

Exploring ligninolytic fungi capabilities for pharmaceuticals detoxification in water effluents

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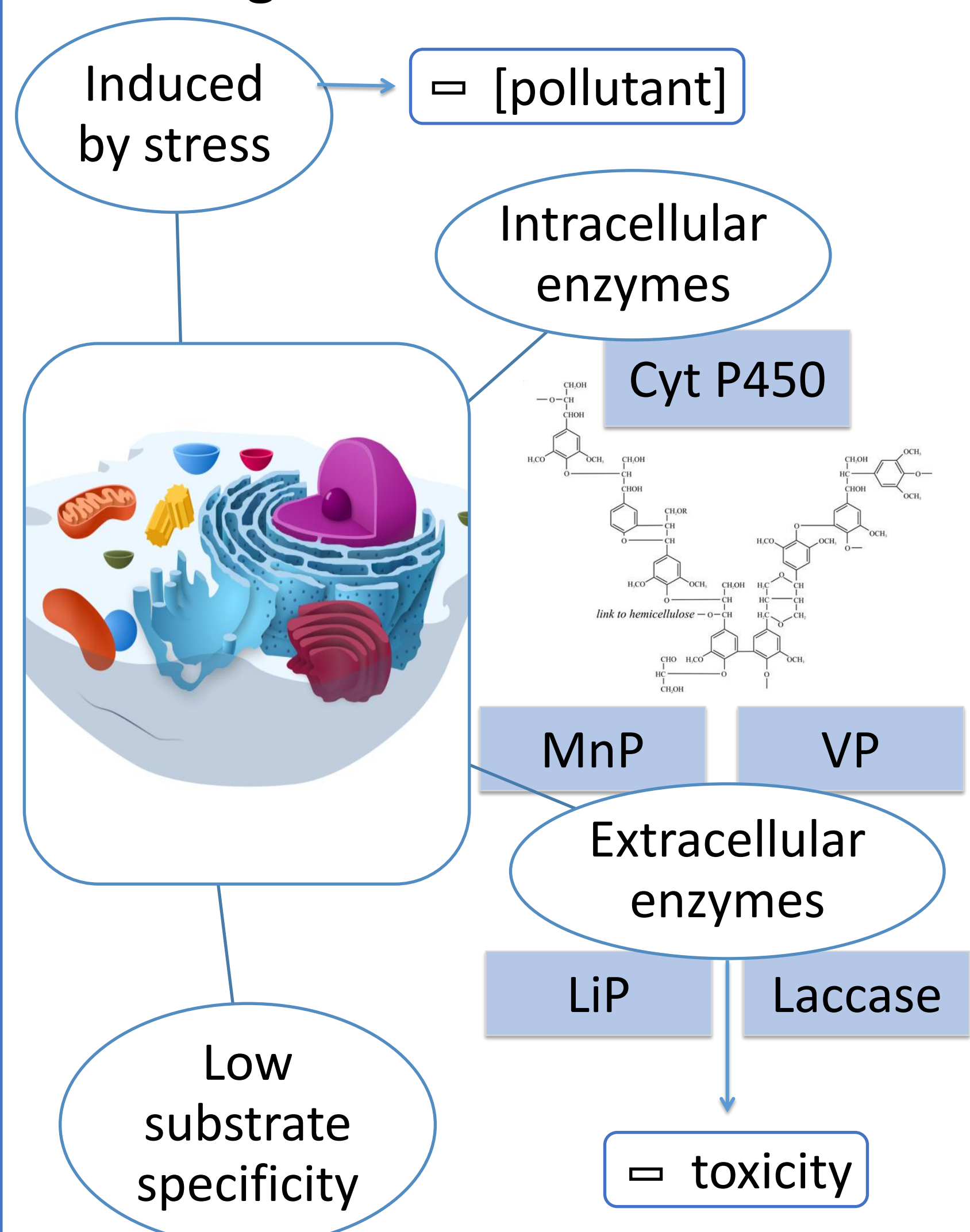
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INTRODUCTION

Statement of the Problem

- Pharmaceuticals consumption rise means contamination rise
- Conventional wastewater treatment plants are not efficient at pharmaceuticals removal

White rot fungi metabolism advantages for bioremediation



OBJECTIVES

Master Thesis specific objectives

- Development and validation of HPLC-HRMS analytical method
- Identification of TPs in water and fungal pellet

Master Thesis main goals

- Ocurrence of selected pollutants in mycoremediation samples
- Characterization of TPs from mycoremediation

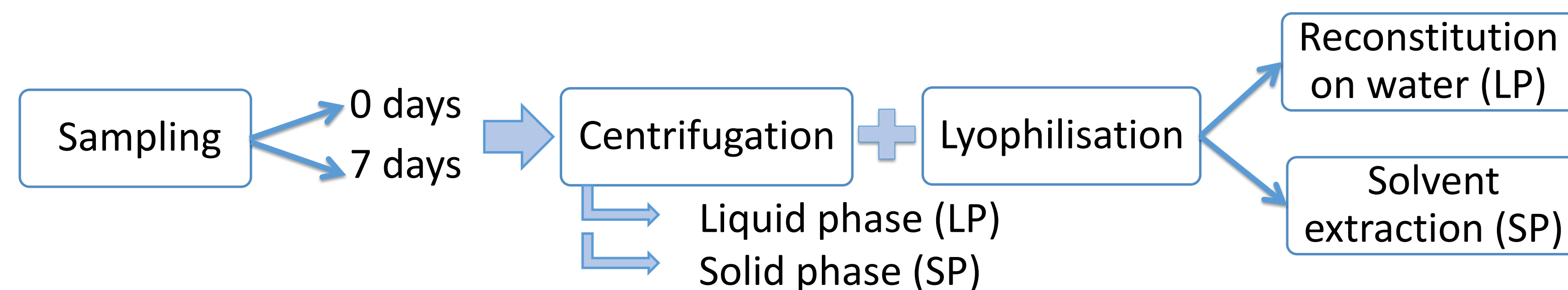
Creation of a spectral library of TPs

Identification of degradation causes of TPs detected

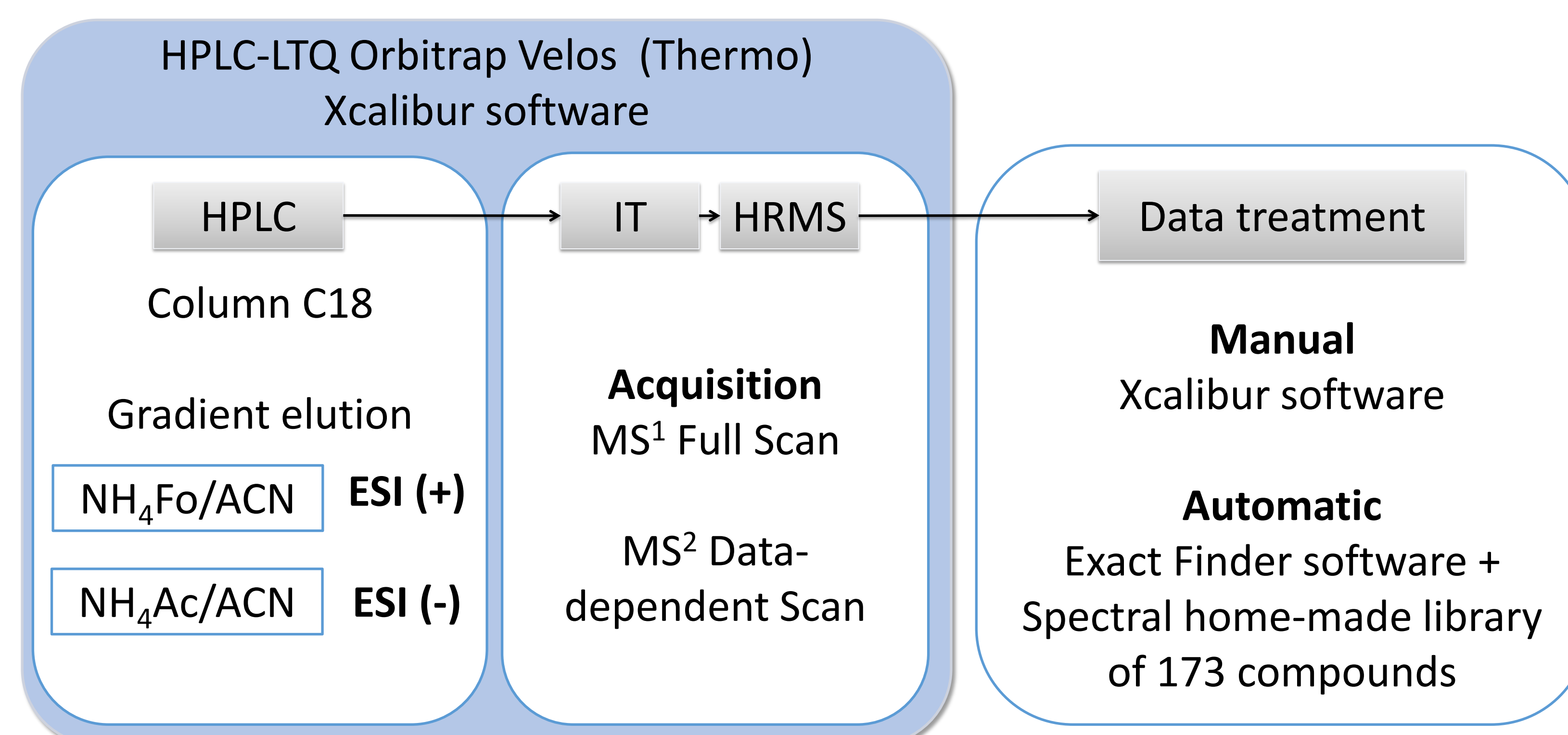
MATERIALS & METHODS

Experimental Design

		Production of fungal pellet + Culture medium triplicates		
Aerobic incubation		Inoculation with fungus (<i>T. versicolor</i> / <i>I. lacteus</i> / <i>G. lucidum</i>)		
		ALIVE FUNGUS	HEAT-INACTIVATED	NO INOCULUM
Spiking with 1000 ppm of pollutants	YES	Experimental batch	Killed Control	Abiotic Control
	NO	Alive Control	N.A.	N.A.



Analysis Method

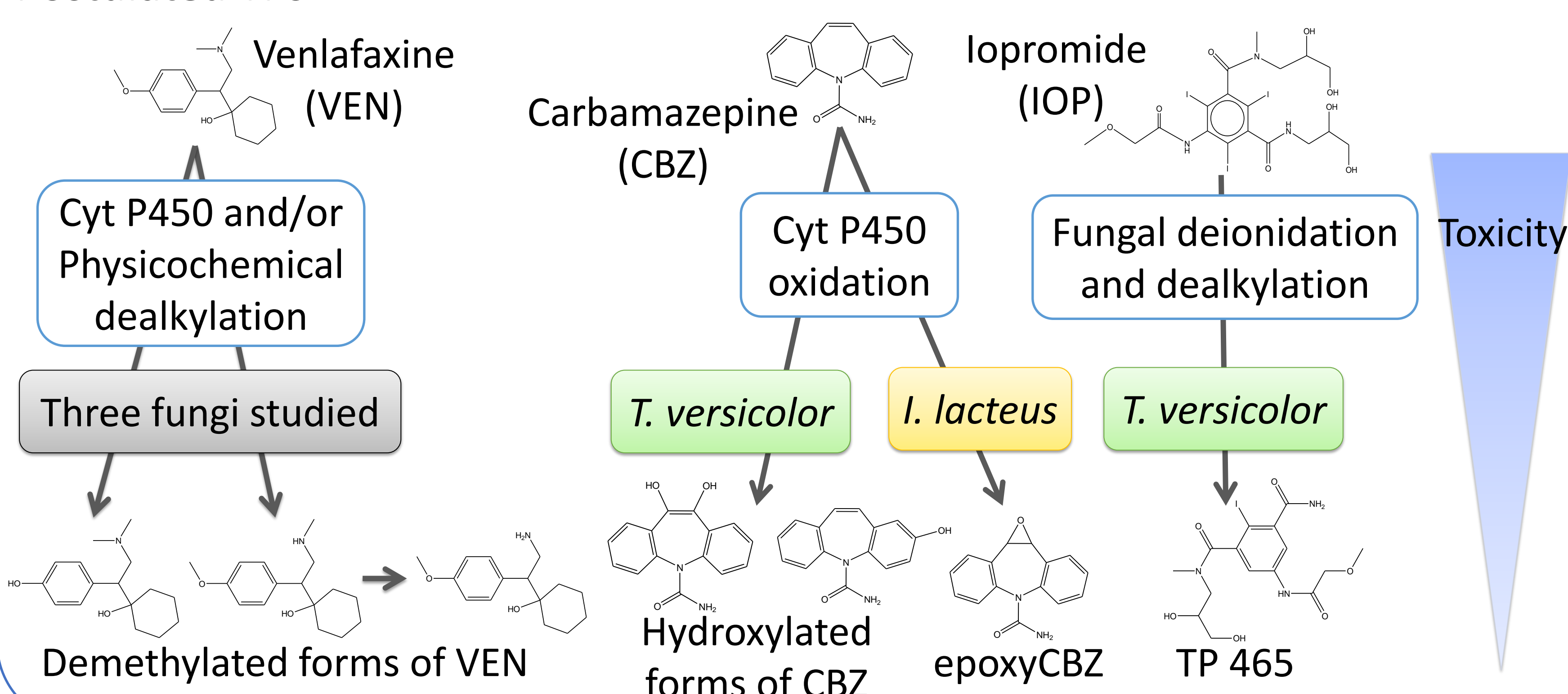


RESULTS

Method validation: UNE-EN-657/2002

Mass accuracy		
Accuracy	Precision	Uncertainty
✓ Bias < ±5ppm	✓ RSD _{intraday} < 10ppm ✓ RSD _{interday} < 10ppm ✓ RSD _T < 10ppm	✓ u _{RSD} < 10ppm ✓ u _{Bias} < 10ppm ✓ u _{exp} < 10ppm

Postulated TPs



CONCLUSIONS

Occurrence of selected pollutants

TPs toxicity < Parental compounds toxicity

Detoxification of water by mycoremediation similar to conventional waste water treatment

Sorption onto fungal pellet ≈ Sorption onto conventional sludge

Characterization of TPs

Specificity of method analysis (mass accuracy) + Physicochemical data from literature (t_R, pK_{OW})

Accurate TPs tentative identification

Degradation pathways

T. versicolor + *I. lacteus* + *G. lucidum*

Biological + Physicochemical

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H2PHARMA project (Fungi, algae and bacteria in pharmaceuticals degradation)

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