

### **Excellence** in Science Innovative Integration of LC-HRMS and In-Silico Libraries for Pharmaceutical Metabolite Annotation and MALDI-HRMS Spatial Residue Analysis in Environmental Samples Fučík Jan<sup>1</sup>, Fučík Stanislav<sup>2</sup>, Jarošová Rea<sup>3</sup>, Rexroth Sascha<sup>4</sup>, Baumeister Andreas<sup>4</sup>, Niehoff Ann Christin<sup>4</sup>, Navrkalová Jitka<sup>1</sup>, Hamplová Marie<sup>1</sup>, Zlámalová Gargošová Helena<sup>1</sup>, Mravcová Ludmila<sup>1</sup>

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## **Workflow for LC-HRMS Metabolite Annotation**

Steps 1-3: Lactuca sativa and Eisenia fetida were exposed to a six-pharmaceutical mix (atenolol, enrofloxacin, erythromycin, ketoprofen, sulfamethoxazole, tetracycline) in soil or water. Root and leaf samples of Lactuca sativa Eisenia fetida, as well as soil and water samples were collected at multiple time points during the exposure experiment and analysed using a validated QuEChERS extraction method. Step 4: Metabolite structures were predicted via literature review, software tools (BioTransformer 3.0, GLORYx, enviPath), and known metabolic pathways. **Step 5:** In-silico MS/MS libraries (ESI+/-) were generated using CFM-ID 4.0, aided by a custom script developed at BUT FCH and FEEC. Step 6: Water and lettuce extracts were analysed at Shimadzu Europa GmbH using LCMS-9050 (DDA & DIA, ESI+/-). Data were processed in MS-DIAL 4.0 and statistically evaluated with MetaboAnalyst 4.0.



Fig. 2 Venn Diagram Comparing Metabolite Identification Across Different MS/MS Modes, Showing Efficiency the Number of Metabolites Identified by Each Mode.

Tetracycline-LS2

Tetracycline-LS3







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Soil Ear	thworm	s		Soil Lettuce						Hydroponics Let			
	EW		21 Days	Soil 28 Days	Lettuce Roots			Lettuce Leaves		Water	Lettuce Roots		
3 Days	7 Days	14 Days			14 Days	21 Days	28 Days	21 Days	28 Days	28 Days	14 Days	21 Days	28 Day
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## SHIMADZU

## MALDI-HRMS Spatial Residue Analysis

Pre-step: Eisenia fetida was initially exposed to a mixture of three pharmaceuticals-enrofloxacin, erythromycin and tetracycline, each representing a distinct class. Samples were collected at 0-, 1-, 3-, and 5-days post-exposure followed by a depuration step

Step 1: Earthworms were euthanized by freezing at  $-20^{\circ}$  C, embedded in OCT compound, and sectioned at a thickness of  $10-15 \,\mu\text{m}$  using a cryostat maintained at  $-20^\circ$  C. The tissue sections were then mounted onto ITO-coated slides.

Tetracycline = Blue and Erythromycin = Green). Time of earthworm exposure to contaminated soil is noted in bottom left, and "H Indicates earthworm head position

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## Conclusion

novel high-throughput workflow annotating pharmaceutica metabolites in environmenta samples (lettuce roots and leaves and earthworms) developed was using open-source and LC-HRMS. Combining software prediction, literature review 3,762 pathways, known metabolites were predicted for six compounds (atenolo parent enrofloxacin erythromycin sulfamethoxazole ketoprofen. silico MS/MS tetracycline). libraries were generated for both ionization modes and analysed using for DDA and DIA data MS-DIAL In total, 26 statistically significant 0.05) were metabolites (p < annotated. Additionally, these results neglecting that demonstrate pharmaceutical metabolites leads to an underestimation of potentia health and environmental risks.

imaging Mass spectrometry (Figure uniform showed of pharmaceuticals distribution in *Eisenia fetida*, indicating uptake via both skin and ingestion. These findings advance understanding of bioaccumulation in soil organisms support improved environmental risk assessment, and represent the first spatial analysis of drug uptake in *E. fetida* from a terrestrial matrix.

## Acknowledgment

This work was measured within Lab4you Shimadzu student program. Moreover, this study was financially supported by the Ministry of Education, Youth, and Sports of the Republic (Project No. Czech FCH-S-25-8807).

