICS

PEA degradation by whole cells



ON GOING



**Incubation 7 and 30 days
15 and 25 °C**

NMR study

Collaboration with C. Sanfilippo ICB-CNR Catania

NEXT STEPS

☐ **Improving** of the enzymatic degradation performances

☐ Setting-up of experiments for **bacterial cell-mediated degradation** suitable with **sea water temperature**

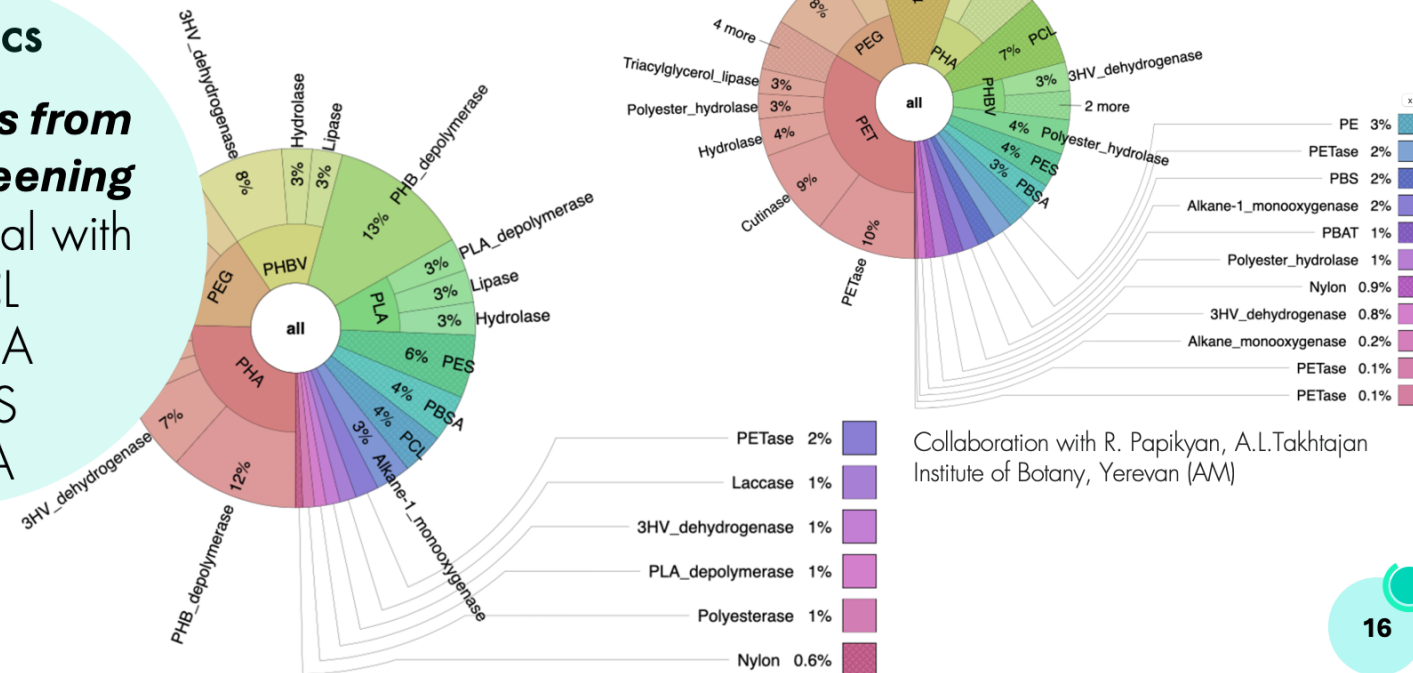
☐ **Gene cloning and expression** to produce enzymes of interest

☐ Evaluating **experiments onto other (bio)plastics**

Results from *in-silico* screening

potential with

- PCL
- PHA
- PES
- PLA



Collaboration with R. Papikyan, A.L.Takhtajan
Institute of Botany, Yerevan (AM)

Extremophilic Research Group

ICB-CNR, Pozzuoli (IT)



Institute of
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Luigi Leone

Maria Vittoria

\$4.5

Antonia D'Amodio

\$4.4

Giovanni Pezzella

\$4.6

**Thank you
for the attention!**

Collaborations

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Roberto Abbamondi ICB-CNR, Pozzuoli (IT)

\$4.7

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European Culture
Collections' Organisation

XLII Meeting – Bari (IT) 18-20 September 2024