



EURACHEM 2019 TARTU

QUANTITATIVE NON-TARGETED ANALYSIS

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Green tea is one of the most commonly consumed beverages worldwide and is claimed to possess numerous health-protective qualities.

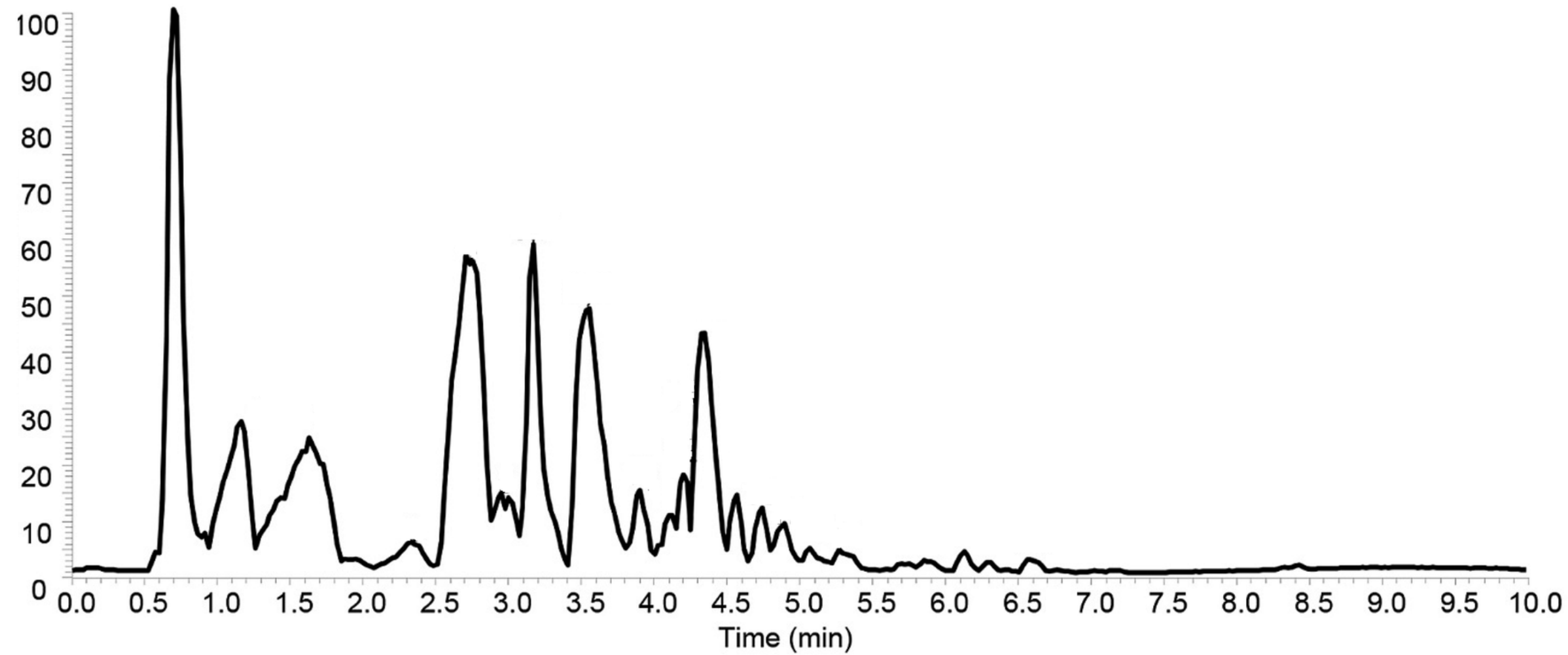


Metabolomic study of green tea

LC/HRMS

aims to correlate changes in the chemical profile of a sample with a corresponding shift in macroscopic phenotype





Non-targeted analysis

LC/ESI/HRMS

2270 marker ions

38 tea samples

19 identified compounds

QUALITATIVE

stop after identification

QUANTITATIVE

purchase or synthesize the standard substances



Quantification

Preliminary assignment

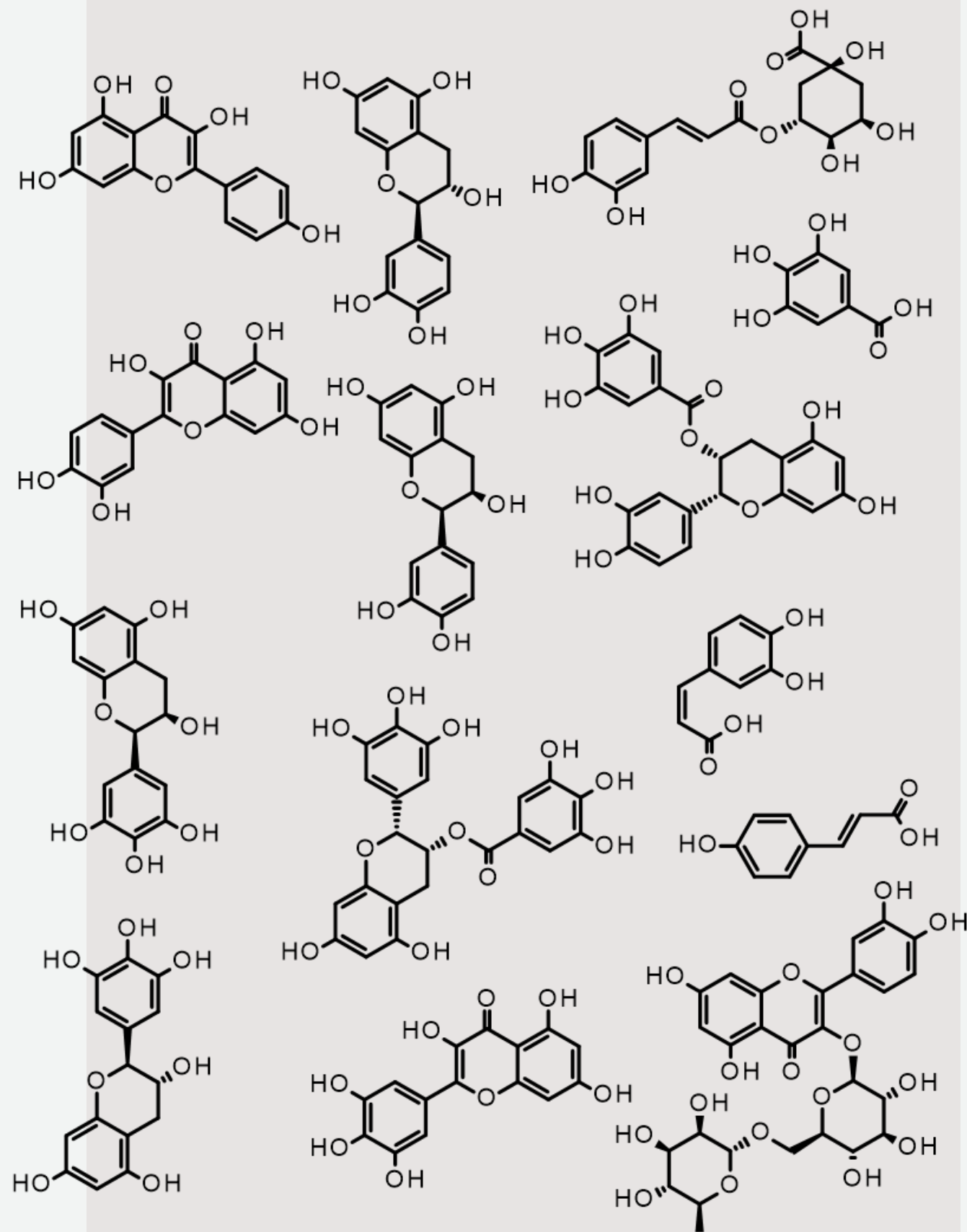
with non-targeted LC/HRMS

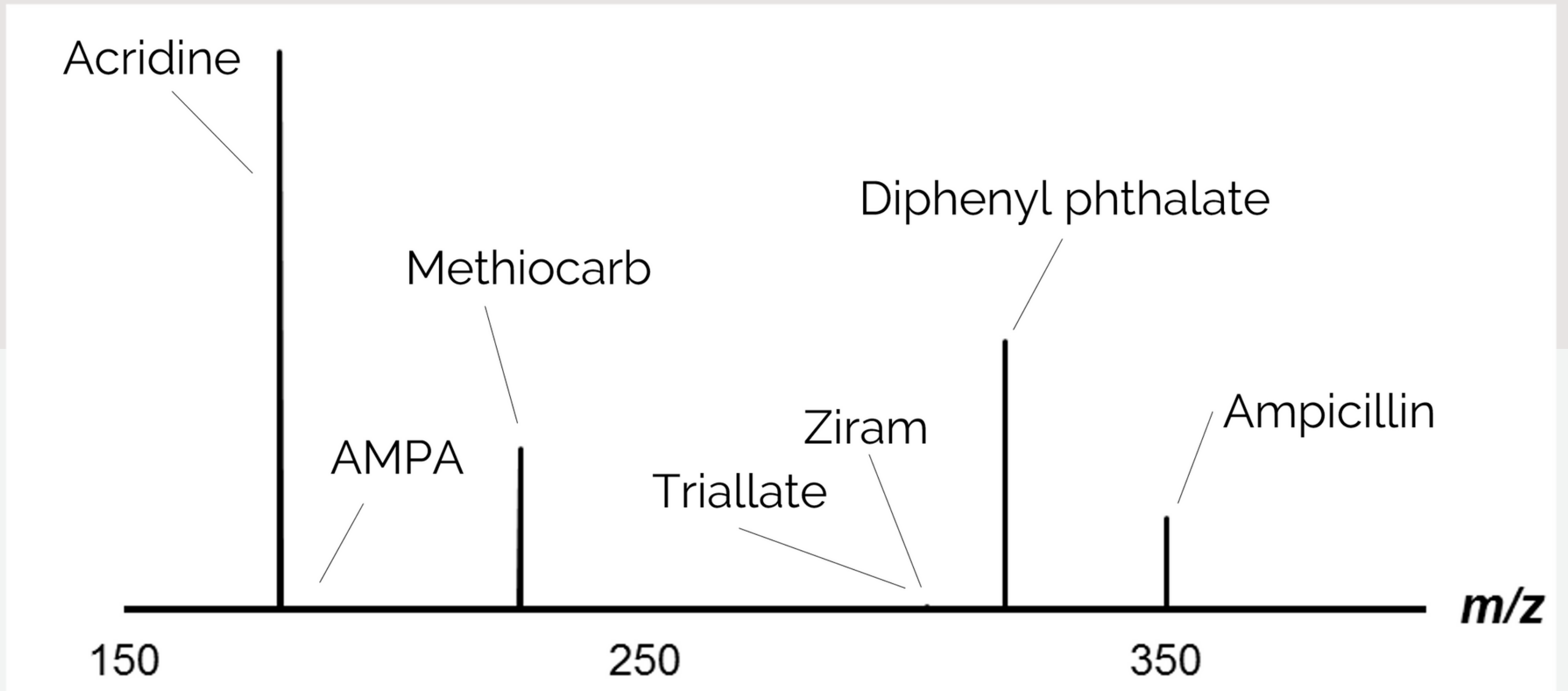
Standard substances

were available for 14 compounds

Quantification possible

with calibration graph method





Mixture of seven compounds at equal concentrations.

Climbing the way to quantitative non-targeted analysis





Ionization efficiency

for ESI source



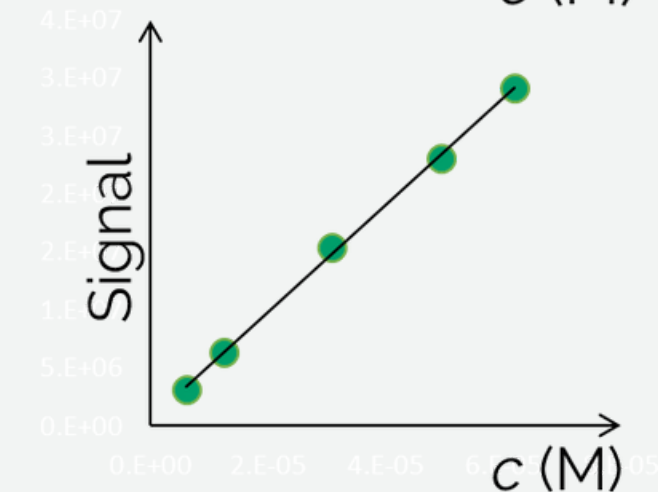
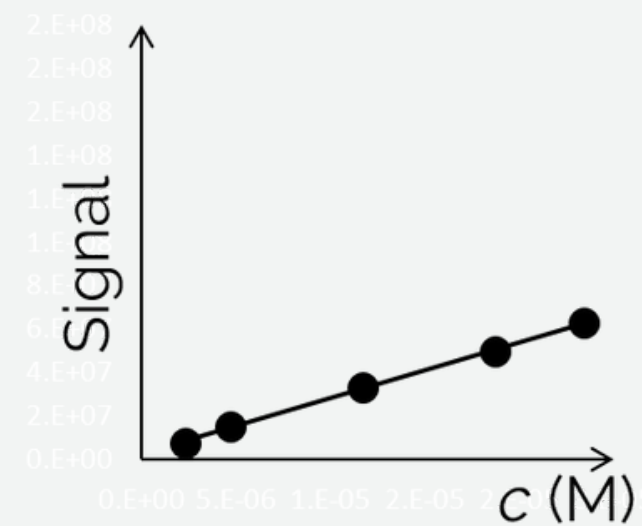
Ionization efficiency

RELATIVE MEASUREMENTS

Flow injections



Calibration graph



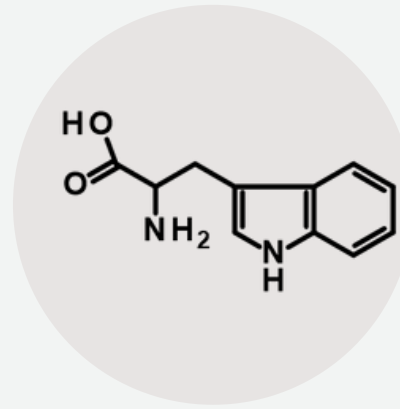
Calculation

$$\log \frac{\text{slope}_1}{\text{slope}_2} \rightarrow \log IE$$



Ionization efficiency

IS AFFECTED BY



Compound structure

hydrophobicity, basicity/acidity

Eluent

organic modifier %, pH, additive type

Instrument

source design, mass analyzer

Matrix

ionization suppression/enhancement

Structure

Hydrophobicity

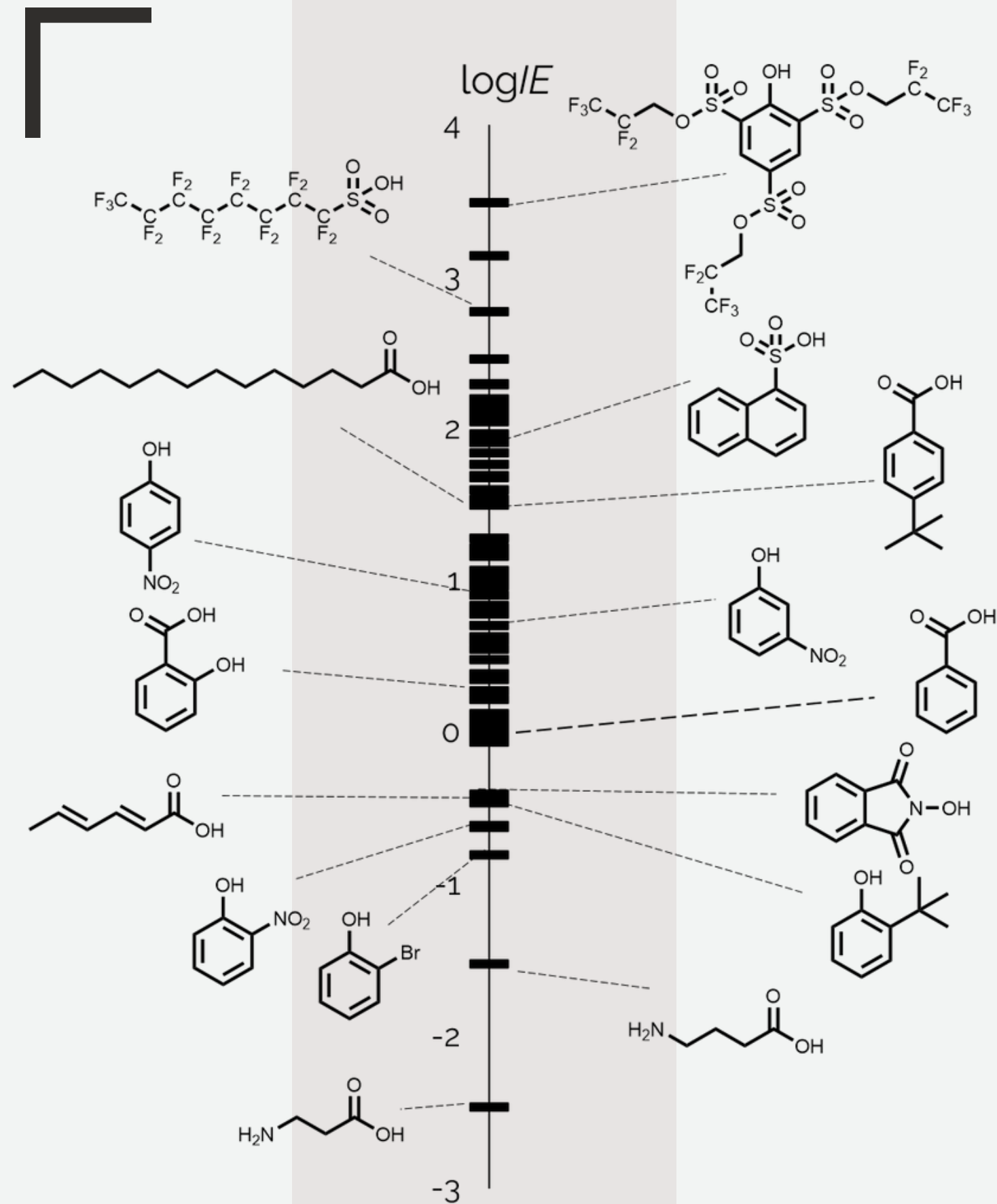
affinity for the droplets surface

Acidity/Basicity

protonation/deprotonation in the solution

Interplay between properties

no single ESI mechanism



Structure

Hydrophobicity

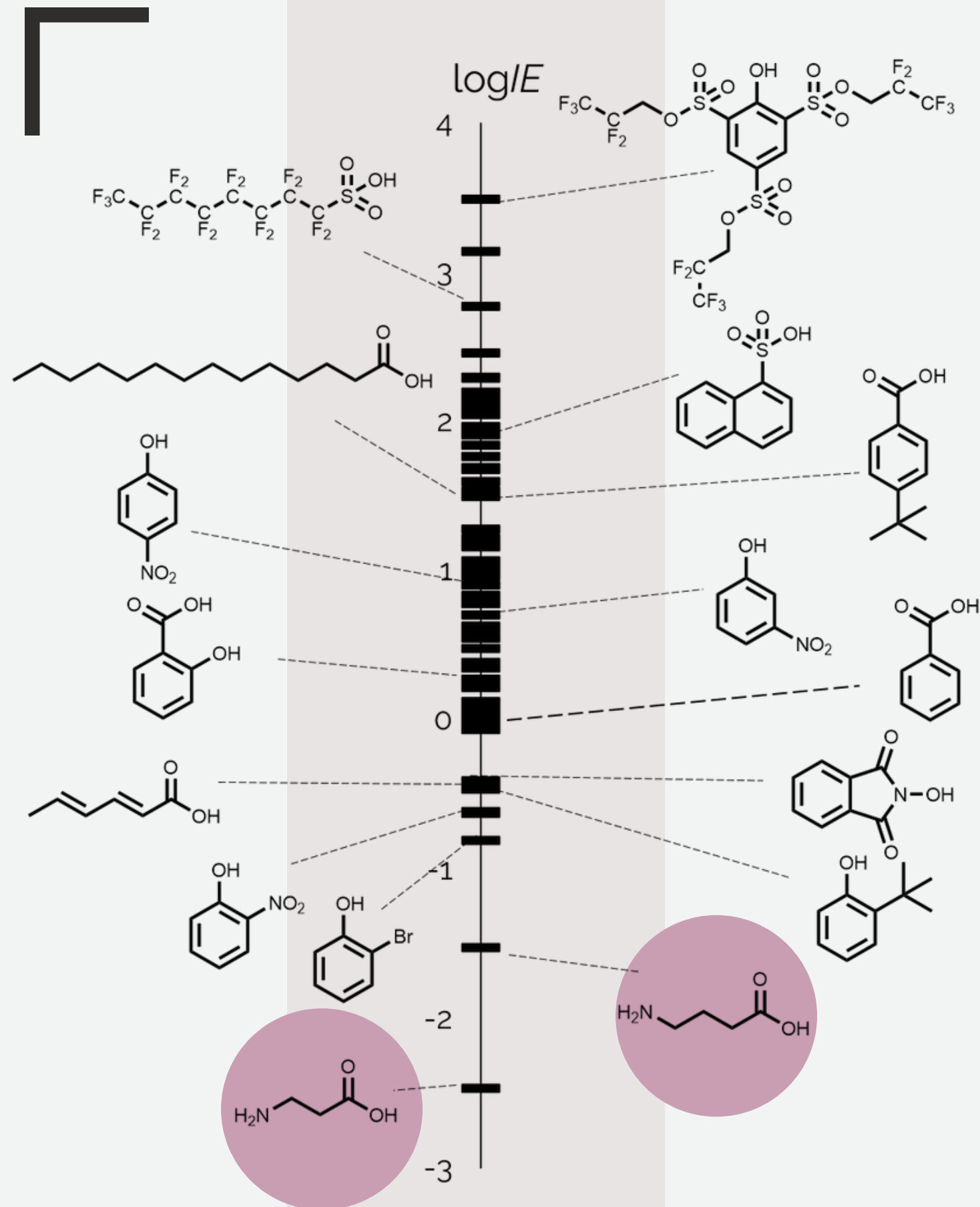
affinity for the droplets surface

Acidity/Basicity

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Interplay between properties

no single ESI mechanism



Structure

Hydrophobicity

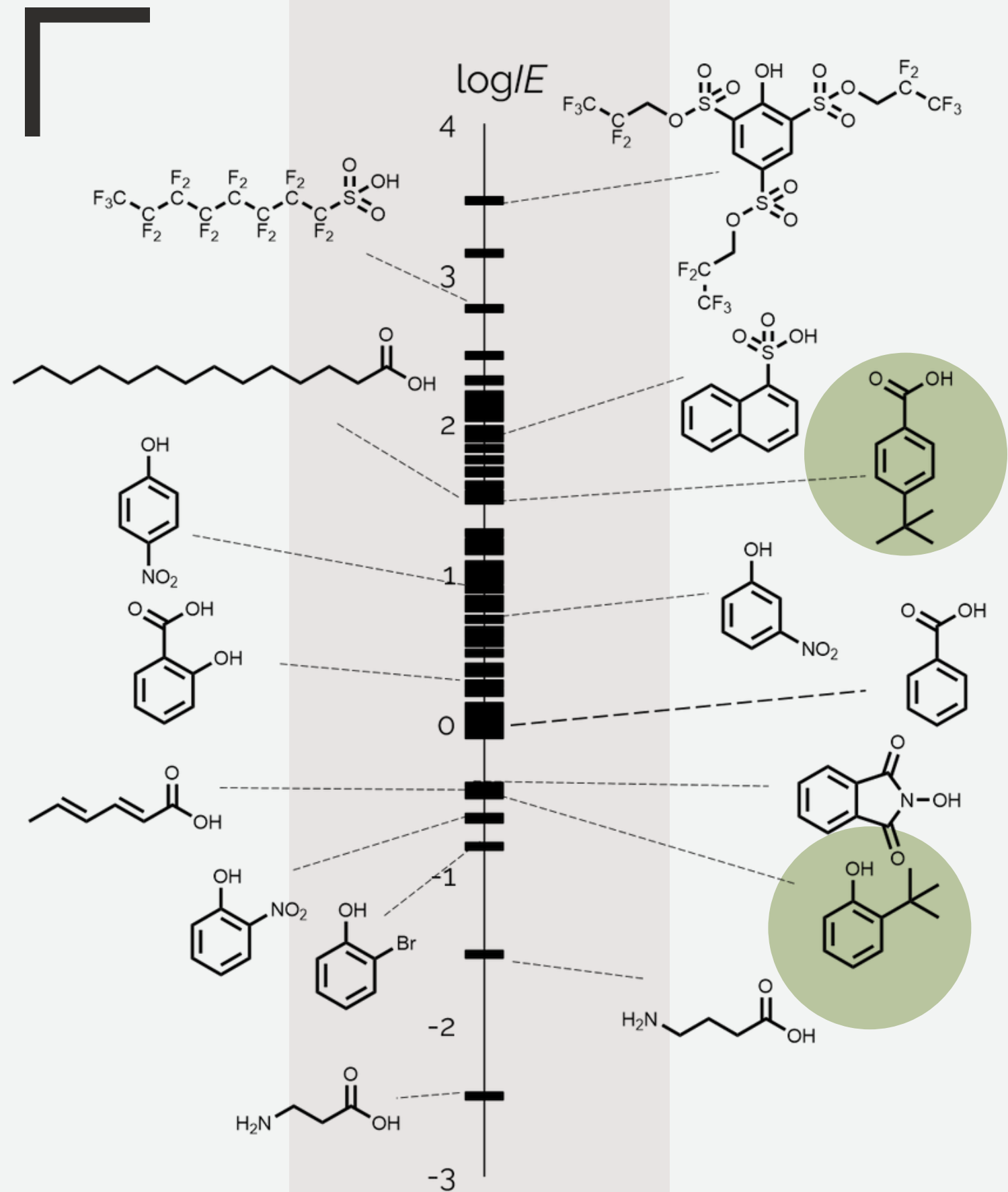
affinity for the droplets surface

Acidity/Basicity

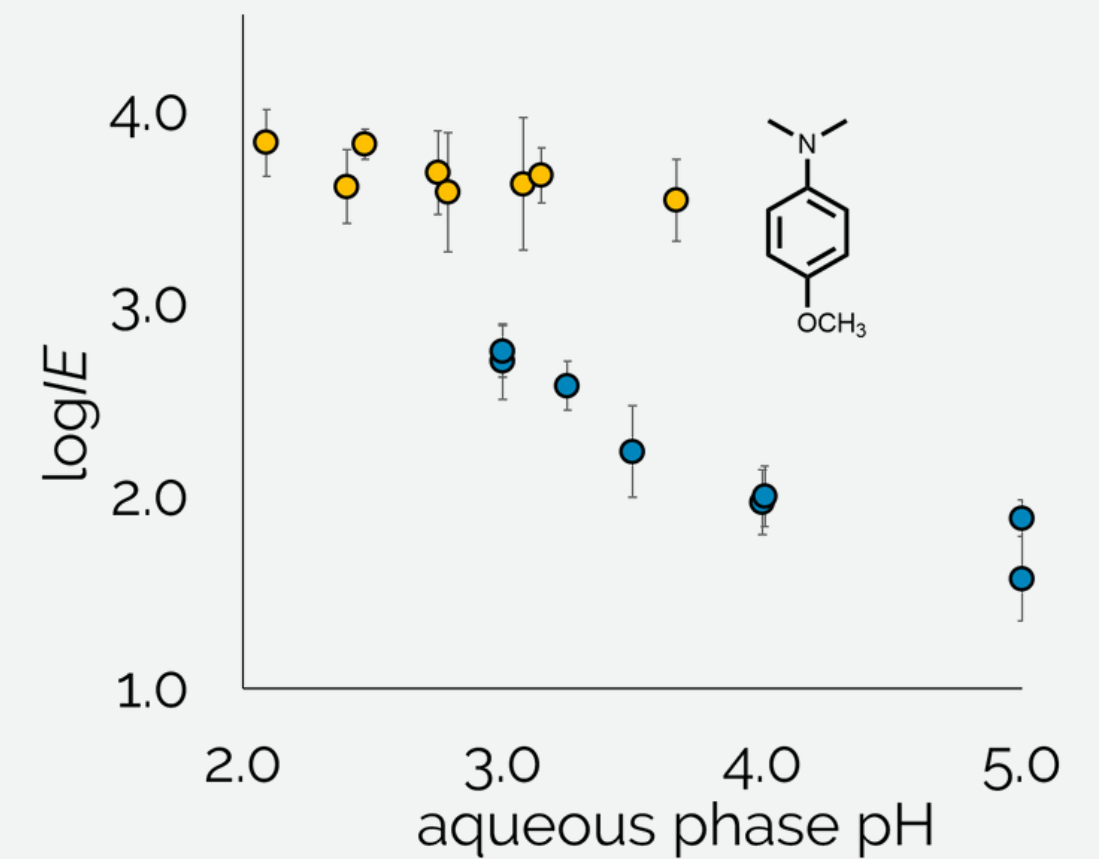
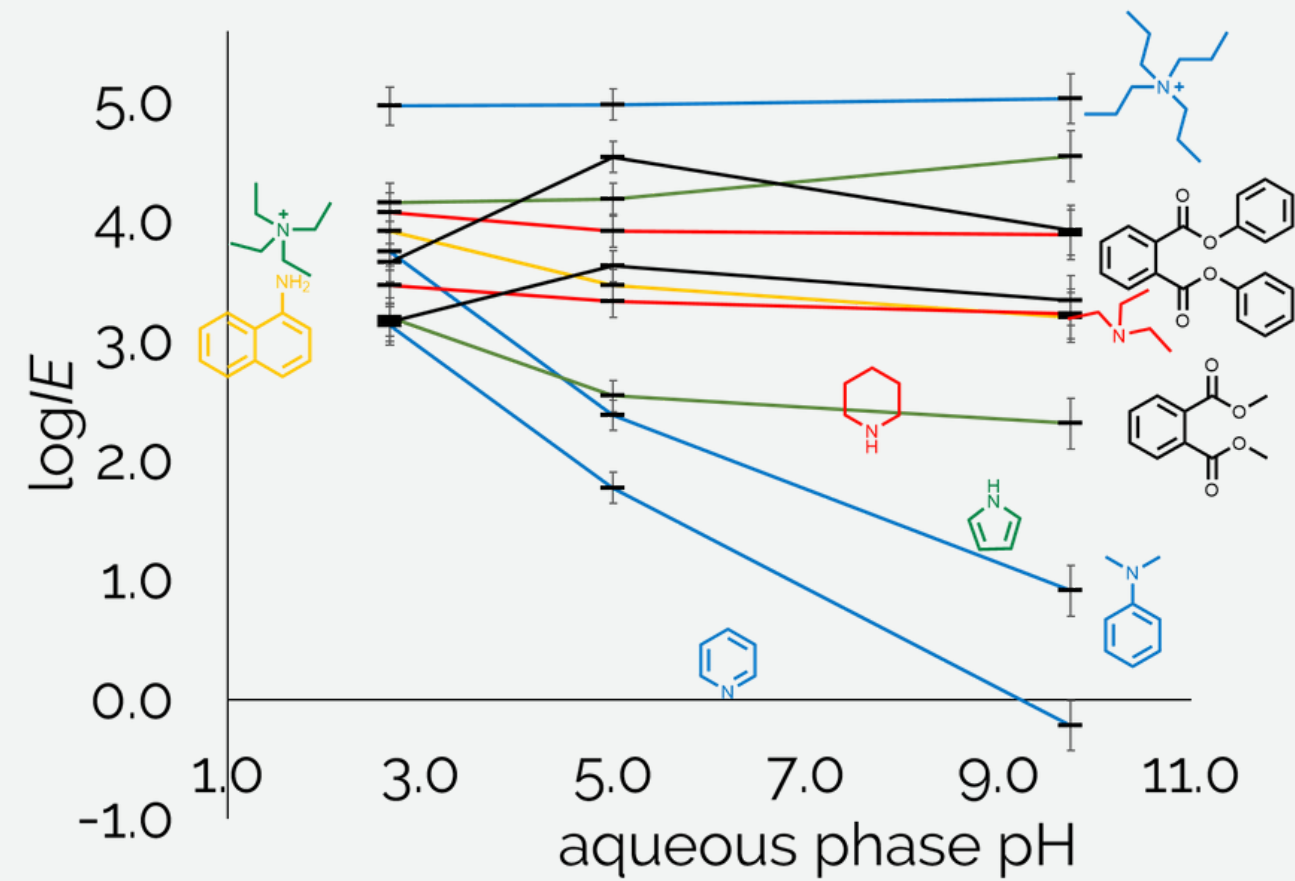
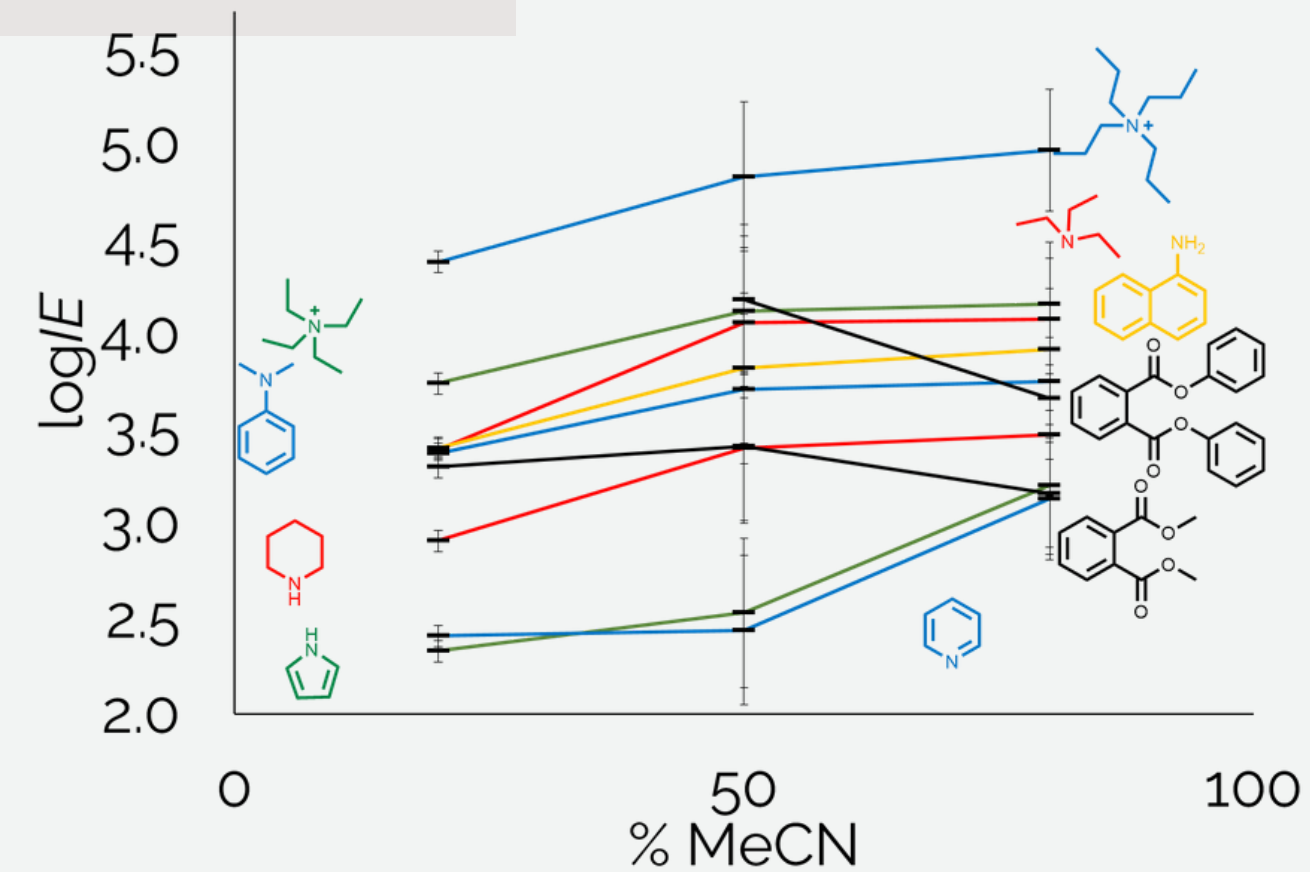
protonation/deprotonation in the solution

Interplay between properties

no single ESI mechanism



Eluent



Organic modifier

higher organic modifier content increases ionization efficiency by >10x

pH

change in ionization efficiency >1000x

Additive type

same pH, different buffer: 20 to 50x different ionization efficiencies

Liigand et al. DOI: 10.1007/s13361-014-0969-x

Liignad et al. DOI: 10.1007/s13361-016-1563-1

Ojakivi et al. DOI: 10.1002/slct.201702269.

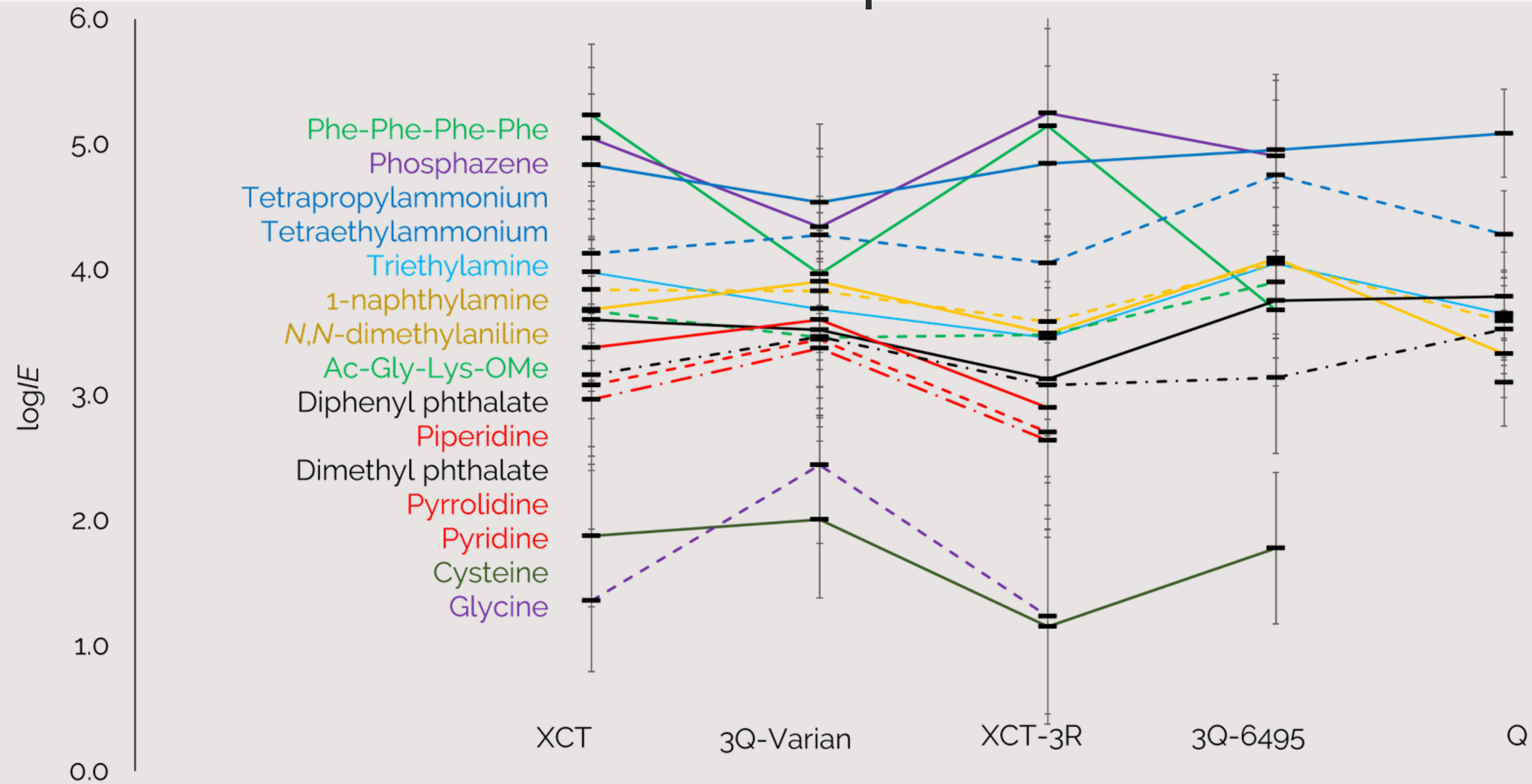
Anneli Kruve 21.05.2019

DIFFERENT INSTRUMENTS

7 different instruments
5 vendors
4 countries

IONIZATION EFFICIENCY

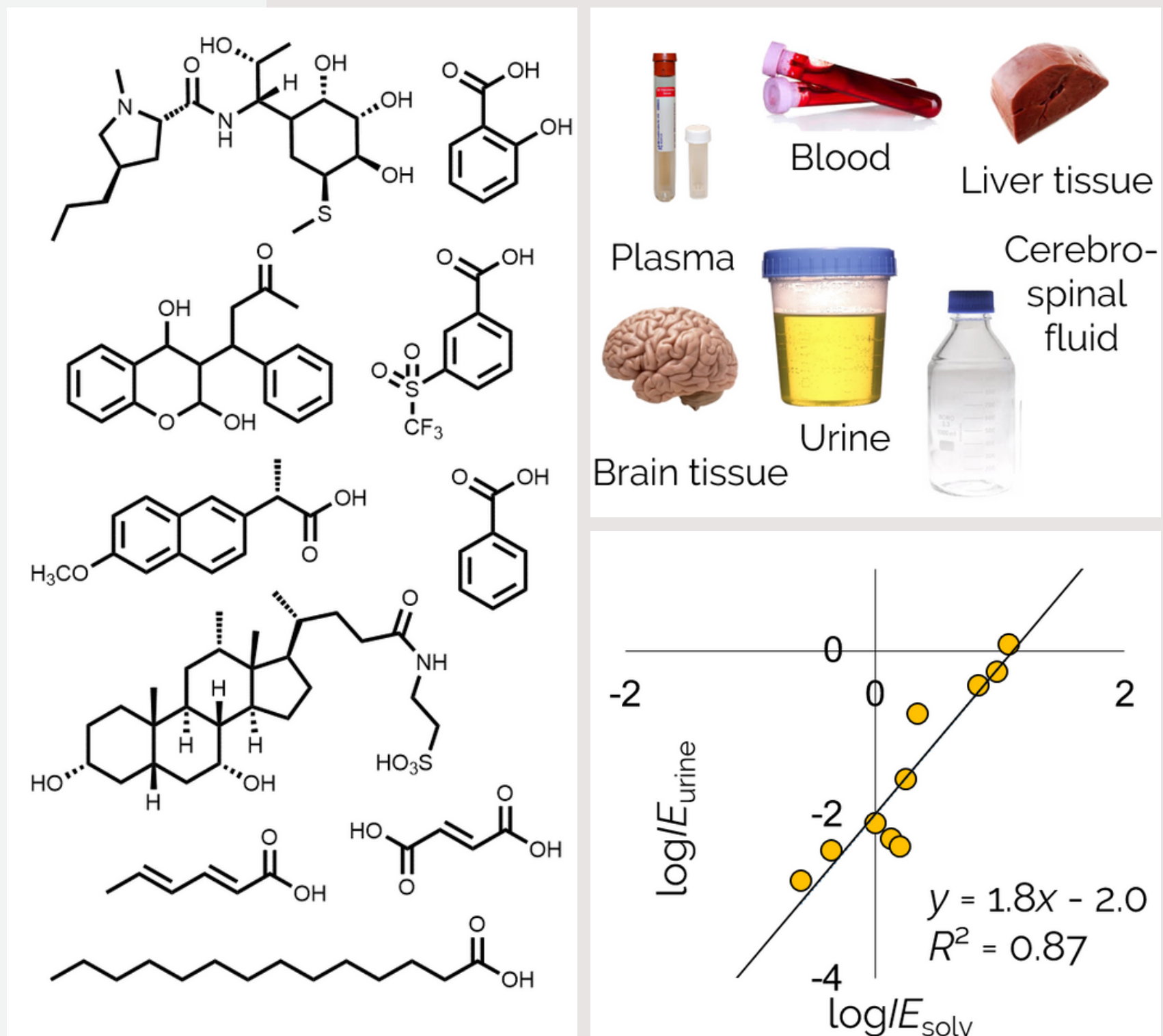
order of compounds is similar
good correlation



Matrix effect in ESI

BIOLOGICAL SAMPLES

order of compounds is the same
good correlation





Modelling

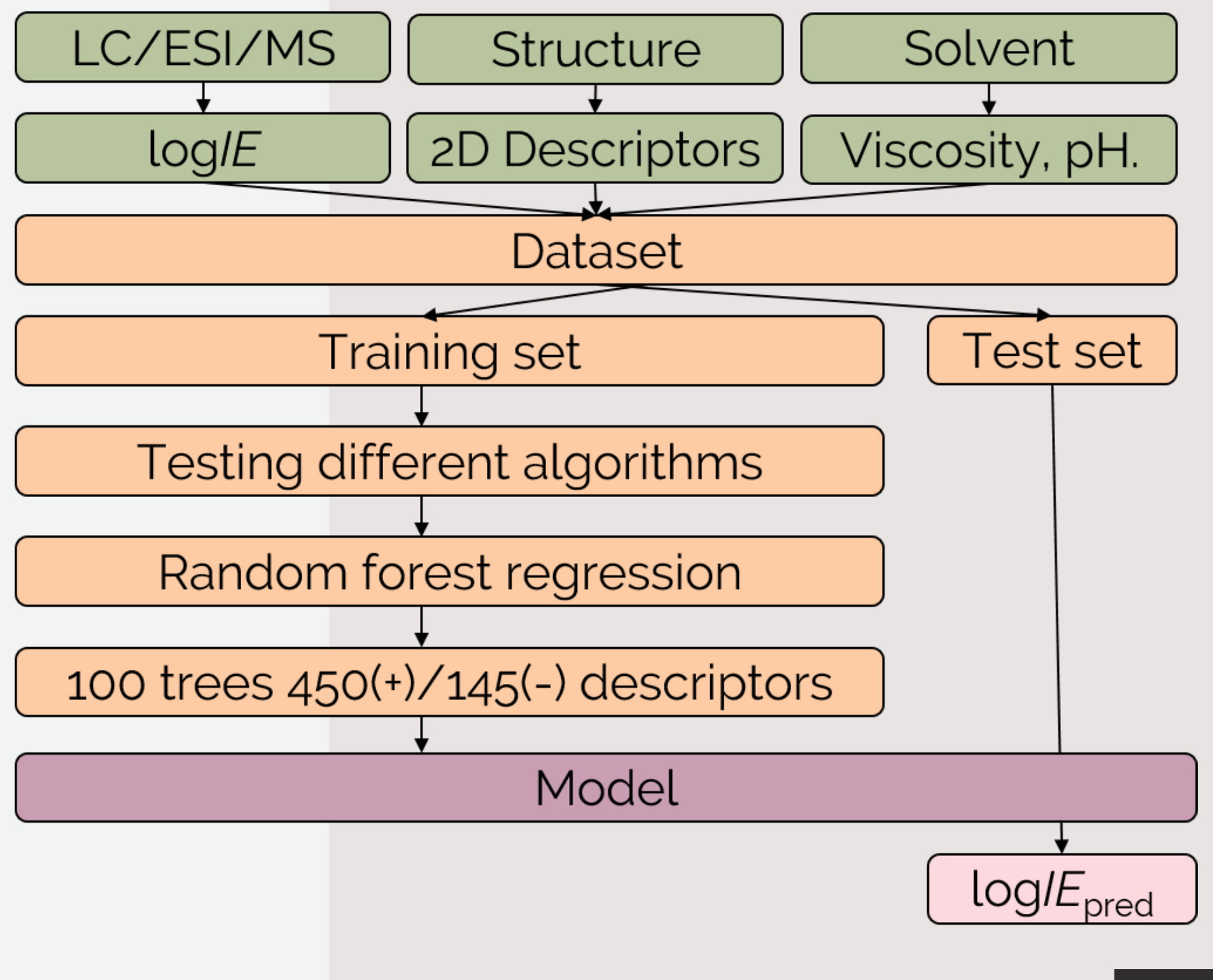
bringing previous knowledge together



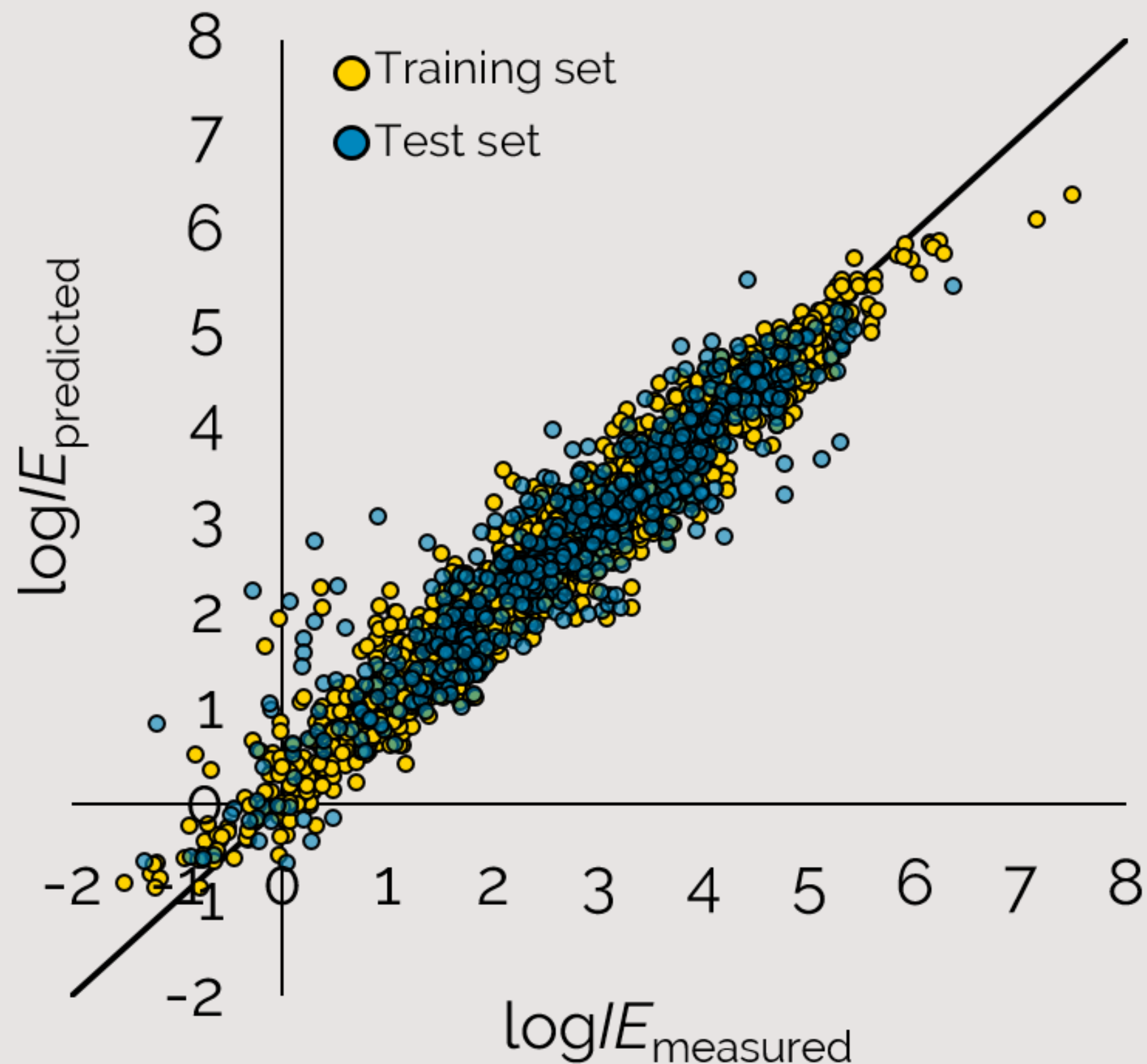
Modelling of ionization efficiency

TRAINING & VALIDATION

different machine learning approaches tested



ESI+



3139 data points

353 small molecules

106 eluent compositions

MeCN/MeOH 0 – 100%

pH = 2.0 – 10.7

Average prediction error

Training set 1.9x

Test set 3.0x

ESI-

1286 data points

101 small molecules

33 eluent compositions

MeCN/MeOH 0 – 100%

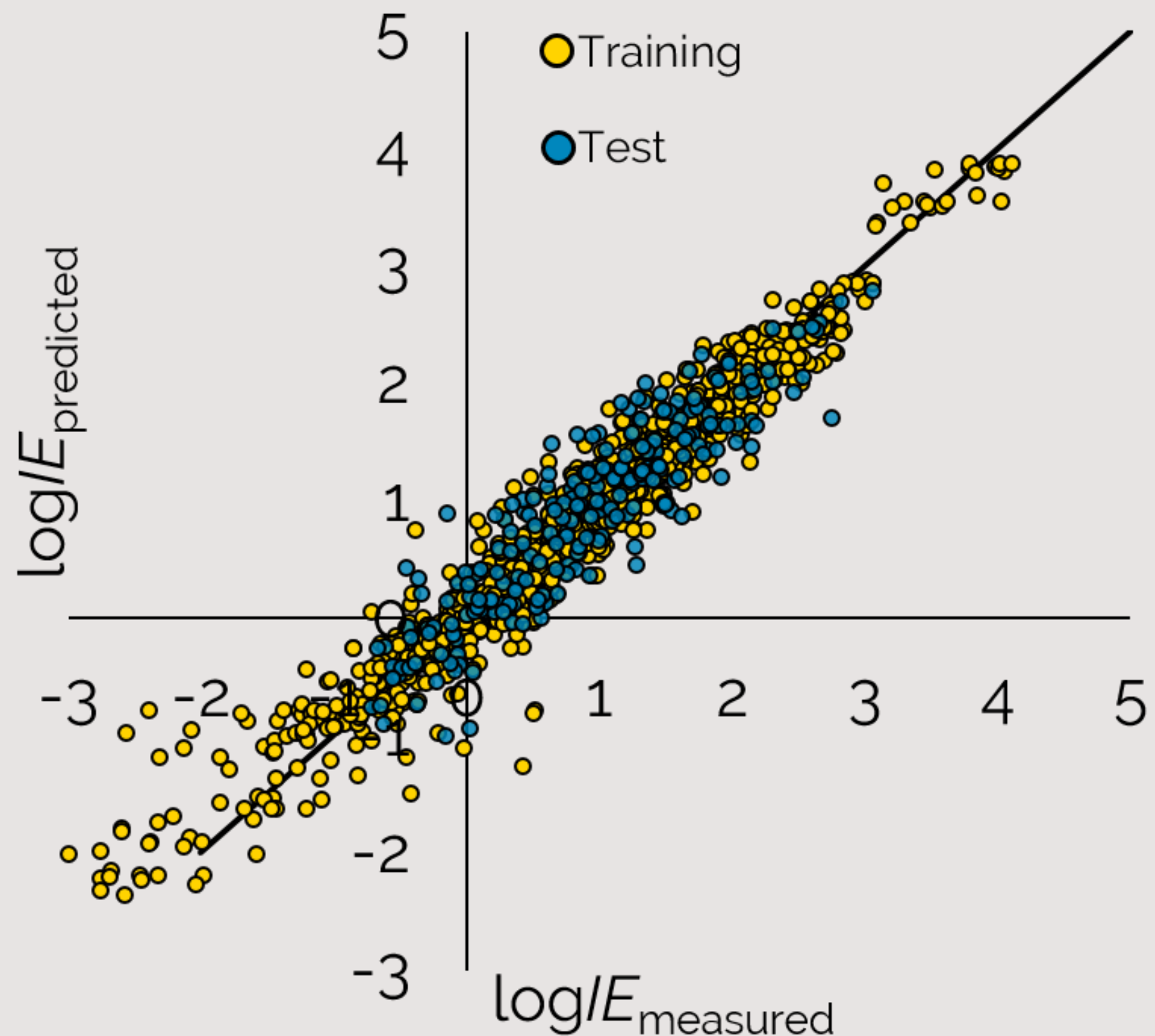
pH = 2.0 – 10.7

Average prediction error

Training set 2.0x

Test set 2.3x

Liignad et al. submitted for publication





Concentrations

application for green tea



How to use it for quantification?



Step 1

predict ionization efficiencies



Step 2

transfer to specific setup



Step 3

calculate



-
- (1) SMILES for identified compounds
 - (2) 5-6 compounds with known concentration
 - (3) gradient parameters
- READY...SET...GO

How does it work in practice?





- (1) SMILES for identified compounds
 - (2) 5-6 compounds with known concentration
 - (3) gradient parameters
- READY...SET...GO

LC/MS web-app

[Download sample input file](#)

Upload a CSV file

pH

Enter a value from 0 to 14.

Solvents

Solvent A

Solvent B

Gradient

T (min)	A %	B %	
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="button" value="Delete"/>





sample - Excel

File Home Insert Page Layout Formulas Data Review View Tell me what you want to do... Anneli Kruve Share

Clipboard Font Alignment Number Styles Cells Editing

D14

	A	B	C	D	E	F	G	H	I	J	K	L	M
1	Compound SMILES	Retention tin	Peak Area	concentration									
2	<chem>CC1=C(N=C(N=C1OC(=O)N(C)C)N</chem>	1.771	7.42E+08										
3	<chem>CN1C(=C(C(=N1)Cl)C(=O)OC)S(=O</chem>	4.982	1.01E+08										
4	<chem>CCOP(=S)(OCC)SCN1C2=C(C=C(C</chem>	6.041	1.66E+07										
5	<chem>COP(=O)(OC)OC1=CC=C(C=C1)[N</chem>	3.492	6.16E+07										
6	<chem>CCOP(=O)(OCC)OP(=O)(OCC)OC</chem>	3.171	1.25E+08										
7	<chem>COP(=O)(N)SC</chem>	0.877	6.22E+06	3.55E-06									
8	<chem>CCCCN(CCCC)SN(C)C(=O)OC1=CC</chem>	7.622	2.88E+07	2.63E-07									
9	<chem>CCCC(CN1C=NC=N1)(C2=C(C=C</chem>	5.263	3.03E+07	1.60E-06									
10	<chem>CCC1=CC=CC(=C1N(COCC)C(=O)C</chem>	5.371	9.36E+06	1.86E-06									
11	<chem>COC(=O)N(C1=CC=CC=C1OC2=N</chem>	5.917	1.16E+08	1.29E-06									
12													
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15													
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- (1) SMILES for identified compounds
 - (2) 5-6 compounds with known concentration
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- (1) SMILES for identified compounds
 - (2) 5-6 compounds with known concentration
 - (3) gradient parameters
- READY...SET...GO

LC/MS web-app

[Download sample input file](#)

Upload a CSV file

pH

Enter a value from 0 to 14.

Solvents

Solvent A

Solvent B

Gradient

T (min)	A %	B %	
<input type="text" value="0"/>	<input type="text" value="95"/>	<input type="text" value="5"/>	<input type="button" value="Delete"/>
<input type="text" value="15"/>	<input type="text" value="0"/>	<input type="text" value="100"/>	<input type="button" value="Delete"/>
<input type="text" value="20"/>	<input type="text" value="0"/>	<input type="text" value="100"/>	<input type="button" value="Delete"/>
<input type="text" value="25"/>	<input type="text" value="95"/>	<input type="text" value="5"/>	<input type="button" value="Delete"/>



- (1) SMILES for identified compounds
 - (2) 5-6 compounds with known concentration
 - (3) gradient parameters
- READY...SET...GO

Results

[Download results](#)

Substance	Concentration
<chem>CC1=C(N=C(N=C1OC(=O)N(C)C)NC)C</chem>	4.39e-05
<chem>CN1C(=C(C(=N1)Cl)C(=O)OC)S(=O)(=O)NC(=O)NC2=NC(=CC(=N2)OC)OC</chem>	1.83e-05
<chem>CCOP(=S)(OCC)SCN1C2=C(C=C(C=C2)Cl)OC1=O</chem>	2.64e-06
<chem>COP(=O)(OC)OC1=CC=C(C=C1)[N+](=O)[O-]</chem>	2.36e-05
<chem>CCOP(=O)(OCC)OP(=O)(OCC)OCC</chem>	1.7e-05
<chem>COP(=O)(N)SC</chem>	3.55e-06
<chem>CCCCN(CCCC)SN(C)C(=O)OC1=CC=CC2=C1OC(C2)(C)C</chem>	2.63e-07
<chem>CCCC(CN1C=NC=N1)(C2=C(C=C(C=C2)Cl)Cl)O</chem>	1.6e-06
<chem>CCC1=CC=CC(=C1N(COCC)C(=O)CC)C</chem>	1.86e-06
<chem>COC(=O)N(C1=CC=CC=C1OC2=NN(C=C2)C3=CC=C(C=C3)Cl)OC</chem>	1.29e-06

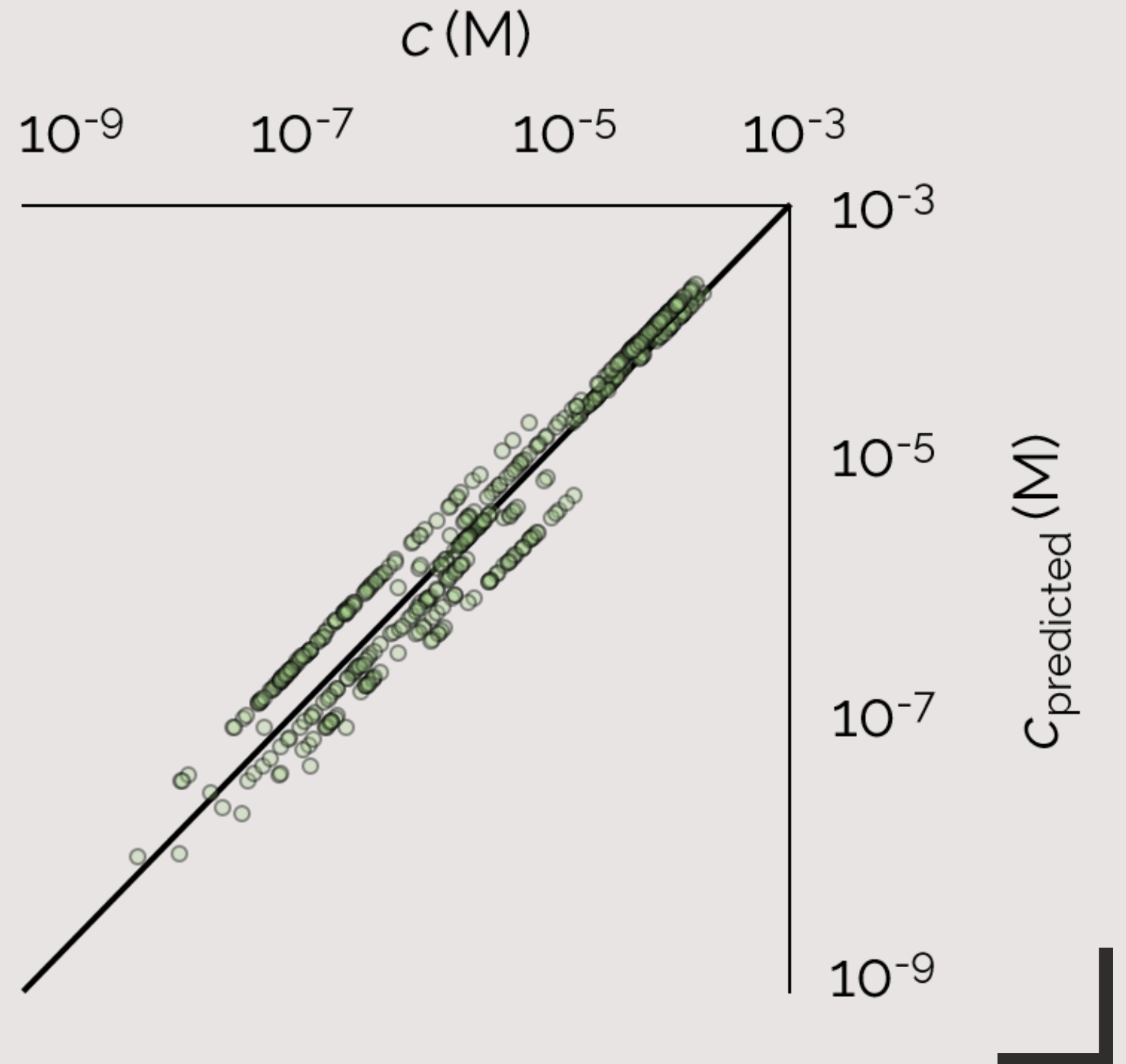
Prediction for green tea

Measurements

UNCG (USA) in 2017

Calculations

UT (Estonia) in late 2018





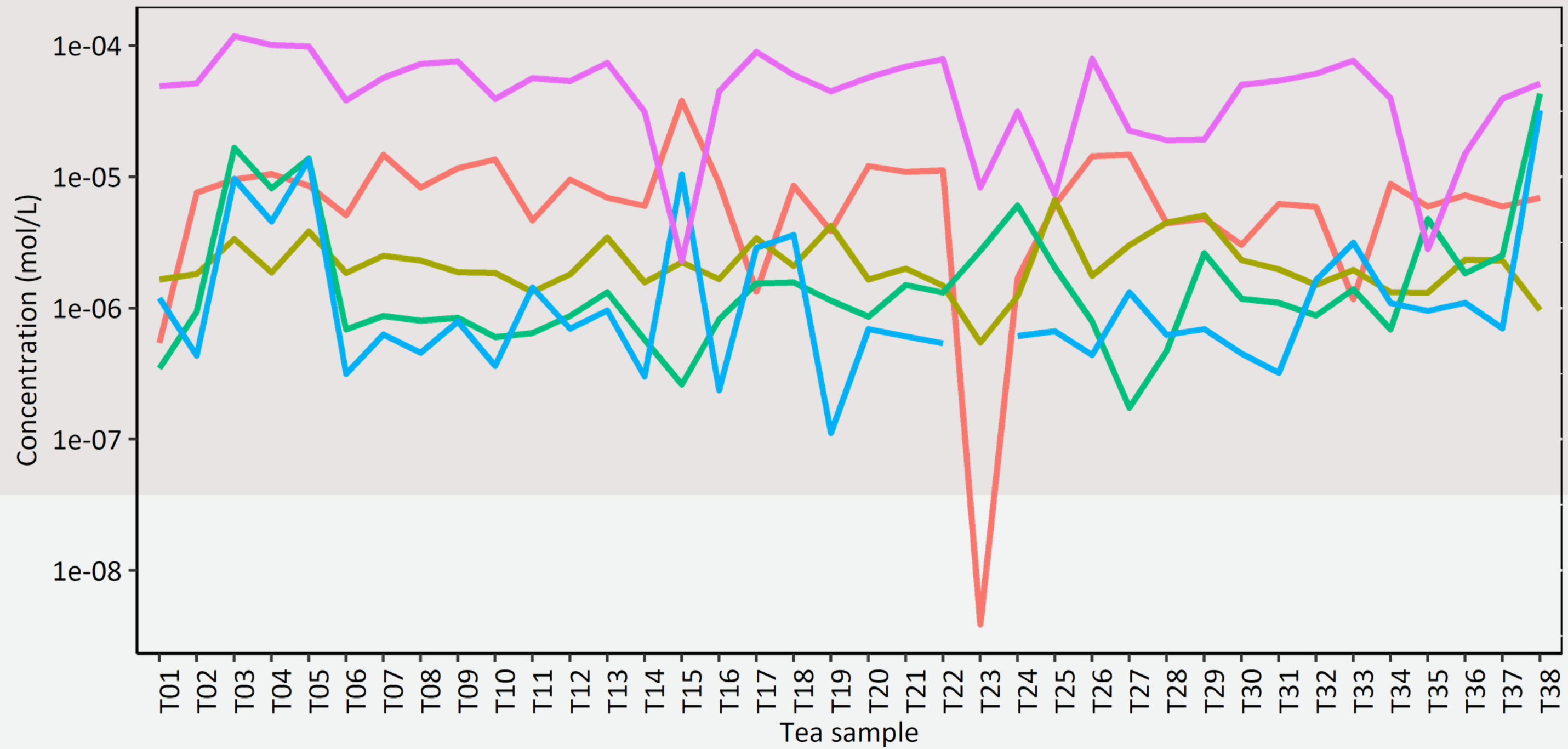
The average prediction
error was

1.7X

predicted concentration
of quercetin $0.8 \mu\text{mol/L}$
the confidence interval
 0.5 to $1.4 \mu\text{mol/L}$



Can the predicted concentrations reveal anything about different tea samples?



Indeed, we can identify the non-green tea samples!



Pesticides in cereals

More than 1000 pesticides and mycotoxins registered in EU.



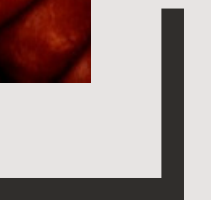
Cereal samples

PESTICIDES AND MYCOTOXINS

35 compounds & 6 transformation
compounds

3.6 nM to 0.35 mM

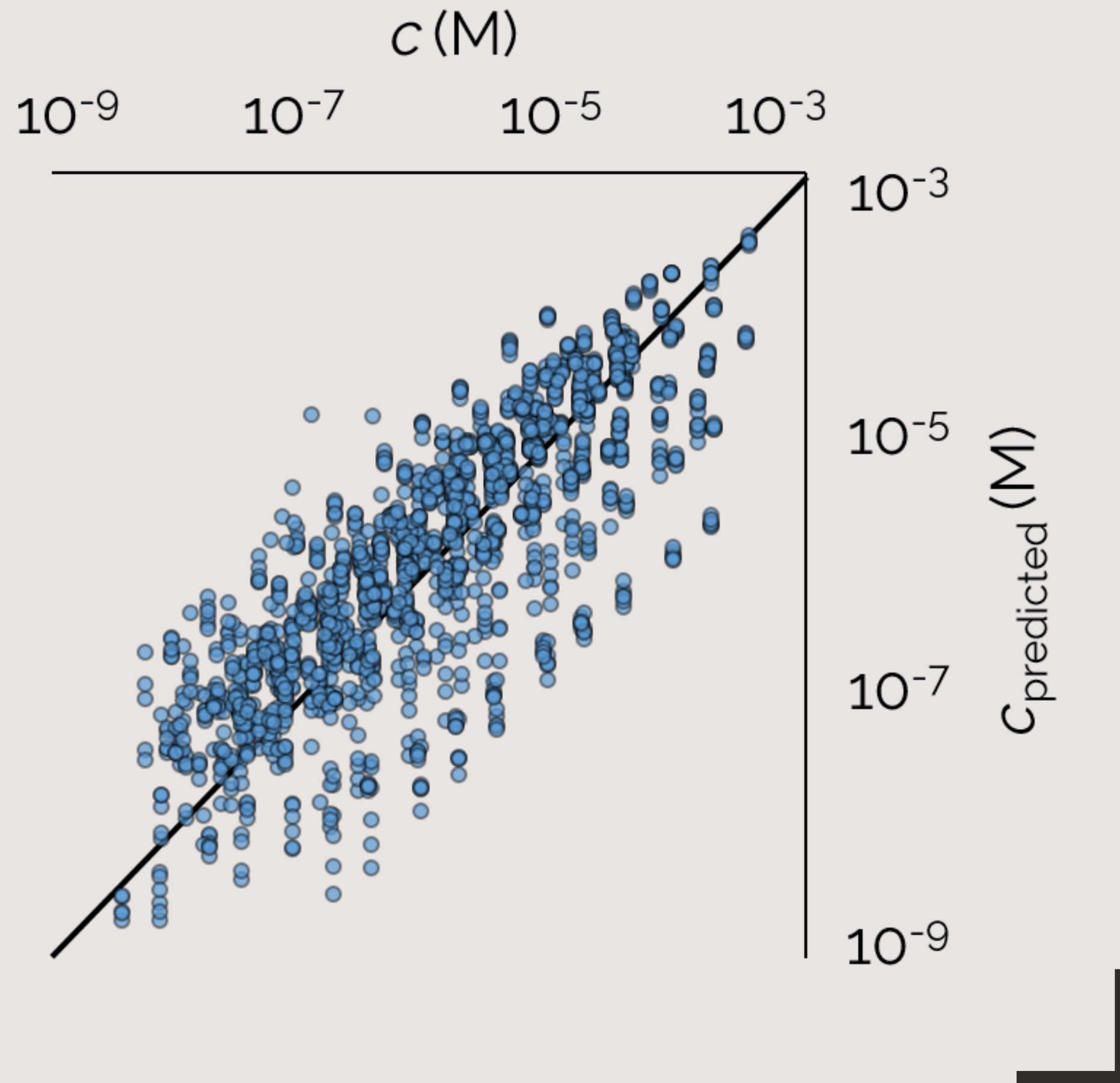
oat, barley, rye, wheat, rice, maize



Cereal samples

PESTICIDES AND MYCOTOXINS

2233 data points
average prediction error 5x.



7 out of 10

prediction error <5x.

9 out of 10

prediction error <10x.

What can we see now?



What does the future hold for non-targeted analysis?



Quantitative results



Directly comparable data from different labs



Retrospective analysis



Quantem
ANALYTICS
quantem.co

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