

Lipase-designed biocatalysts for silymarin derivatization

Dissertation Thesis: Experimental report
Chemistry of Advanced Materials

Student: Giulia-Roxana Gheorghiu

Scientific Coordinator: Assoc. Prof. Dr. Mădălina Tudorache

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1.

OUTLINE

Biocatalyst preparation and characterization

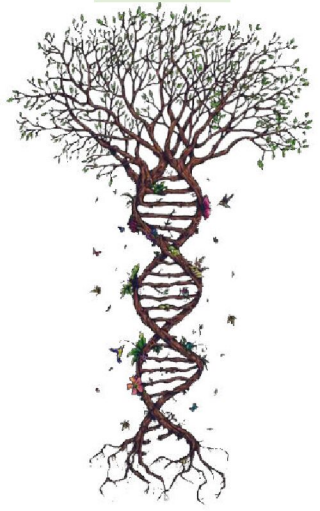
Immobilization approaches and performance

Relative enzyme activity

Design of the biocatalytic system

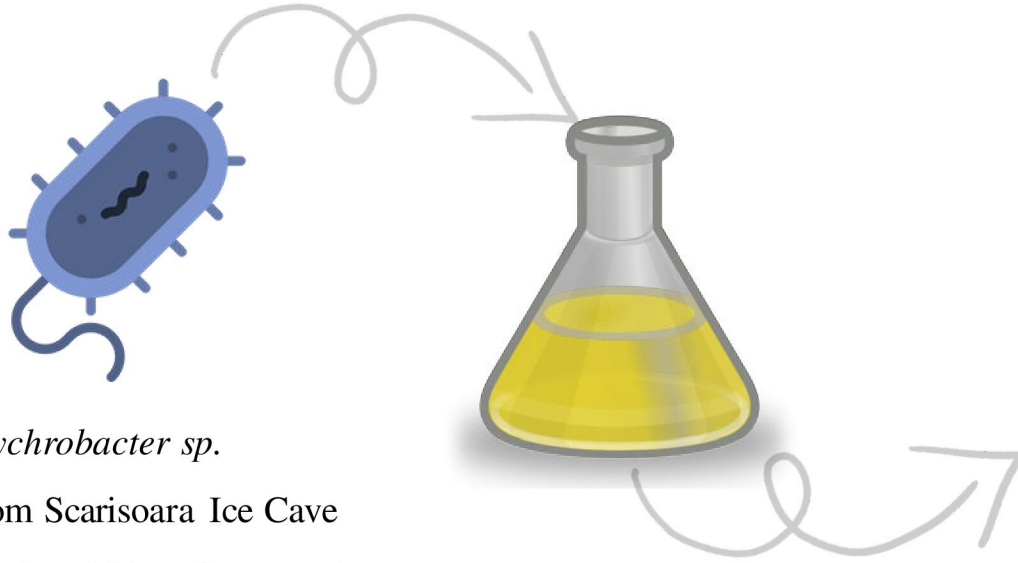
Testing the biocatalytic system

Conclusions



2.

BIOCATALYST PREPARATION AND CHARACTERIZATION



Psychrobacter sp.

From Scarisoara Ice Cave

Psychrophilic microorganism

Producing cold-active
extracellular lipases

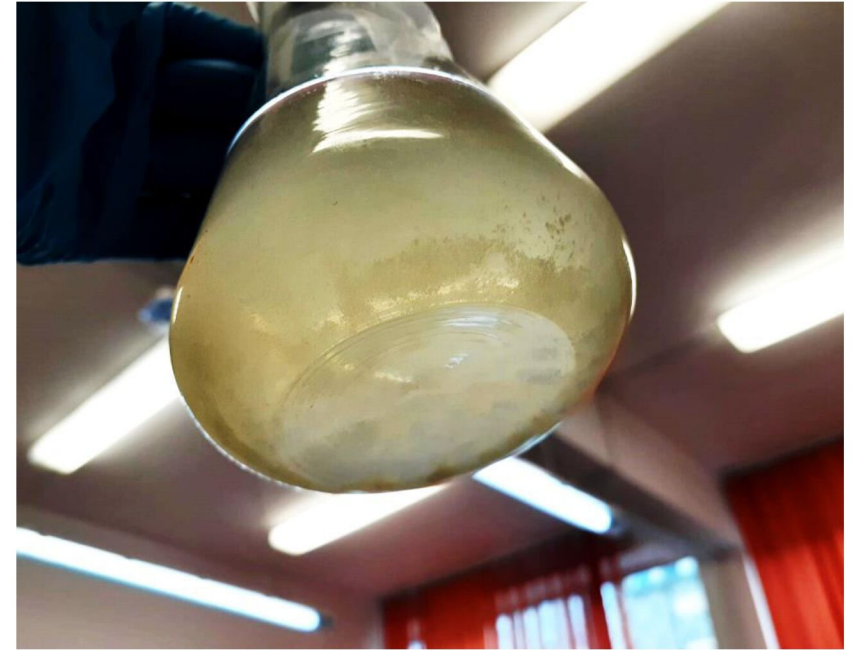
Reasoner's or Tryptic soy media

Proper cultivation media for *Psychrobacter*

15°C for 3 or 4 days under stirring

Supplimented with olive oil*

*carbon source for lipase production

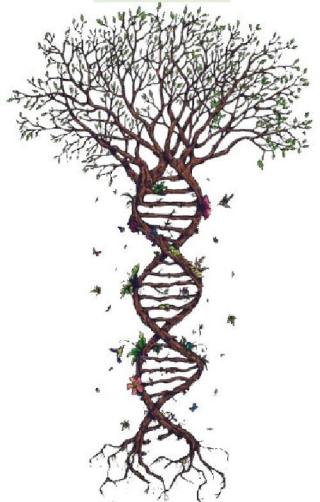


After bacterial growth: cell deposition and
supernatant collection.

Supernatant contains extracellular lipases.

Extraction with organic solvent: 80% cold
acetone added drop by drop onto the
supernatant.

Why is precipitation needed?



3.

BIOCATALYST PREPARATION AND CHARACTERIZATION

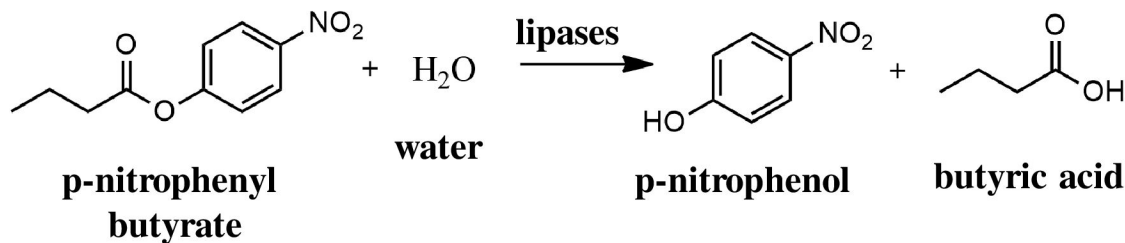
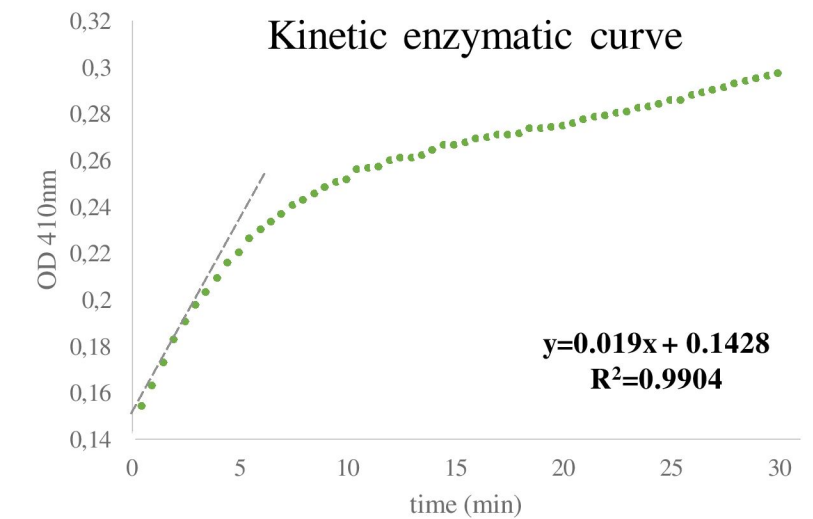
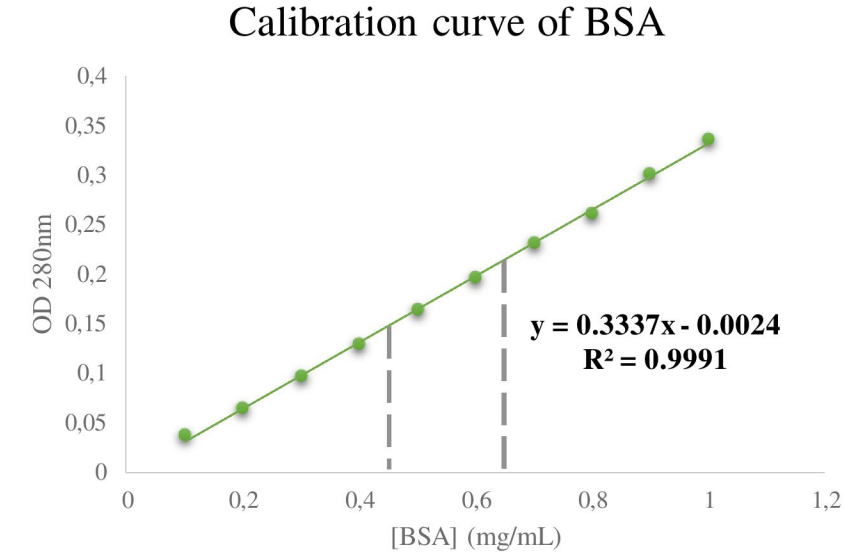
Protein concentration

After protein powder finally obtained,
resuspension in 100mM TRIS-HCl pH 8.
Sample diluted 1 to 10 for UV concentration analysis.

Concentration of the protein content
4.96 mg/ml - 6.6 mg/ml.

Enzyme kinetics

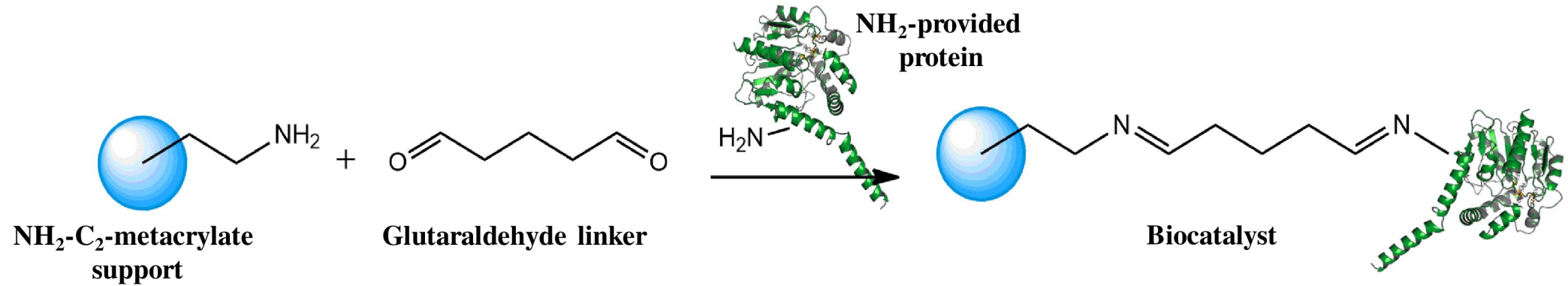
Substrate mix sonicated for emulsification:
4 750 μ l 100 mM TRIS-HCl pH 7.55,
90 μ l Triton100, 250 μ l p-NPB.
100 μ l protein extract + 900 μ l substrate mix.
Measuring continuously for 30 minutes at 410nm.
Lambert Beer's Law is applied.



Enzyme activity
4.25 μ M/min/mg protein

4.

IMMOBILIZATION OF THE BIOCATALYST



Procedure

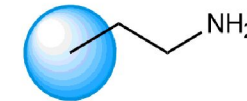
0.1g support

0.1% glutaraldehyde

100 μ L protein extract

in MES (pH 4.7) or PBS (pH 7.4) buffers

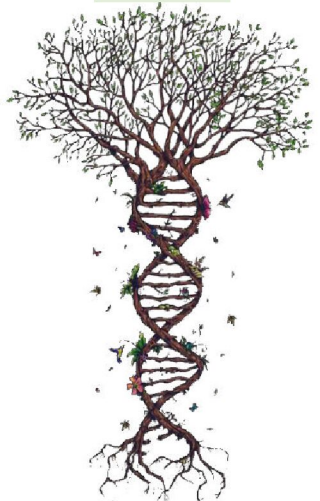
Legend:



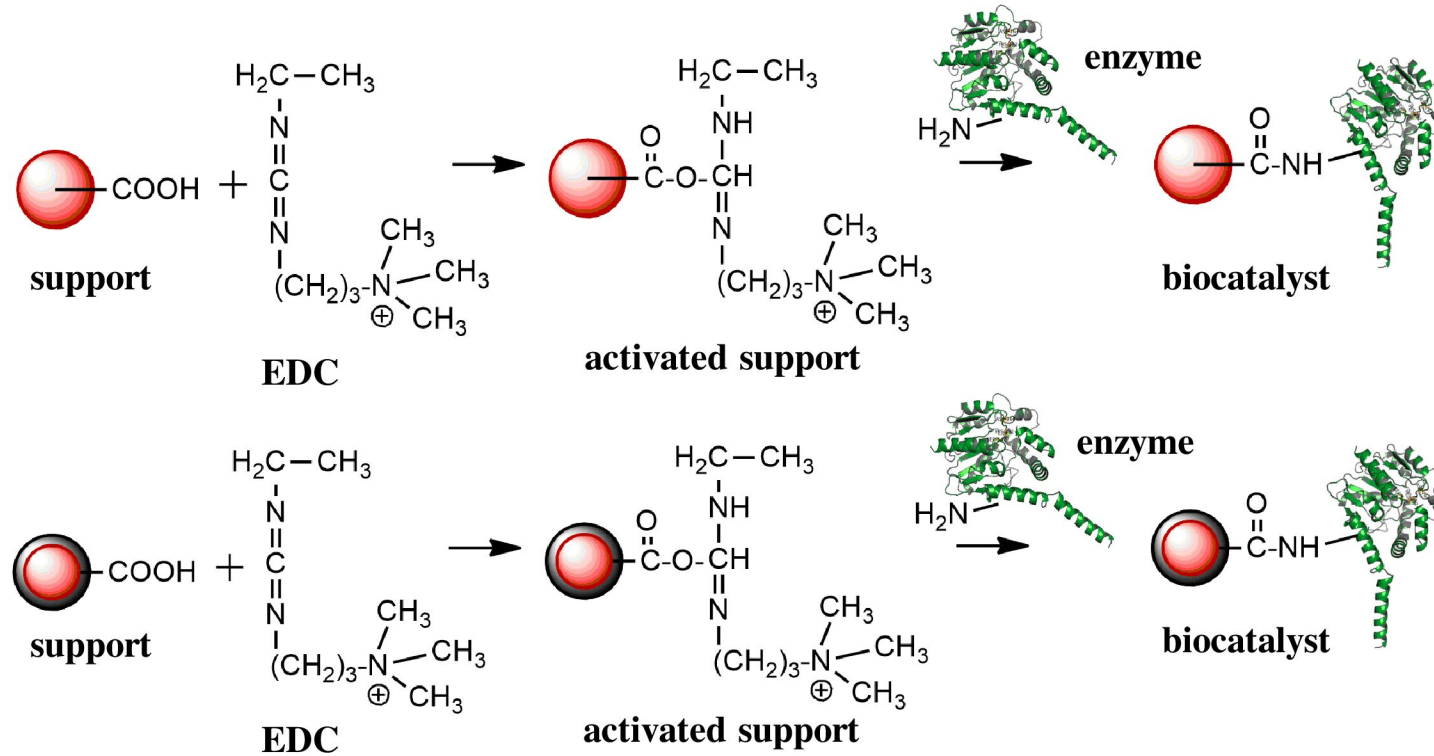
spherical beads (150–300 μ m) of methacrylate cross-linkers polymer functionalized with -NH_2 groups.

Immobilization performance

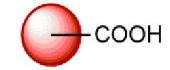
Starting concentration	Immobilized	Protein loading	Protein recovery
4.96 mg/mL	4.382 mg/mL	41.73 mg protein/g support	88.35%
	4.47 mg/mL	42.33 mg protein/g support	90.12%



IMMOBILIZATION OF THE BIOCATALYST



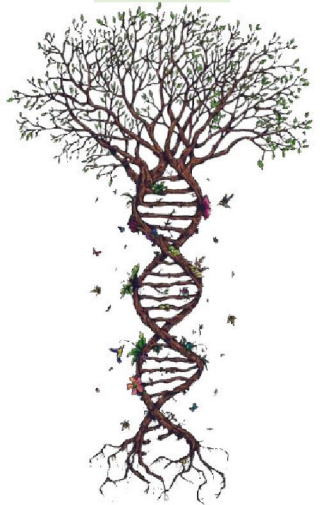
Legend:



0.5 μm to 1 μm magnetic cores coated with non-porous silica provided with functional groups.



50 nm, 100 nm and 200 nm beads of magnetic iron oxides covered with hydrophilic polymers with different functional groups.



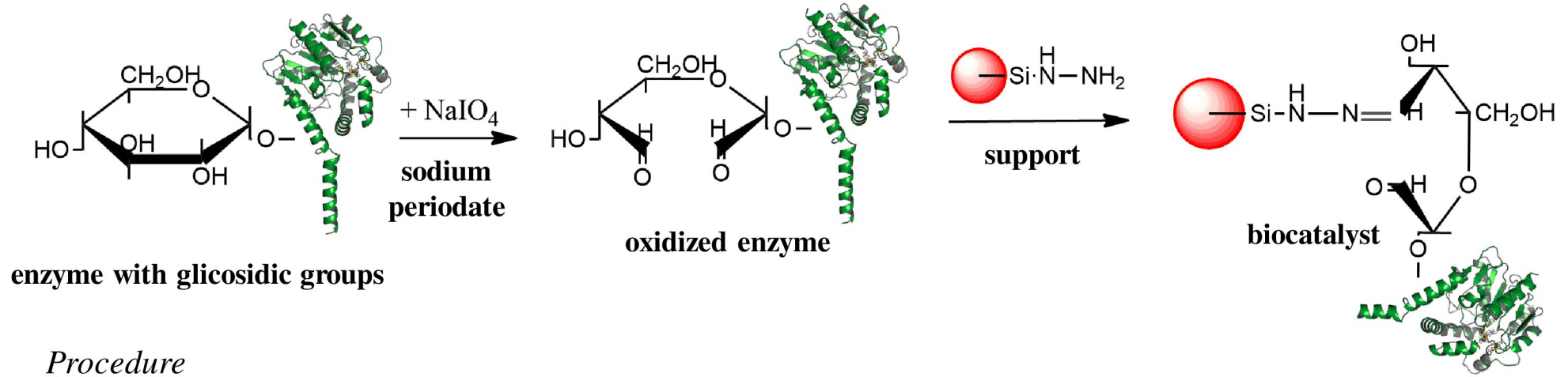
Carbodiimide method
 1mL support
 0.25M EDC in MES buffer
 200 μL protein extract

Immobilization performance

Starting concentration	Immobilized	Protein recovery	Biocatalyst
6.6 mg/mL	6.545 mg/mL	99.16%	Si-MAG-Carboxyl x Protein
	6.33 mg/mL	95.9%	Fluid-MAG-ARA x Protein

6.

IMMOBILIZATION OF THE BIOCATALYST



Procedure

500 μ L protein extract

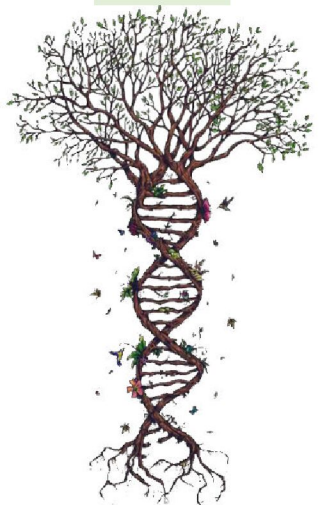
5mg NaIO₄

1mL support

250 μ L protein extract (oxidized)

Immobilization performance

Starting concentration	Immobilized	Protein recovery	Biocatalyst
6.6 mg/mL	6.53 mg/mL	98.94%	Si-MAG-Hydrazine x Protein



7.

RELATIVE ENZYME ACTIVITY

Enzyme activity of both free and immobilized protein

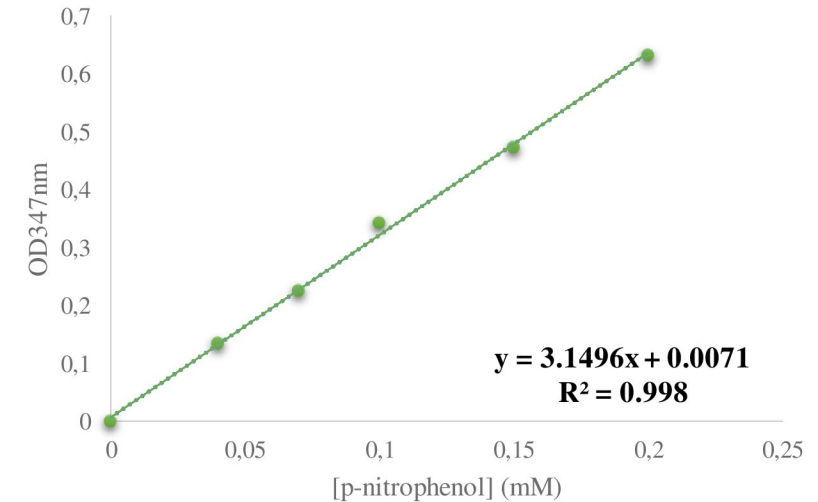
Reaction mix:

0.25 mL protein extract, 0.65 mL 0.1M TRIS-HCL pH 7,

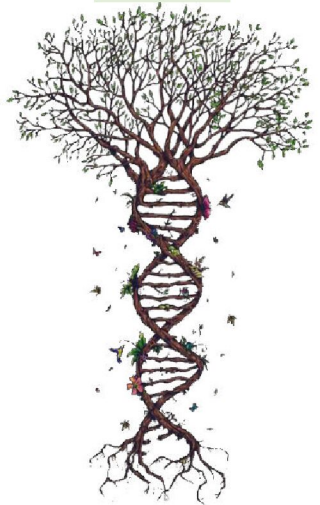
0.1 ml 0.025M substrate dissolved in acetonitril

30 minute at 37°C

Blocking solution: 0.1M Na₂CO₃

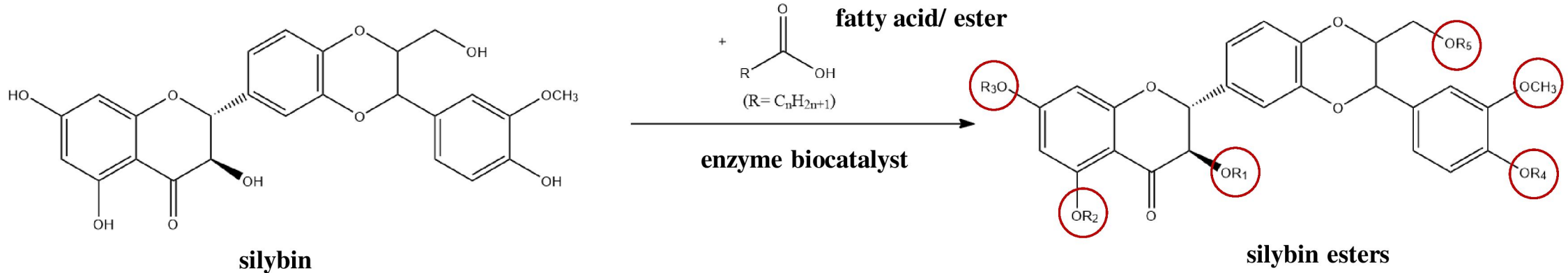


Relative enzyme activity	Biocatalysts	μM/min/mg protein
	Si-MAG-carboxyl x Protein	3.6
Si-MAG-hydrazine x Protein	3.6	
Fluid-MAG-ARA x Protein	5.9	
NH ₂ -C ₂ -metacrylate in MES x Protein	4.8	
NH ₂ -C ₂ -metacrylate in PBS x Protein	4.4	
Free protein extract	4.6	



8.

DESIGN OF THE BIOCATALYTIC SYSTEM



Reaction mixture

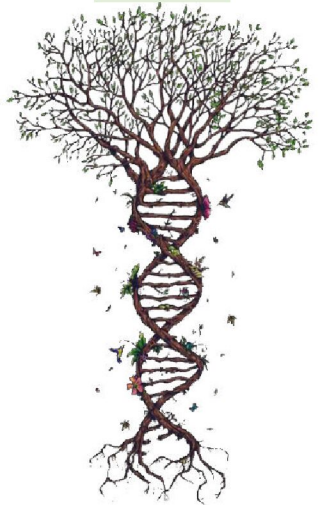
1mg of silybin in 1mL THF
 10 μ L of fatty acid/ester
 100 μ L of protein specimen
 25 $^{\circ}$ C for 24h at 1000rpm

centrifuged for 15 minutes at 1500rpm
 0.22 μ m milipore filtration
 Solvent evaporation at 70 $^{\circ}$ C
 Redissolving into mobile phase

HPLC analysis

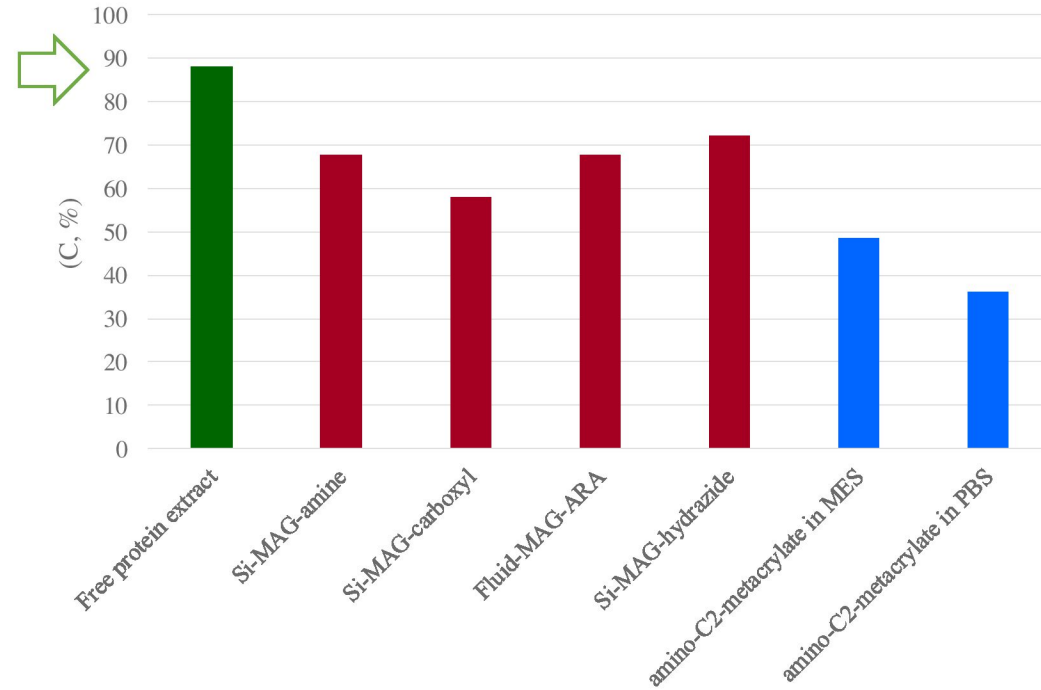
Poroshell 120 EC-C18 column
 acetonitrile and water (41/59, v/v) as mobile phase
 1mL/min flow rate and 25 μ L injection volume
 30 minutes reading
 DAD detection

$$C(\%) = \frac{\text{mass of converted substrate}}{\text{initial mass of substrate}} \times 100$$

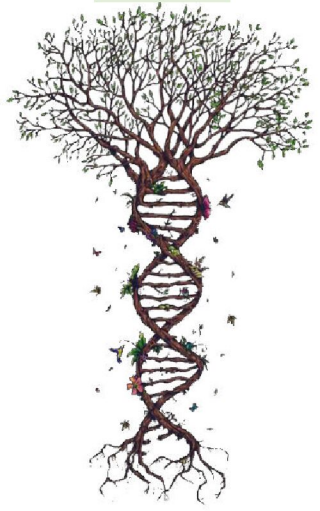
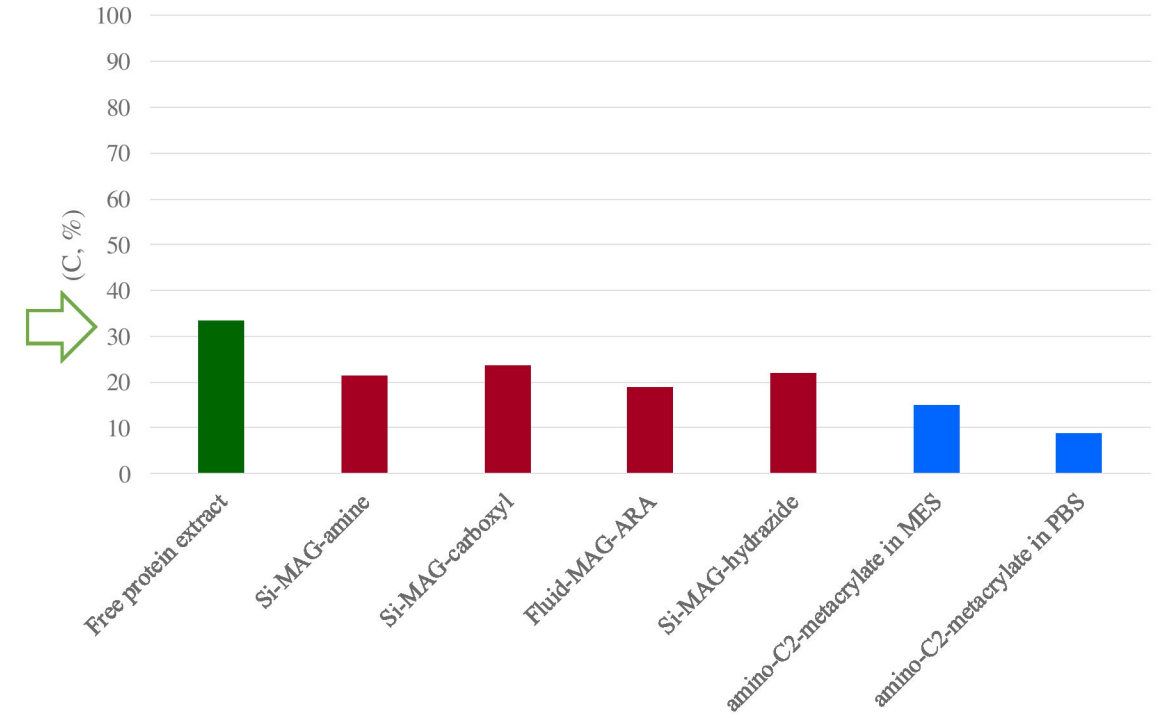


TESTING THE BIOCATALYTIC SYSTEM

Esterification of silybin with octanoic acid

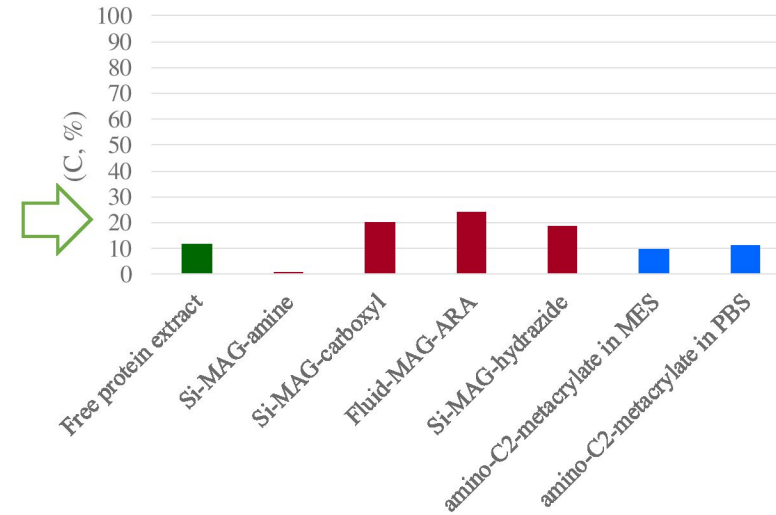
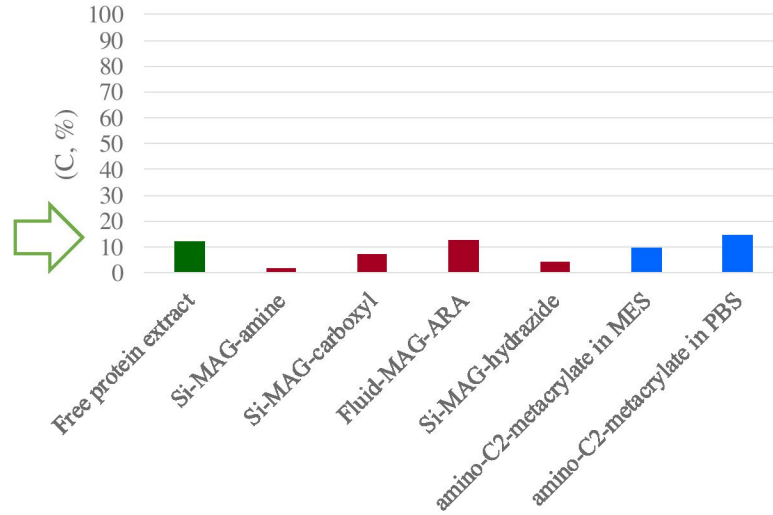


Esterification of silybin with oleic acid

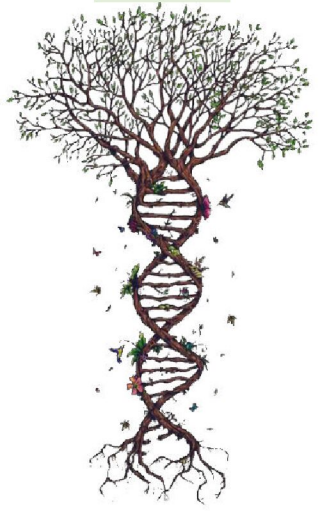
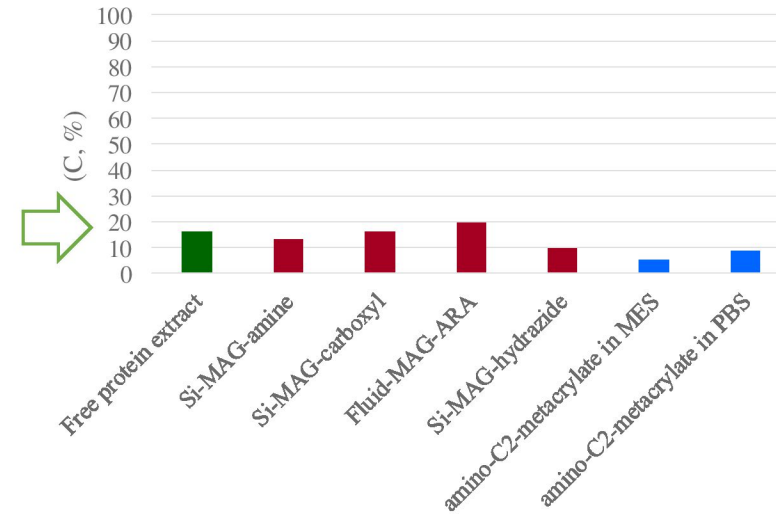
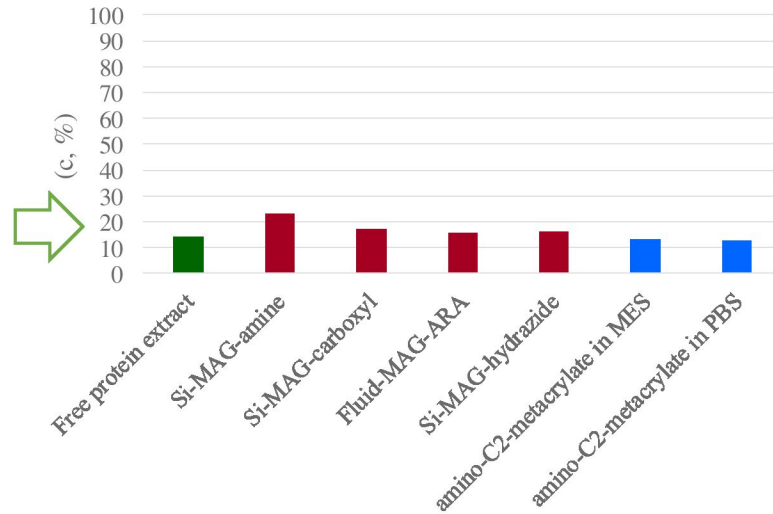


TESTING THE BIOCATALYTIC SYSTEM

Transesterification of silybin with methyl decanoate vs. methyl laurate



Transesterification of silybin with methyl myristate vs. methyl palmitate

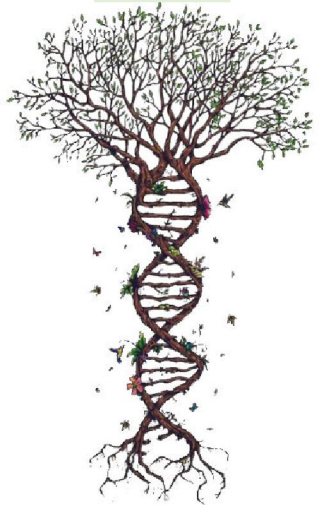


CONCLUSIONS

- Following the extracellular production of proteins, thus suspended in the culture medium of the microorganism, a purification step is absolutely necessary, in order to eliminate the interferences imposed by the composition of the environment or by the production of other biomolecules by the microorganism.
- Immobilization of the protein was achieved as a compromise between stability and activity, particularly imposed by the covalent bonding.
- The protein material obtained showed a good activity in catalyzing the esterification reaction between silybin and octanoic acid.

Perspectives

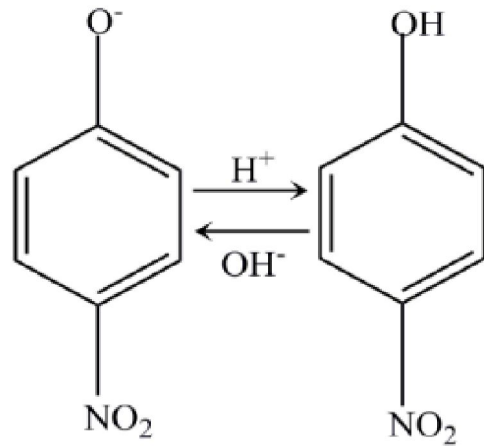
- Characterization of the free and immobilized biocatalyst through FTIR, SEM, CD.
- Optimizing the biocatalytic system in terms of good substrate conversion.
- Establishing if the cold-active character of the biocatalyst remains unaltered.



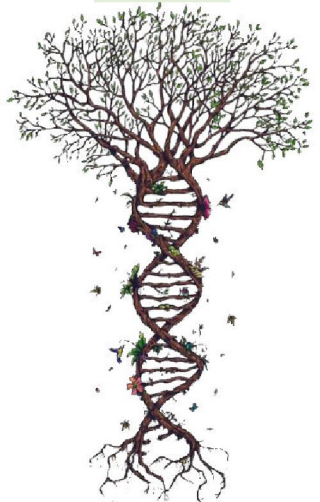
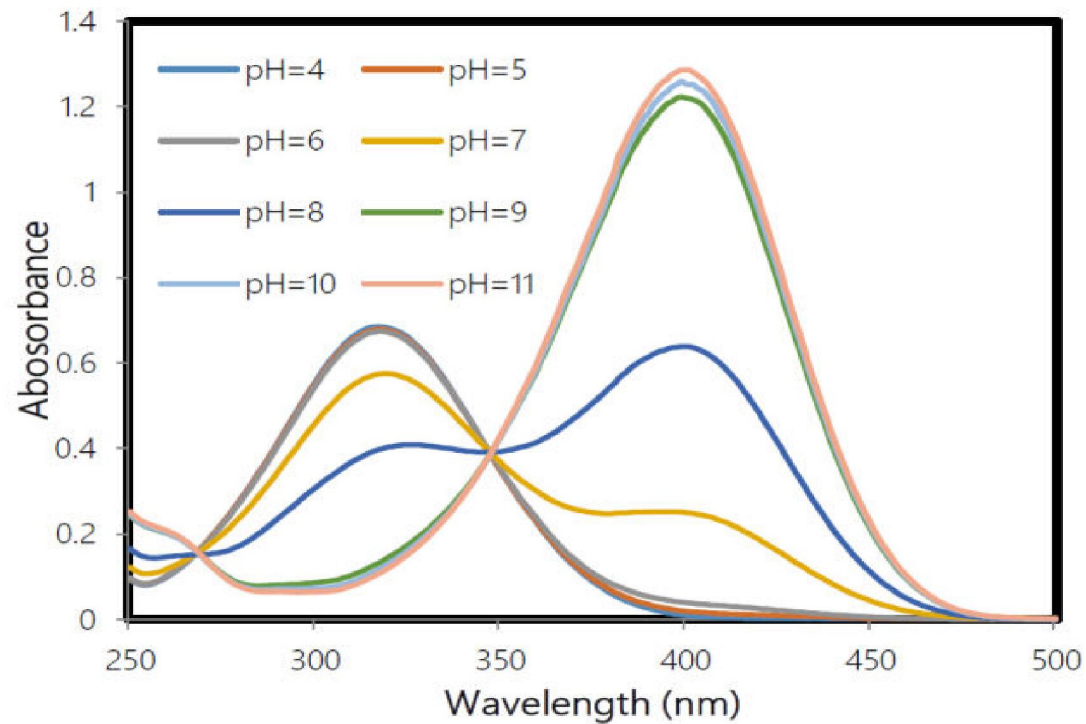


Thank you for your attention!

ANNEX



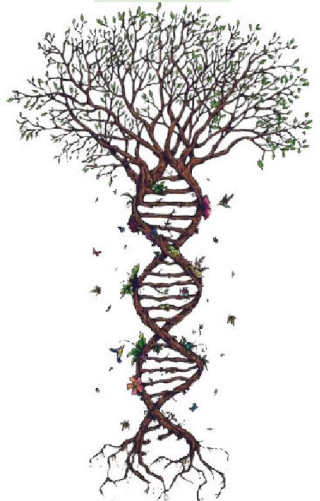
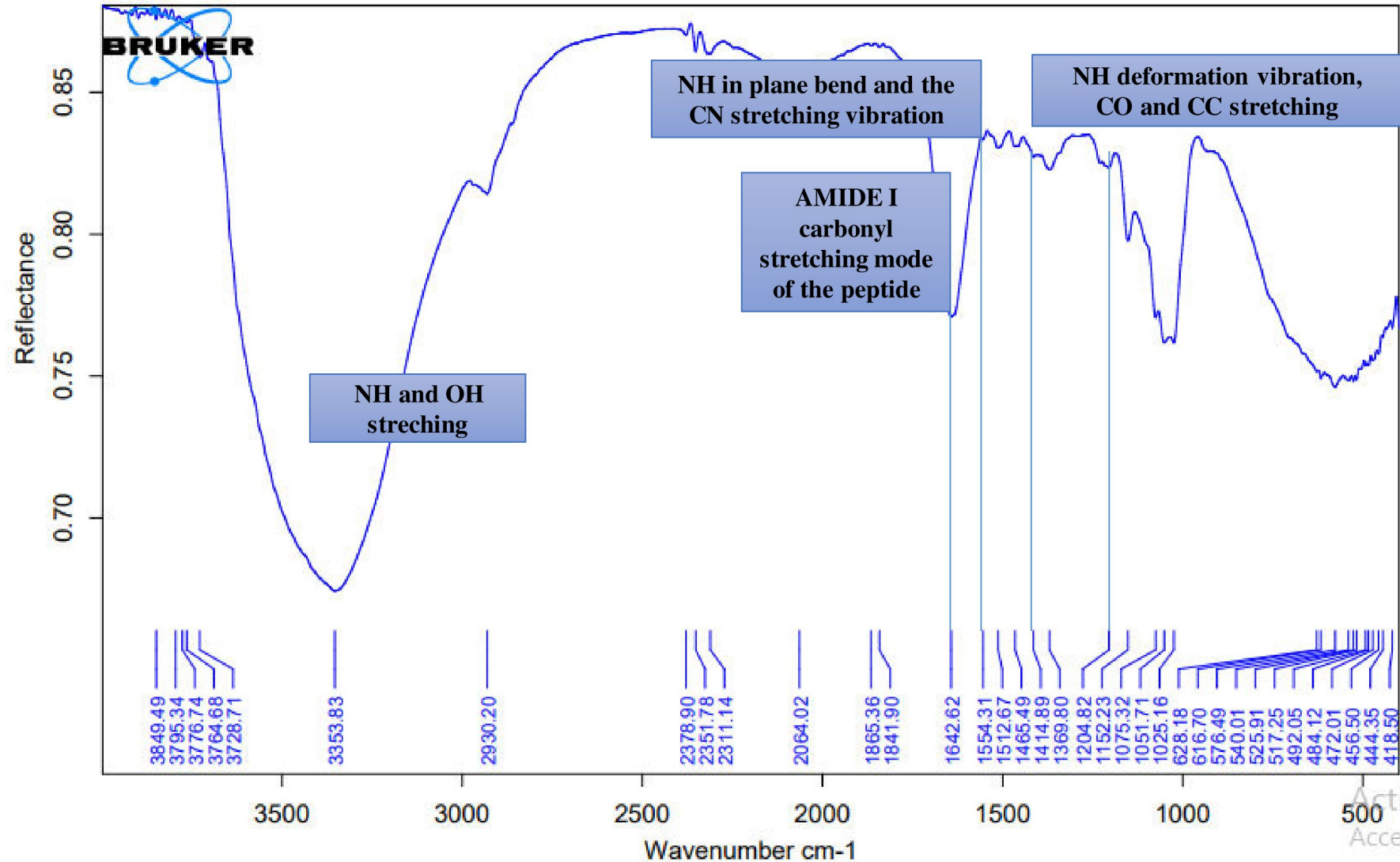
Ionized and non-ionized forms of p-nitrophenol



IMMOBILIZATION PERFORMANCE

FT-IR analysis in the context of biological materials

- *Free protein extract*



IMMOBILIZATION PERFORMANCE

FT-IR analysis in the context of biological materials

- Immobilization via magnetic particles support*

Legend:

Immobilized protein
free protein extract

support

