



# High throughput separation of chiral molecules using non-covalent complexes and SIM2 analysis on a TIMS-ToF

Sandra Alves, Clément Chalet, Estelle Rathahao-Paris, Chenqin Cao, Denis Lesage, Annelaure Damont, Ekaterina Dariy, Alain Perret, François Fenaille, Jean-Claude Tabet

## ► To cite this version:

Sandra Alves, Clément Chalet, Estelle Rathahao-Paris, Chenqin Cao, Denis Lesage, et al.. High throughput separation of chiral molecules using non-covalent complexes and SIM2 analysis on a TIMS-ToF. 15émes journées scientifiques du RFMF, May 2023, Perpignan, France. hal-04862041

HAL Id: hal-04862041

<https://hal.science/hal-04862041v1>

Submitted on 2 Jan 2025

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

# High throughput separation of chiral molecules using non-covalent complexes and SIM<sup>2</sup> analysis on a TIMS-ToF

Sandra Alves<sup>1</sup>, Clément Chalet<sup>1</sup>, Estelle Rathahao-Paris<sup>1,2</sup>, Chenqin Cao<sup>2</sup>, Denis Lesage<sup>1</sup>, Annelaure Damont<sup>2</sup>, Ekaterina Dariy<sup>3</sup>, Alain Perret<sup>3</sup>, François Fenaille<sup>2</sup>, Jean-Claude Tabet<sup>1,2</sup>

<sup>1</sup> Institut Parisien de Chimie Moléculaire (IPCM), Faculté des Sciences et de l'Ingénierie, Sorbonne Université, UPMC Paris VI, 75005 Paris, France;

<sup>2</sup> CEA, Département Médicaments et Technologies pour la Santé (DMTS), LIAA, INRAE, Université Paris-Saclay, 91191 Gif-sur-Yvette, France;

<sup>3</sup> Génomique Métabolique, Genoscope, Institut François Jacob, CEA CNRS, Univ Evry, Université Paris-Saclay, 91057 Evry, France

## INTRODUCTION

- Chiral molecules pose a unique challenge in life sciences, since enantiomers may exhibit different biological activity/toxicity while having similar physical and chemical properties.
- Separation of enantiomers using **mass spectrometry** (MS) mostly relies on derivatization or formation of **non-covalent complexes** (NCCs) in the gas phase (with various metal/alkali cations and chiral selectors such as cyclodextrins or carbohydrates [1,2]). Alternatively, **ion-mobility spectrometry** (IMS) separates ions based on their *m/z* ratio and their collision cross section (CCS), which makes it a promising tool for the rapid analysis of isomeric compounds.
- In this work, chiral analyses were conducted on a trapped ion mobility spectrometry time-of-flight (TIMS-ToF) instrument, using serial acquisitions by applying single ion mobility monitoring (**SIM<sup>2</sup>**) at high mobility resolving power and at different ion mobility ranges [3]. The method was applied on various natural products such as amino acids (AAs) and biologically relevant molecules.

## RESULTS

### Fast IM analysis of chiral AAs

Table 1: Chiral separations expressed as ΔCCS% values obtained in positive ESI for each AA in enantiomer solutions (L+D)LL, in SIM<sup>2</sup> mode.

(x) No complex found in individual and enantiomer solutions (LLL, DLL and (L+D)LL); (-) No mobility separation or several peaks found in the enantiomer solutions (L+D)LL; Orange Chiral separation in the enantiomer solutions (L+D)LL only.

[...-H+Cu <sup>II</sup> ] <sup>+</sup> adduct	Positively charged			Negatively charged			Polar			Special			Hydrophobic							
Arg	Arg	His	Lys	Asp	Glu		Asn	Gln	Ser	Thr	Cys	Pro	Ala	Ile	Leu	Met	Phe	Trp	Tyr	Val
AA,Pro	1.3%	-	-	x	-		x	x	x	x	x		-	0.4%	-	x	0.9%	-	-	-
AA,Phe	1.4%	1.0%	-	1.1%	0.8%		-	1.0%	-	-	x	1.0%	-	-	-	0.7%	1.8%	1.6%	-	-
AA,AA	1.1%	-	1.0%	-	2.6%		x	-	x	x	-	-	-	1.3%	0.6%	-	1.6%	-	1.5%	x
AA,Pro,Pro	-	x	2.2%	-	-		x	1.7%	-	0.5%	x		-	-	-	-	1.8%	2.4%	2.0%	-
AA,Phe,Pro	0.9%	x	-	-	-		0.4%	1.1%	0.7%	-	x		-	-	-	2.1%	2.0%	1.3%	-	-
AA,Phe,Phe	-	x	0.6%	-	-		1.2%	-	-	-	x	-	-	0.9%	-	-	1.3%	-	1.0%	-
AA,AA,Pro	x	x	x	x	x		-	-	x	0.9%	-	-	x	-	-	-	1.4%	1.6%	1.6%	-
AA,AA,Phe	x	x	x	x	x		x	1.1%	0.5%	-	1.3%	1.8%	-	x	x	1.4%	-	0.5%	-	
AA,AA,AA	x	x	x	x	x		-	1.3%	-	x	-	x	-	x	-	0.8%	-	-	-	

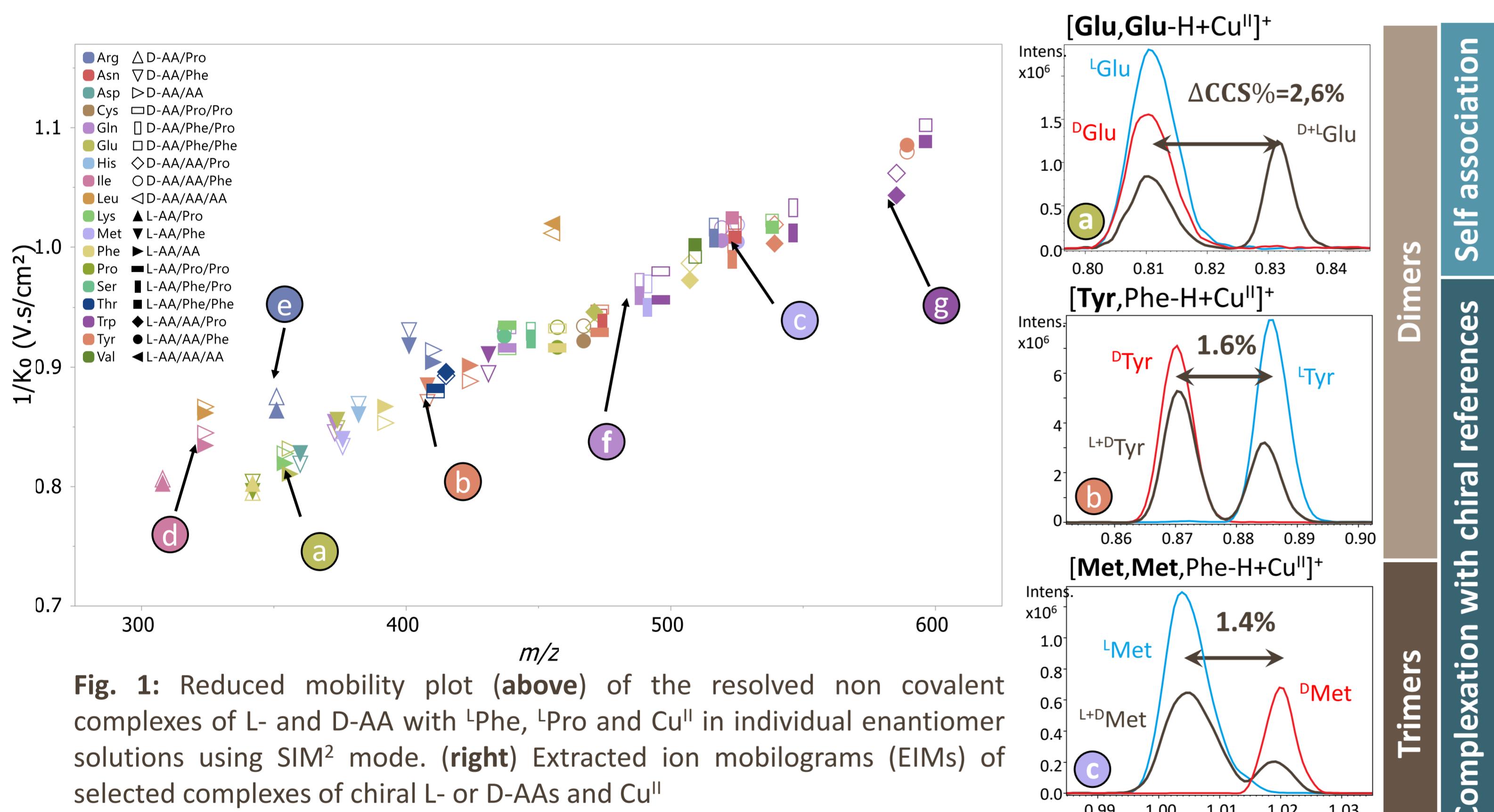
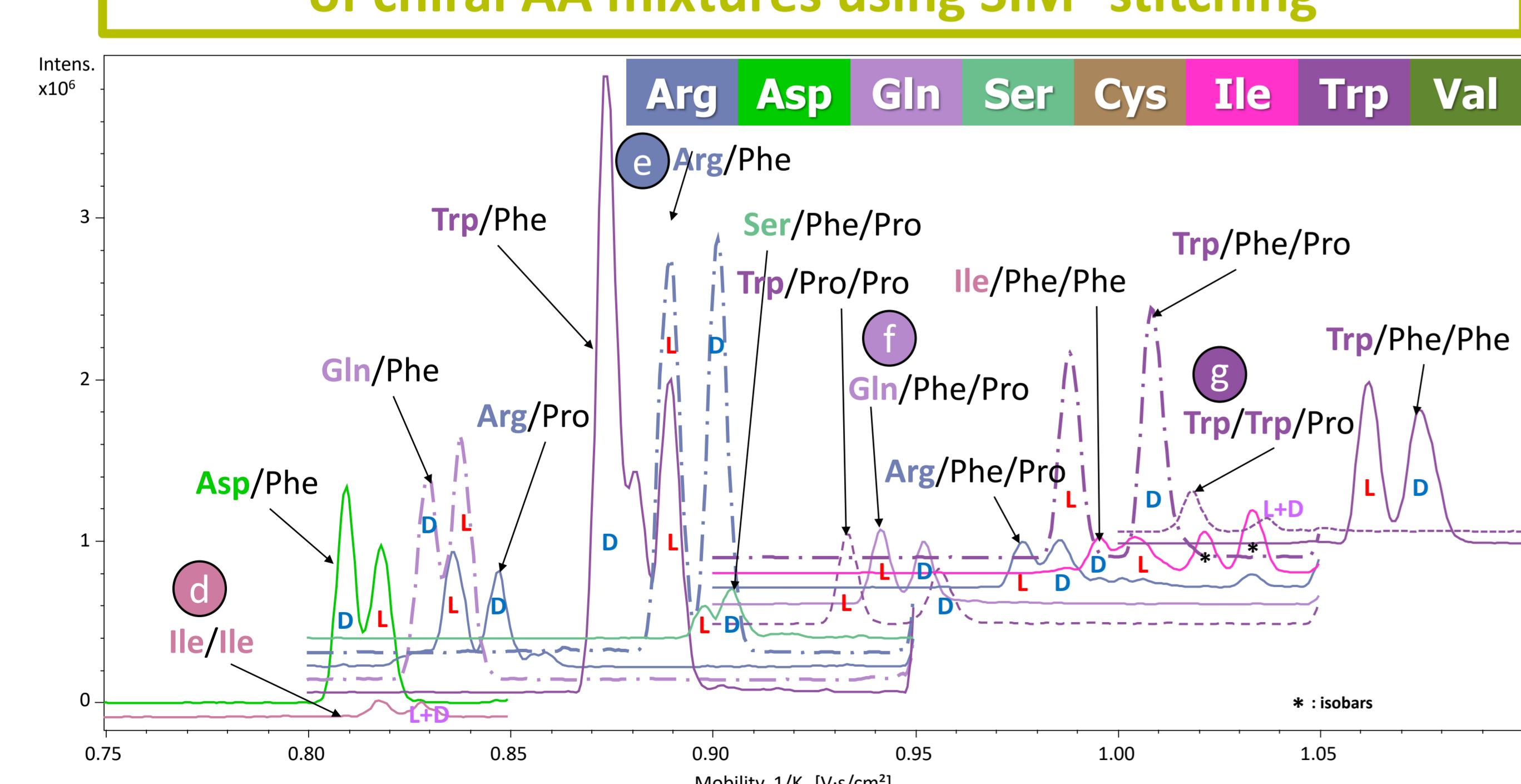


Fig. 1: Reduced mobility plot (above) of the resolved non covalent complexes of L- and D-AA with <sup>1</sup>Phe, <sup>1</sup>Pro and Cu<sup>II</sup> in individual enantiomer solutions using SIM<sup>2</sup> mode. (right) Extracted ion mobilograms (EIMs) of selected complexes of chiral L- or D-AAs and Cu<sup>II</sup>

### High resolution and high throughput IM analysis of chiral AA mixtures using SIM<sup>2</sup> stitching



SIM<sup>2</sup> provides the highest mobility resolution and high sensitivity on a limited range, while SIM<sup>2</sup> stitching\* maintains these performances on a broad mobility range

\* SIM<sup>2</sup>-stitching: collection of multiple adjacent SIM<sup>2</sup> ranges

More details about TIMS and SIM<sup>2</sup> stitching in poster by Chalet et al.

## CONCLUSION

- Most proteinogenic L- and D-AAs can be discriminated using non-covalent complexes with Phe, Pro and Cu<sup>II</sup> in positive mode with at least one combination of references and analyte and/or through self-association.
- Chiral separations are successful for chiral AA mix solutions, although not all complexes of interest may be detected. SIM<sup>2</sup> stitching acquisition ensures high resolution and high throughput onto large reduced mobility ranges.
- Chiral analyses of other biologically relevant molecules using complexation were less efficient. The choice of references, adducts and analytical conditions is thus important to ensure proper chiral separations with NCCs.

## MATERIALS

- Direct infusion of chiral compounds with addition of Cu<sup>II</sup> and <sup>1</sup>Phe & <sup>1</sup>Pro as chiral references in H<sub>2</sub>O/MeOH (5 μM).
- Chiral mix solutions consists of each AA at 1 μM, references at 10 μM and Cu<sup>II</sup> at 20 μM.
- TIMS-ToF:** ESI in positive and negative mode – **SIM<sup>2</sup>** with maximum mobility resolving power (narrow mobility range detection centered on each ion of interest)
- Mobility separation evaluated with:

$$\Delta CCS\% = 2 \frac{(CCS_B - CCS_A)}{(CCS_A + CCS_B)} \times 100$$

### Chiral analysis of natural products

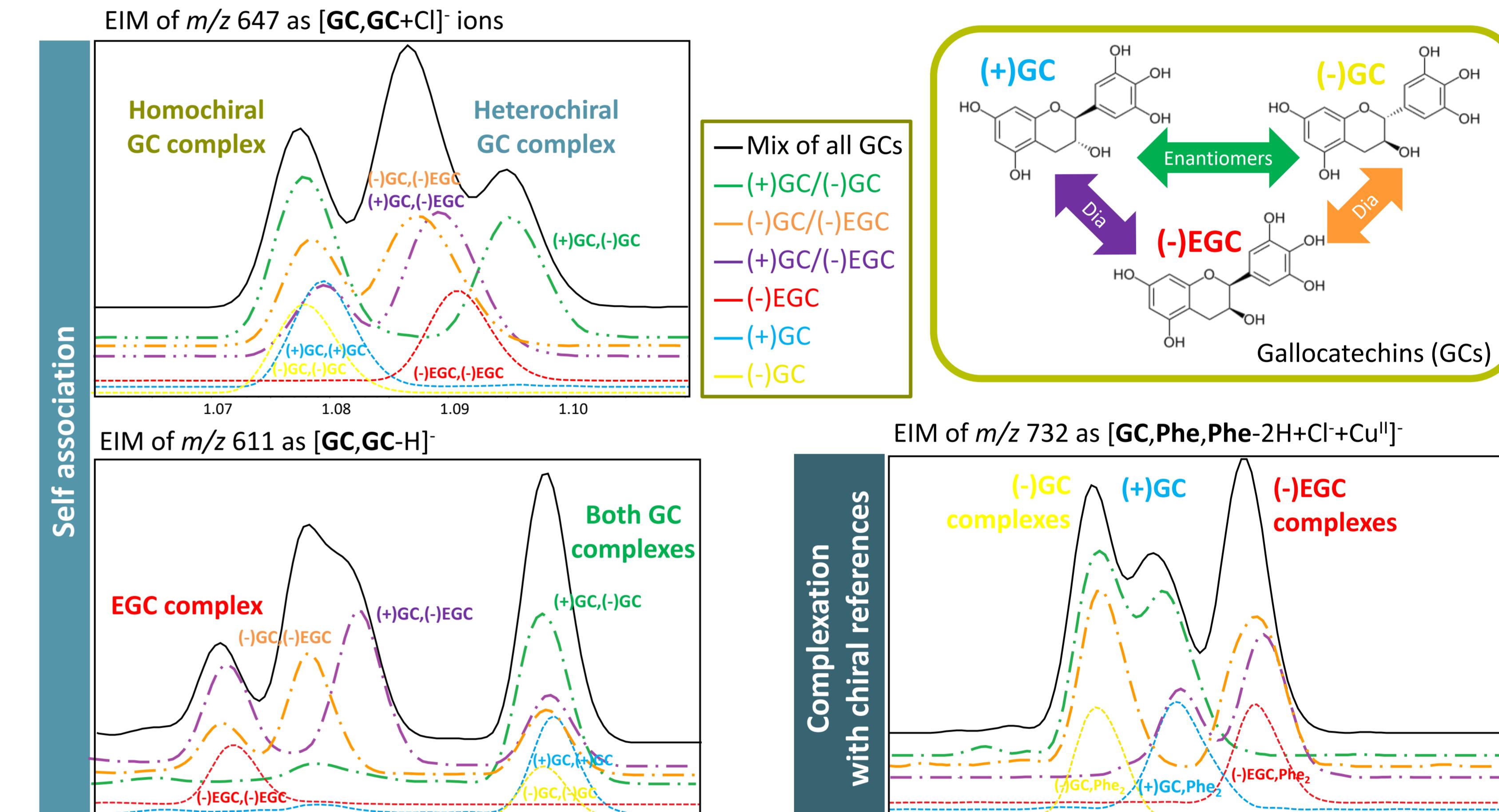


Table 2: Chiral separations expressed as ΔCCS% values obtained for biologically relevant chiral molecules (structures above and right) in individual or mix solutions; SIM<sup>2</sup> mode, positive and negative ionization. (x) No complex detected; (-) Complex detected but no mobility separation; ( ) no tested

Complex	ESI	Adducts	(-)-GC (-)-EGC	(+)-GC (-)-EGC	(+)-GC (-)-GC	All GCs	MA	PhG	EQ	LA	EPD
X,Pro	POS	+Cu <sup>II</sup>	-	-	-	-	1.2%	1.0%	( )	( )	-
	NEG	+Cu <sup>I</sup>	( )	( )	( )	( )	( )	( )	( )	( )	-
		+Cu <sup>II</sup>	3.1%	2.8%	-	2.8%	1.8%	-			
X,Phe	POS	+Cu <sup>I</sup>	2.8%	1.8%	2.4%	2.6%	1.6%	1.4%	1.3%	( )	( )
		+Na <sup>+</sup>	4.5%	4.7%	2.9%	-	4.6%	3.1%			
		+Cu <sup>II</sup>	3.9%	4.6%	2.9%	-	3.9%	0.7%	2.8%	1.8%	2.3%
		+Cu <sup>I</sup>	2.8%	4.9%	1.1%	1.4%	2.7%	2.1%	1.1%	( )	( )
		-H <sup>-</sup>	0.9%	1.4%	2.0%	-	0.9%	2.4%	-		
		+Cl <sup>-</sup>	0.8%	0.9%	1.6%	-	0.9%	0.7%			
X,Pro,Pro	POS	+Cu <sup>II</sup>					x	0.9%	( )	( )	-
	POS	+Cu <sup>I</sup>					-	0.9%			
X,Phe,Pro	NEG	+Cu <sup>II</sup> Cl	1.0%	1.0%	-	1.0%	( )	( )	-	-	-
	POS	+Cu <sup>II</sup>					x	-	( )	( )	-
X,Phe,Phe	NEG	+Cu <sup>II</sup> Cl	1.6%	0.9%	0.7%	0.6%	( )	( )	-	-	-
	+Cu <sup>I</sup>										
	+Cl <sup>-</sup>										
X,X,Pro	POS	+Cu <sup>II</sup>					x	1.5%	( )	( )	-
X,X,Phe	POS	+Cu <sup>II</sup>					x	-	( )	( )	-
	NEG	+Cu <sup>II</sup>					( )	( )	-	-	
X,X,X	POS	+Cu <sup>II</sup>					-	1.4%	( )	( )	-
	NEG	-H <sup>-</sup>	2.7%	2.2%	-	2.4%	( )	( )			

## REFERENCES

- [1] S. Yang et al.; Talanta, **243** (2022)
- [2] J. Czekner et al.; J Am Mass Spectrom, **33** (2022)
- [3] E. Rathahao-Paris et al.; J Mass Spectrom, **57** (2022) 0017 and Metabohub infrastructure (ANR-11-INBS-0010)

## ACKNOWLEDGMENTS

This research was performed within funding from the CHIRAMICS project (ANR-20-CE29-0017) and Metabohub infrastructure (ANR-11-INBS-0010)