

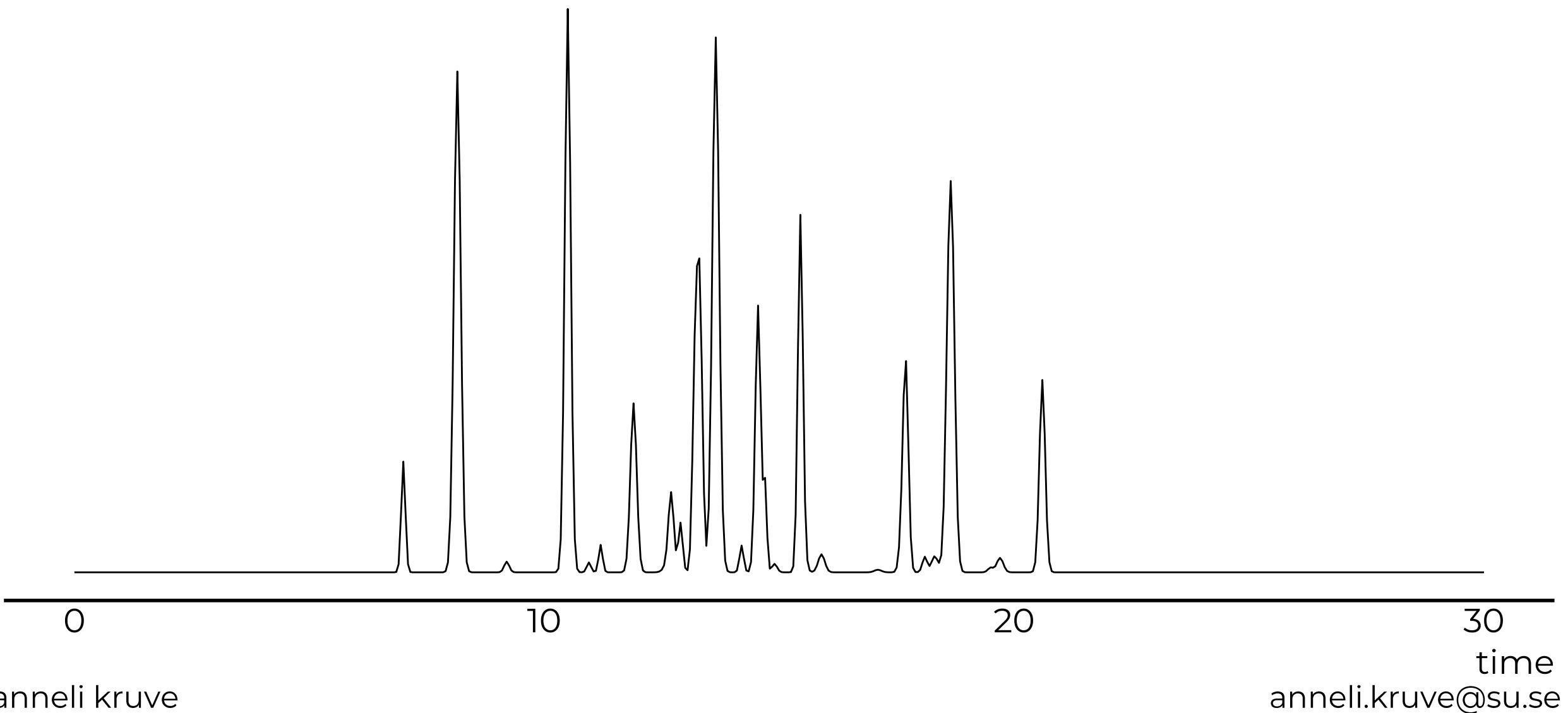
# semi-quantification in LC/HRMS non-targeted screening: the efforts of the community

anneli kruve

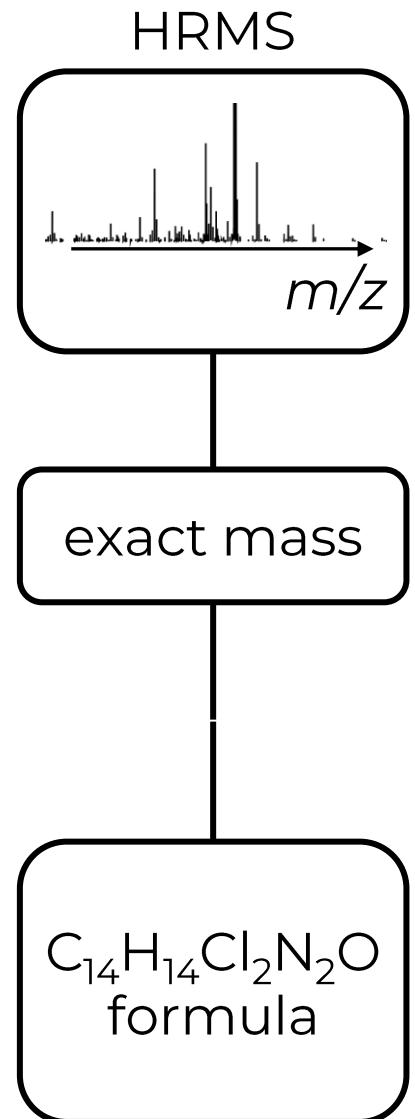
anneli.kruve@su.se

kruvelab.com

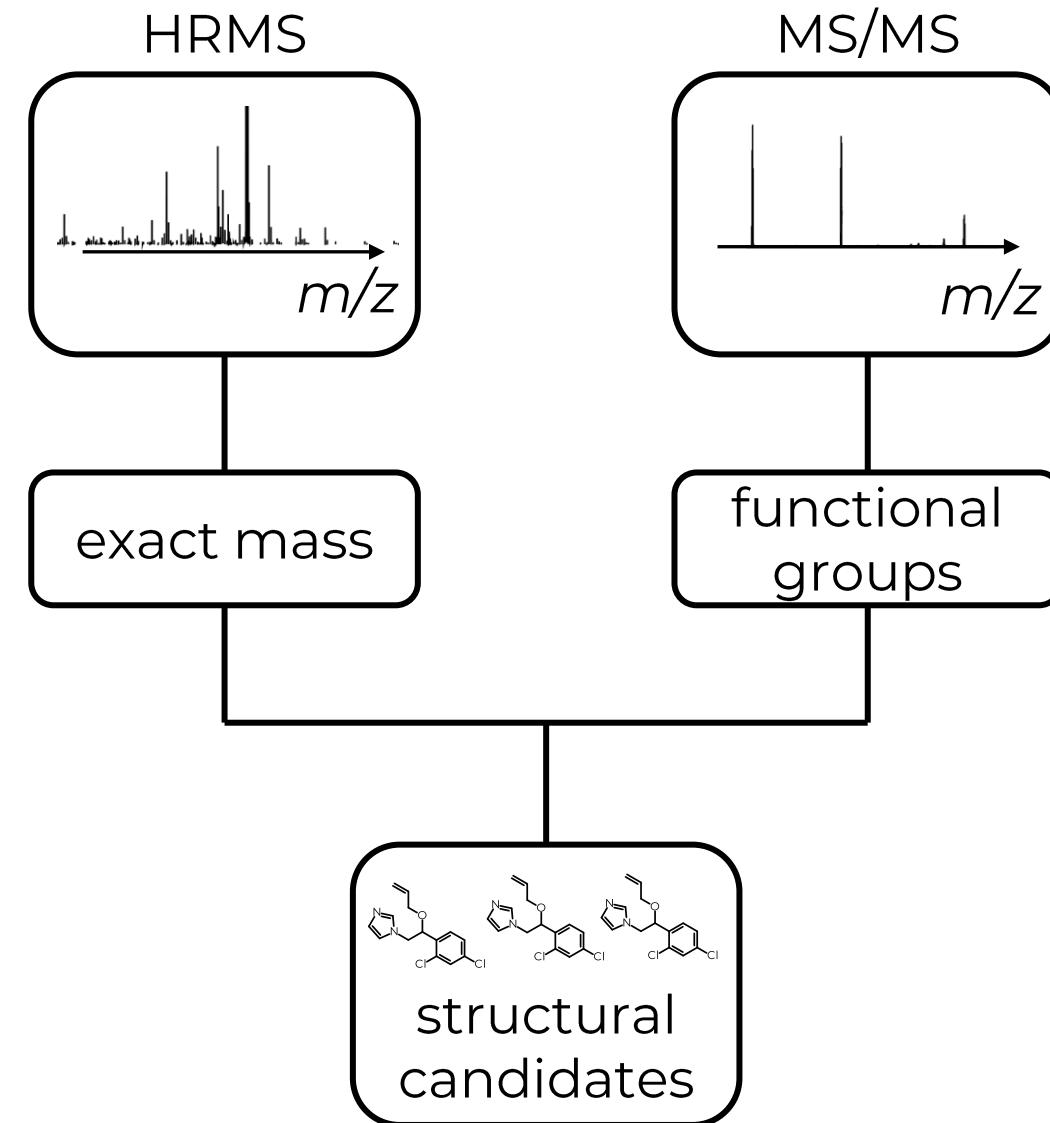
# nontarget screening



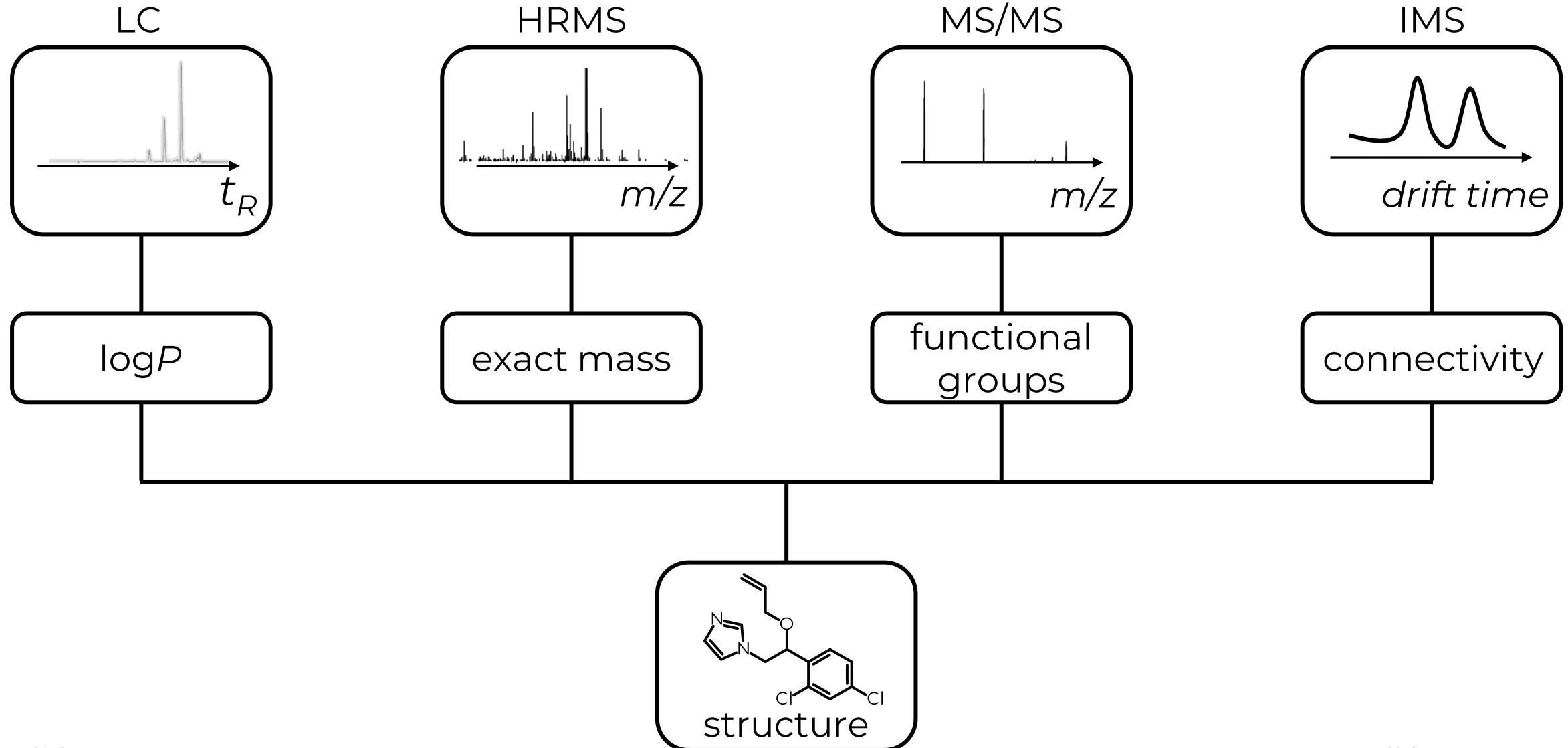
# identification



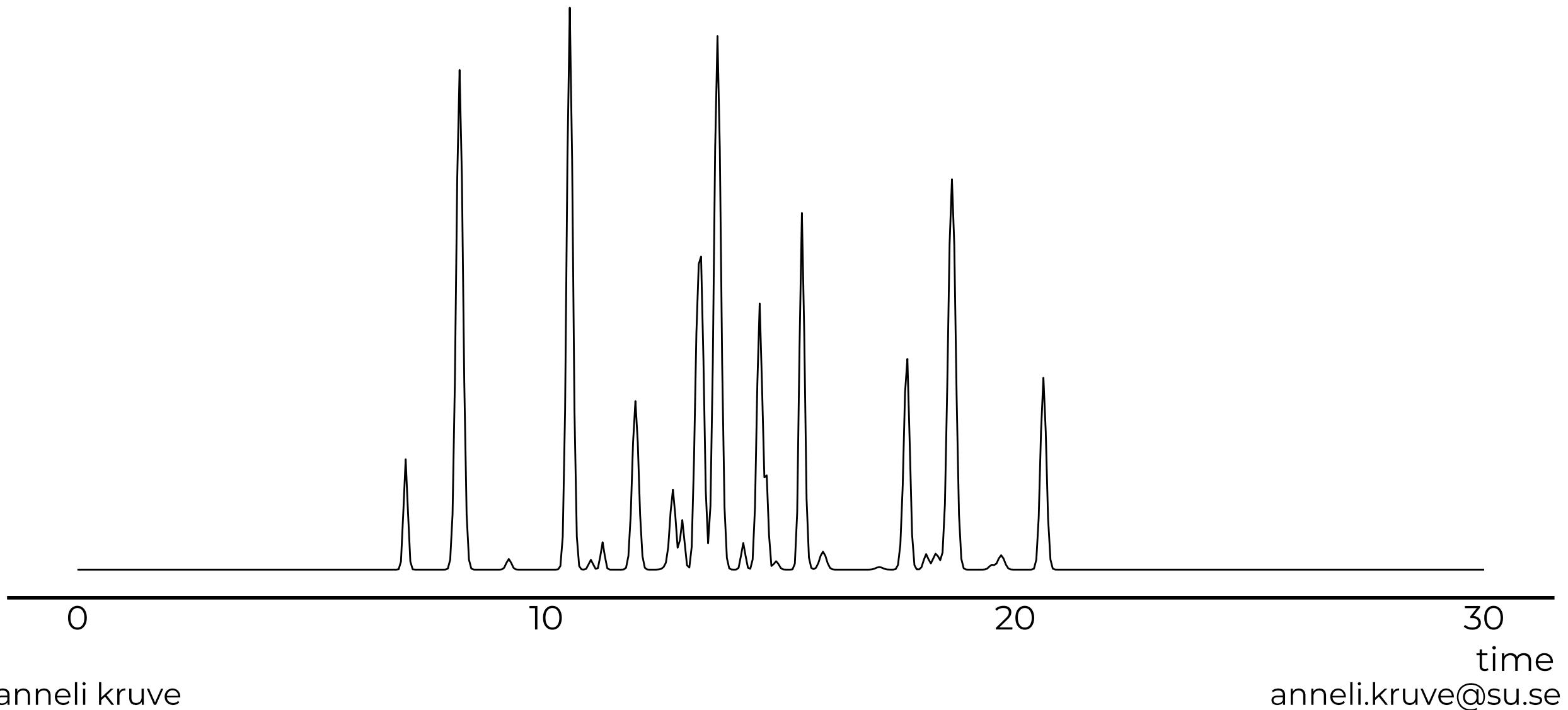
# identification



# identification



# what next?



# prioritization



toxicity

# prioritization



toxicity



concentration

# prioritization



toxicity

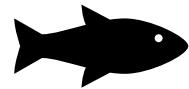


concentration



risk

# prioritization



toxicity



concentration



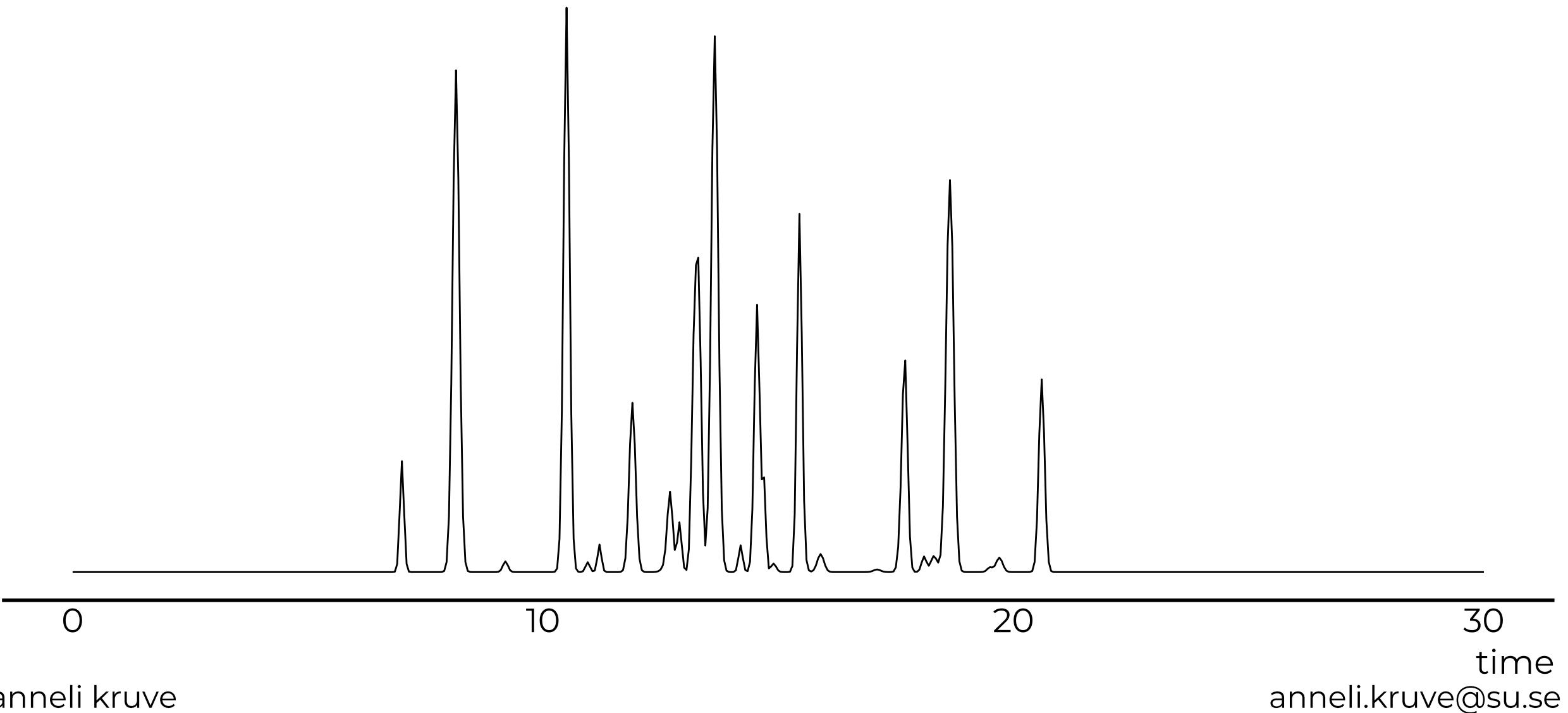
risk

$$\text{PriorityScore} = \frac{c_{\text{predicted}}}{AC_{50}^{\text{5th percentile}}}$$

# quantification in ESI/HRMS

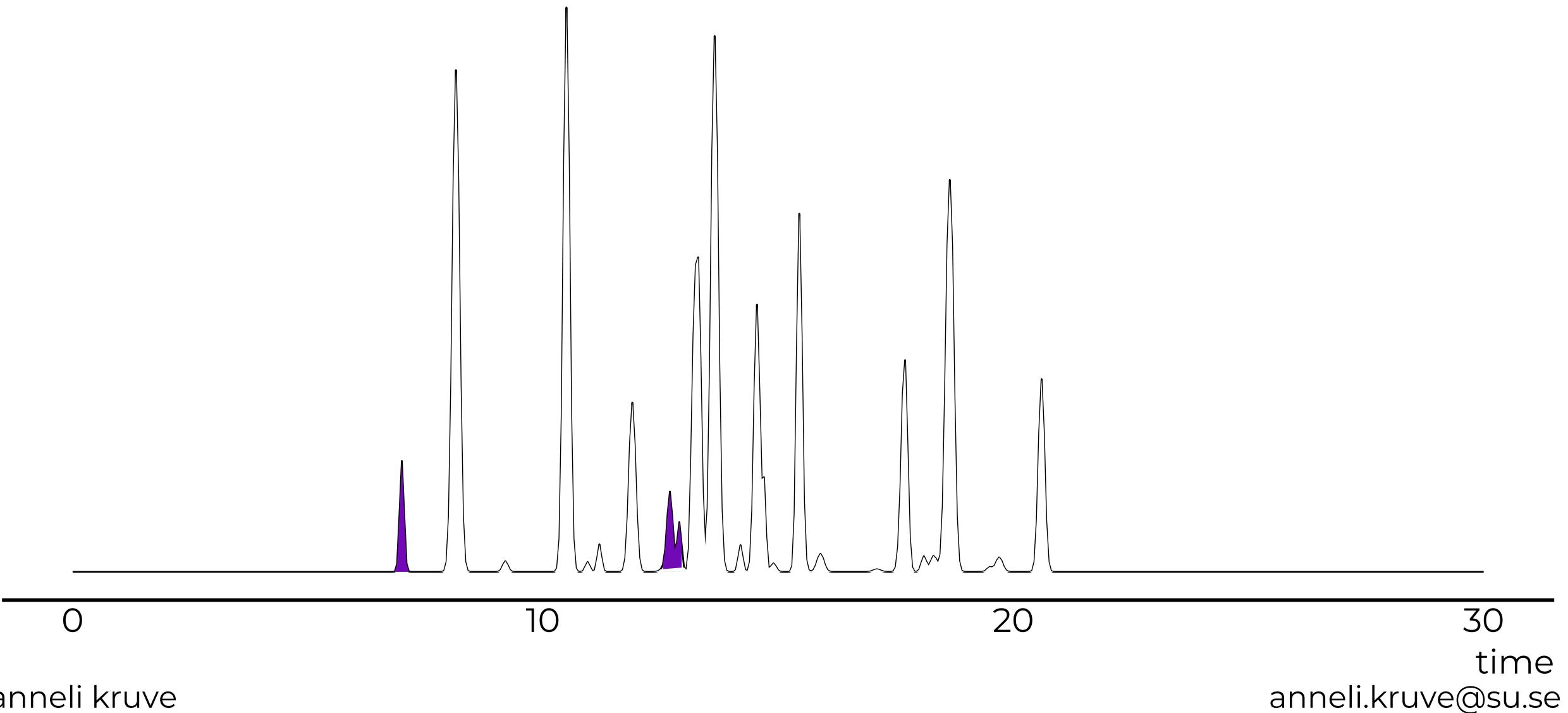
# quantification in ESI/HRMS

Malm et al. Molecules 2021



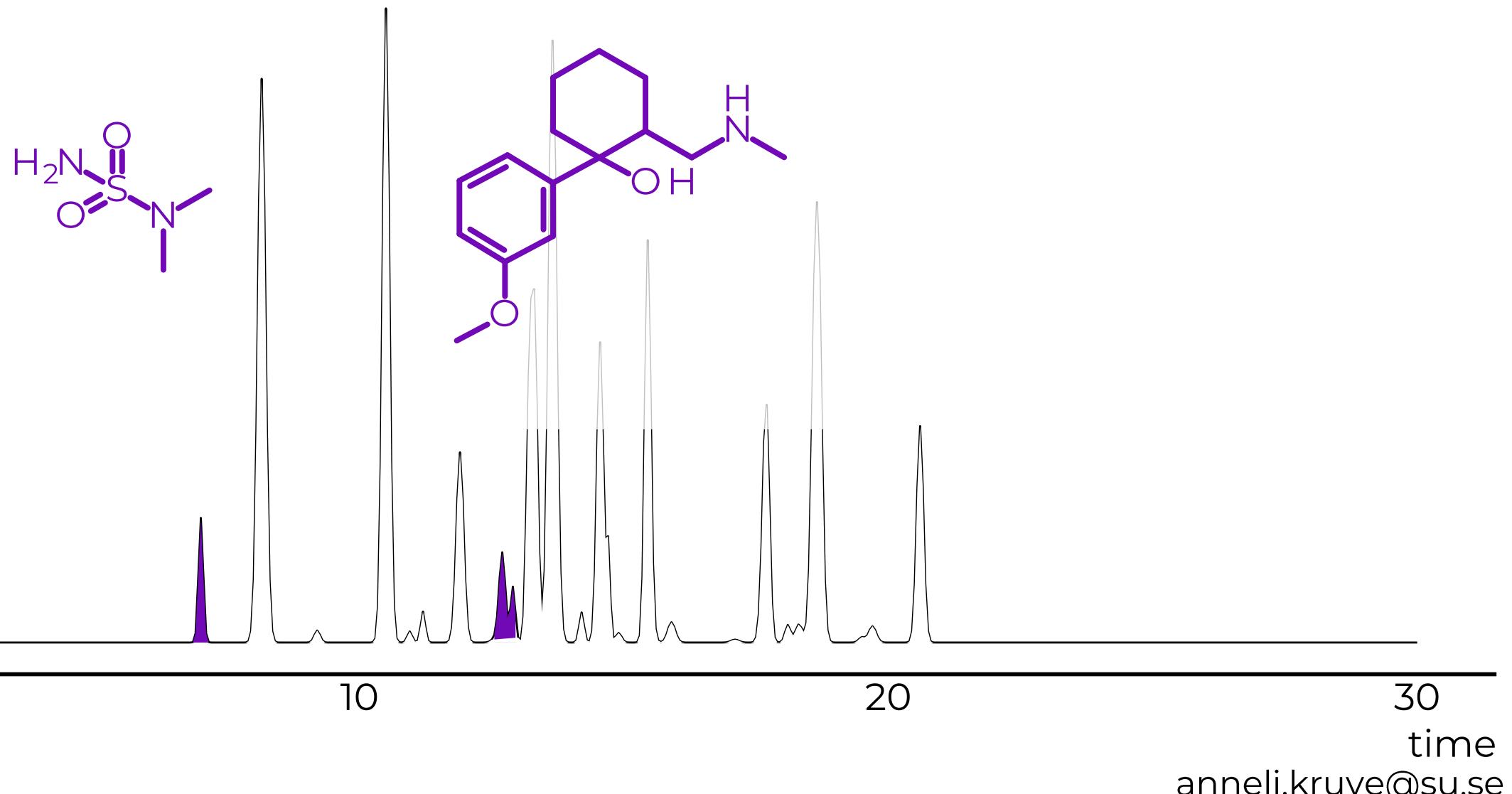
# quantification in ESI/HRMS

Malm et al. Molecules 2021



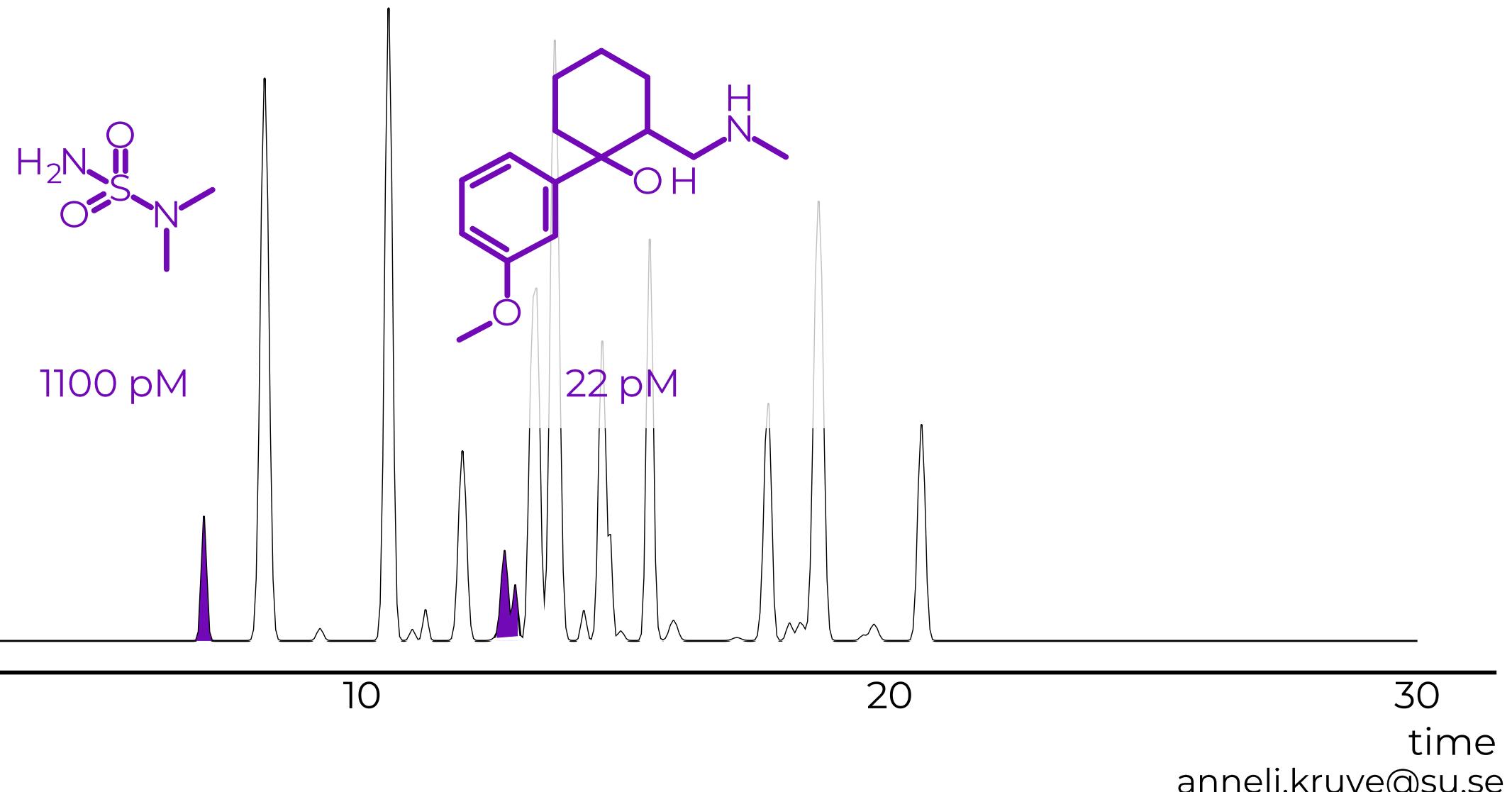
# quantification in ESI/HRMS

Malm et al. Molecules 2021

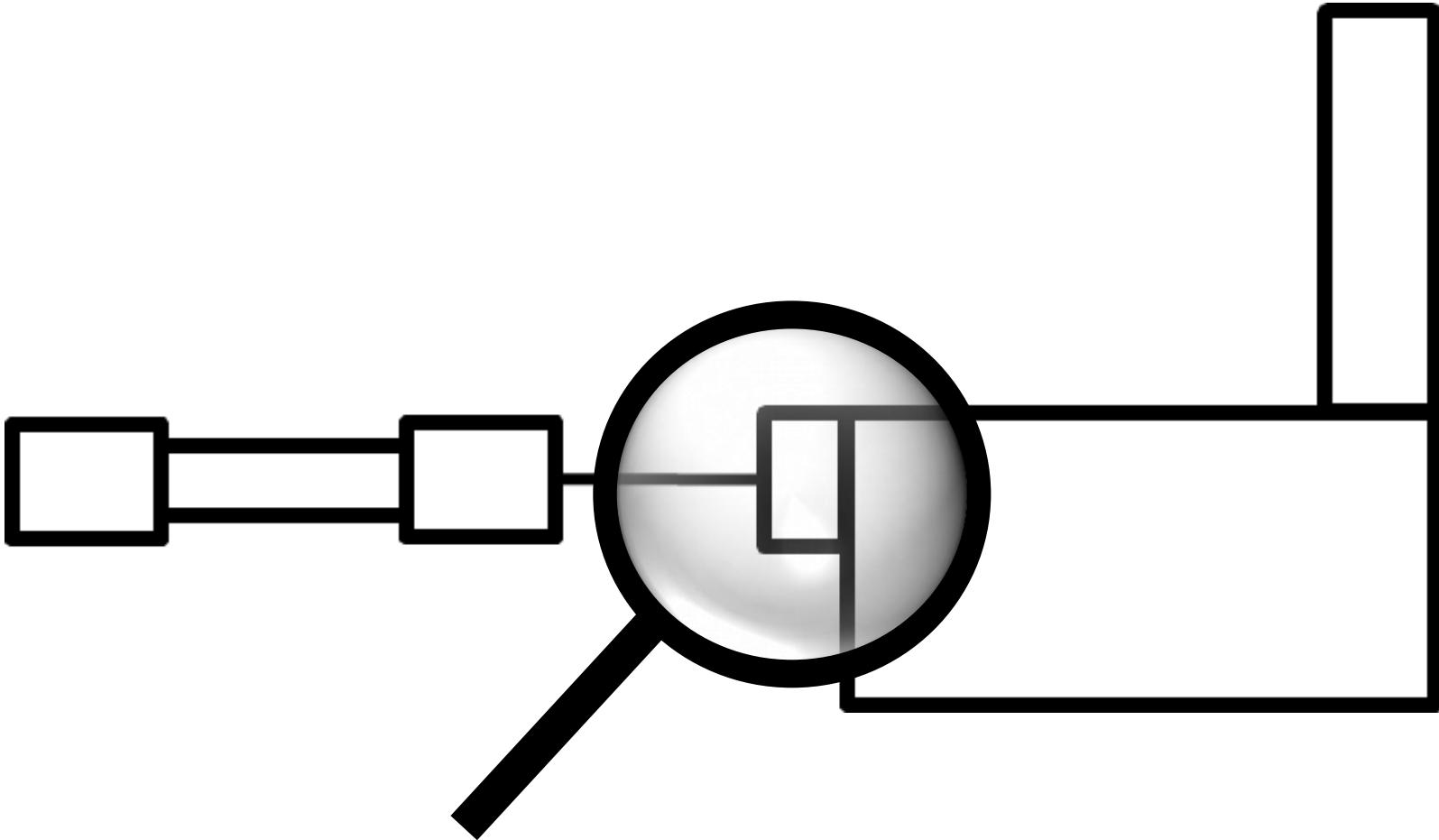


# quantification in ESI/HRMS

Malm et al. Molecules 2021

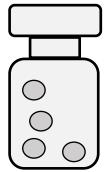


# electrospray



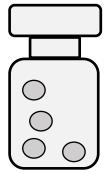
ionization efficiency

# workflow

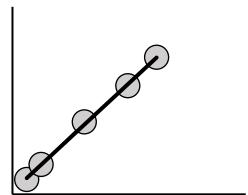


flow injections

# workflow

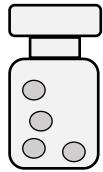


flow injections

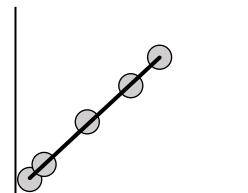


calibration graph

# workflow



flow injections



calibration graph

$$\frac{slope_1}{slope_2} \rightarrow IE$$

relative measurements

# structure

# structure

one solvent, purely analyte properties

377 chemicals

# structure

ionization efficiency

$1 \times 10^{+1}$

$1 \times 10^{+3}$

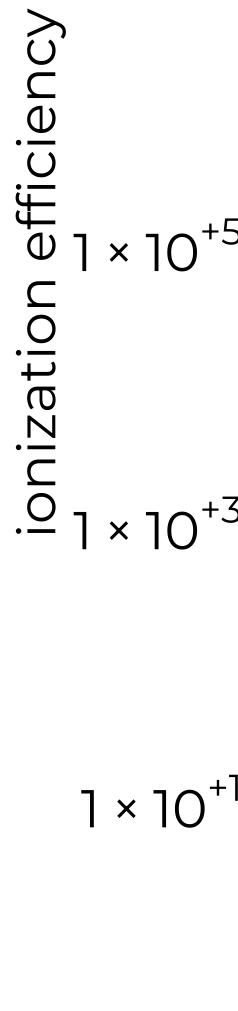
$1 \times 10^{+5}$



one solvent, purely analyte properties

377 chemicals

# structure



one solvent, purely analyte properties

377 chemicals

10,000,000x difference in ionization efficiency

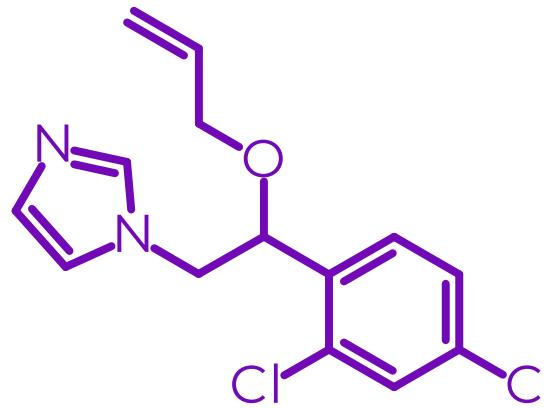
# structure

ionization efficiency

$1 \times 10^{+5}$

$1 \times 10^{+3}$

$1 \times 10^{+1}$



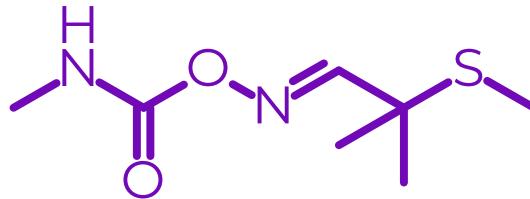
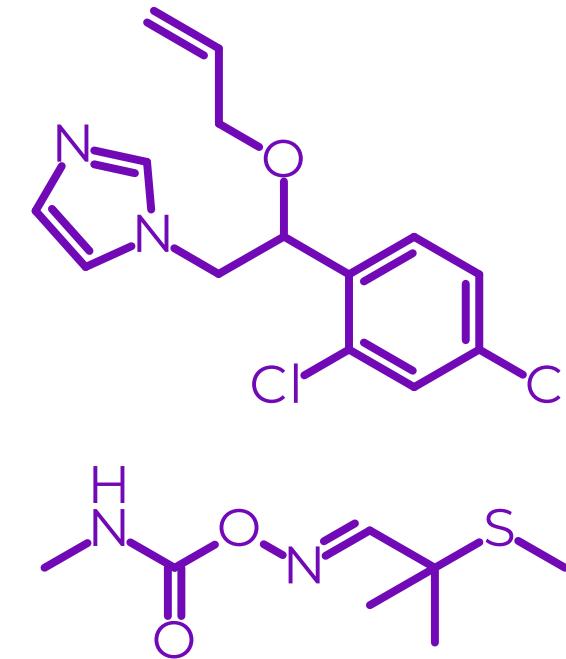
# structure

ionization efficiency

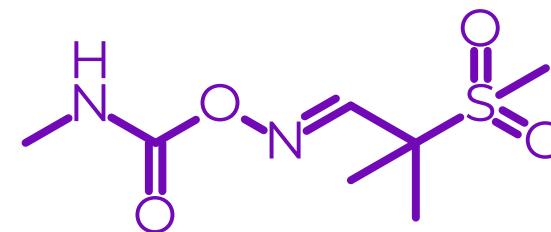
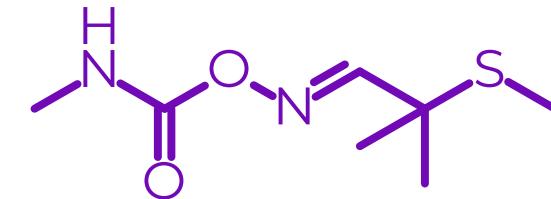
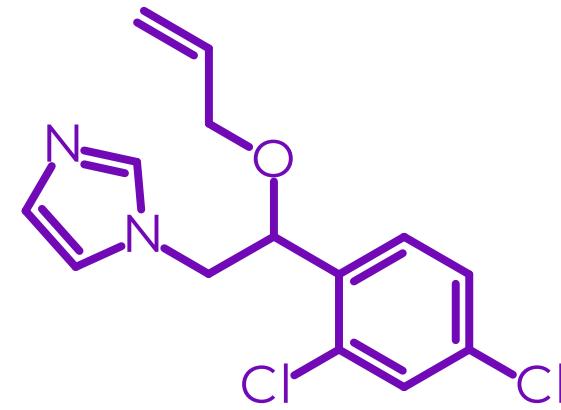
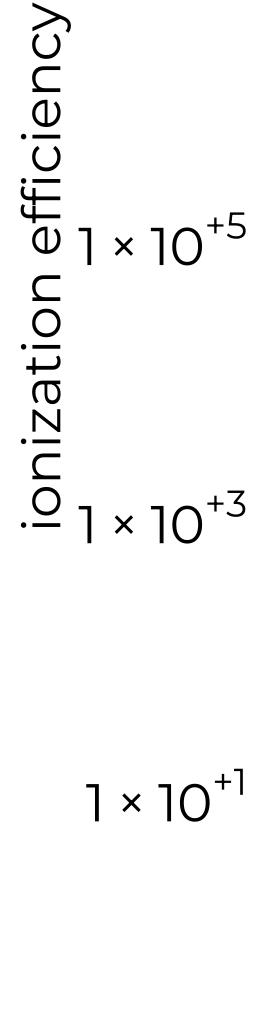
$1 \times 10^{+1}$

$1 \times 10^{+3}$

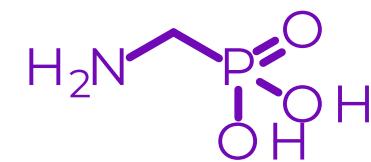
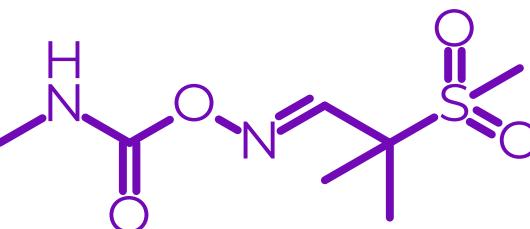
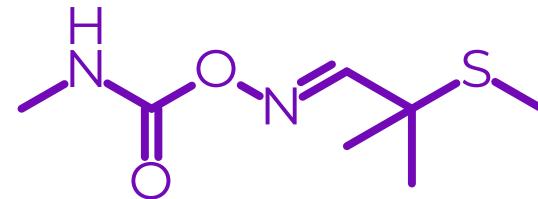
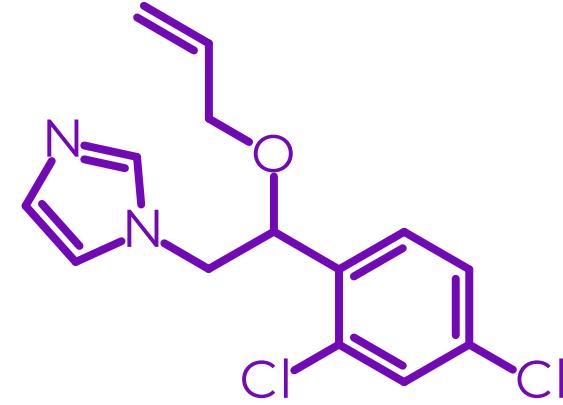
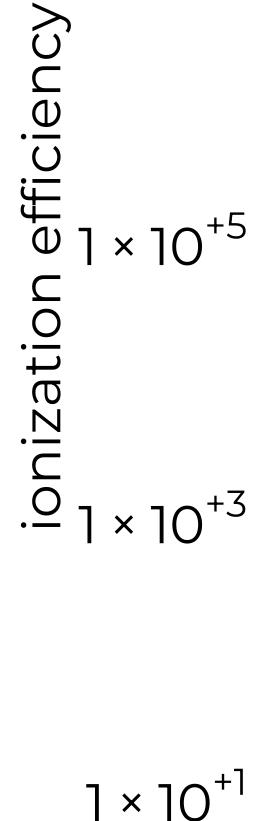
$1 \times 10^{+5}$



# structure



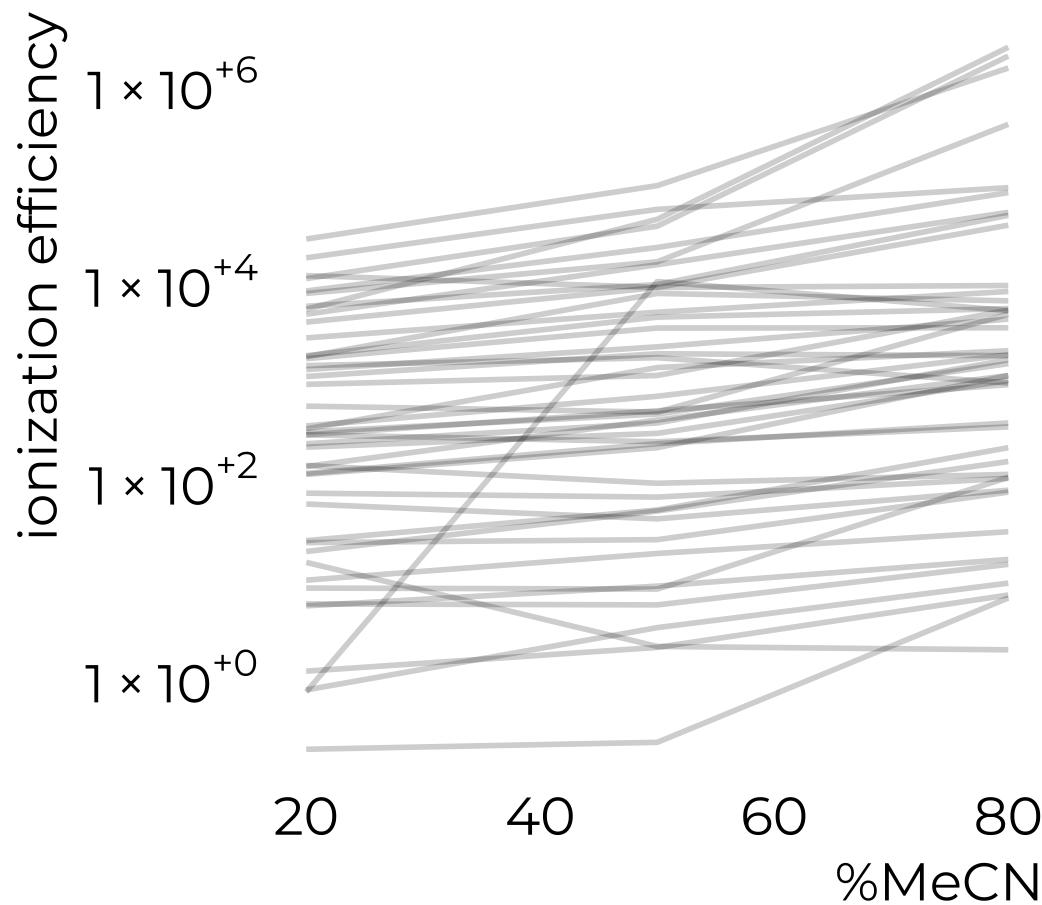
# structure



# mobile phase: organic modifier

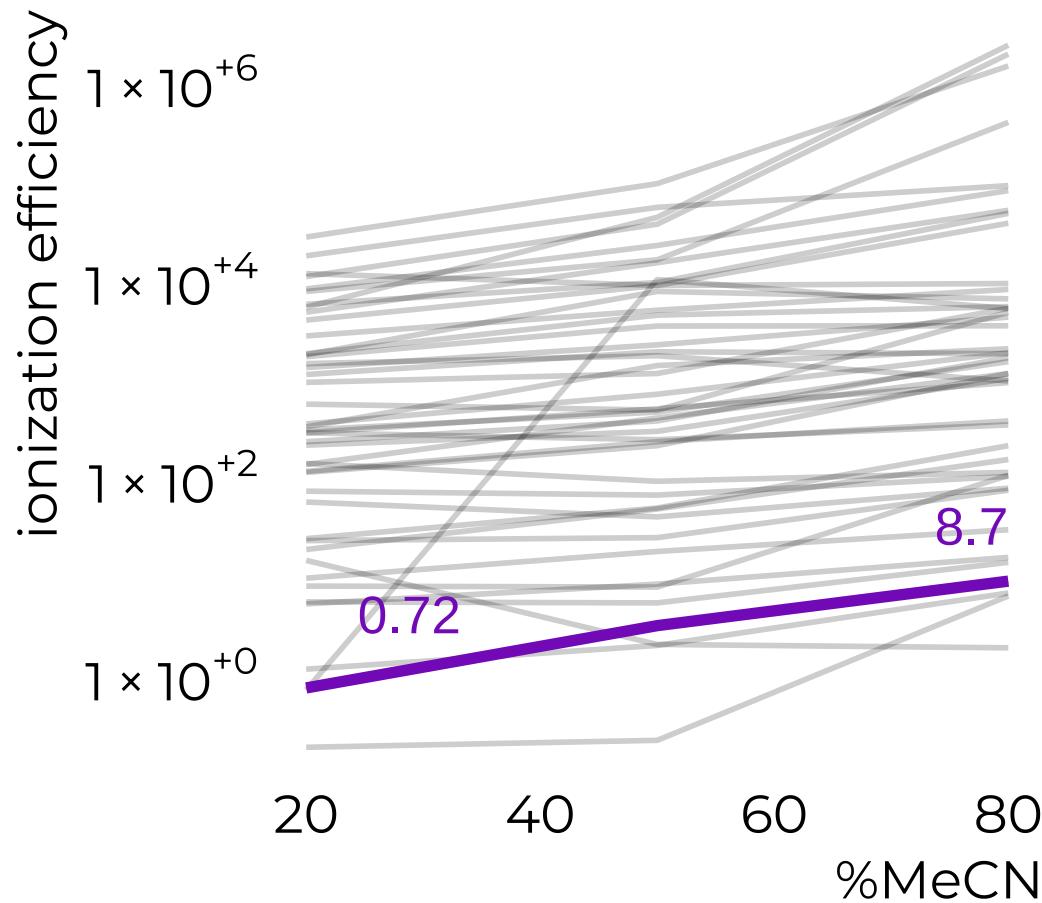
# mobile phase: organic modifier

Liigand et al. JASMS 2014



# mobile phase: organic modifier

Liigand et al. JASMS 2014



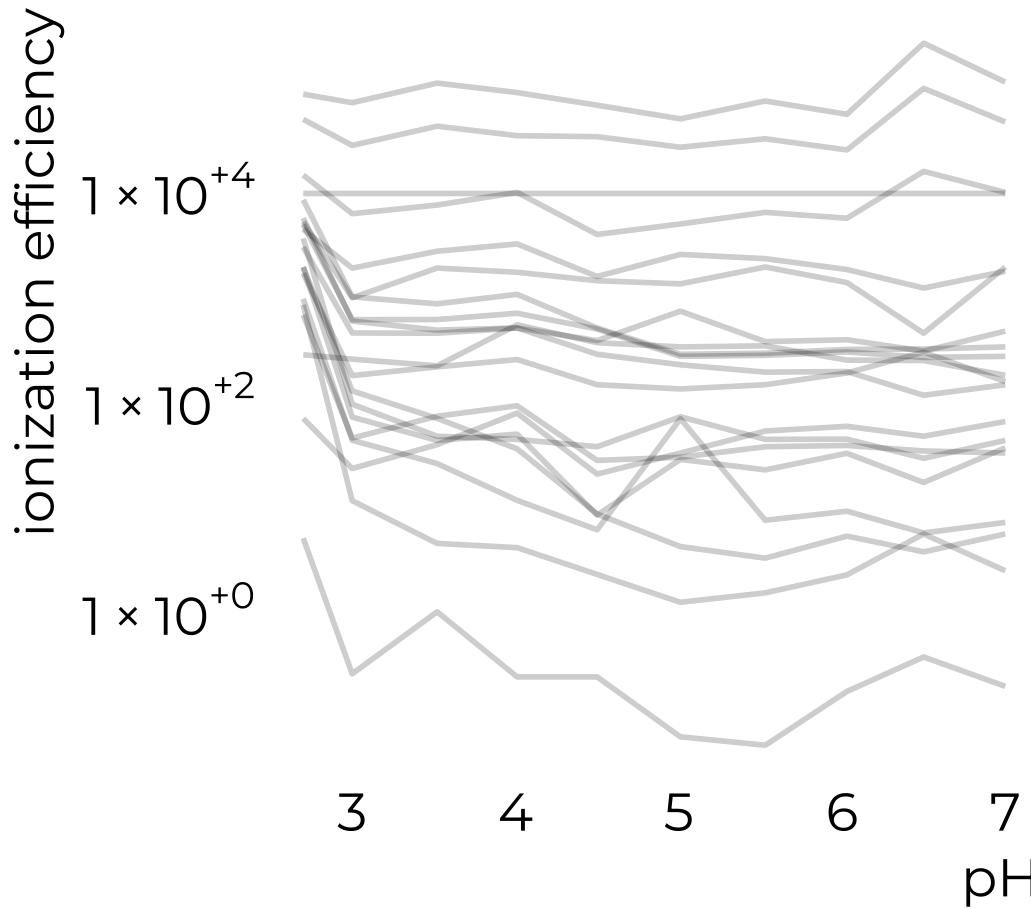
$\% \text{MeCN} \uparrow \sim \text{ionization efficiency} \uparrow$

# mobile phase: pH

# mobile phase: pH

Liigand et al. JASMS 2017

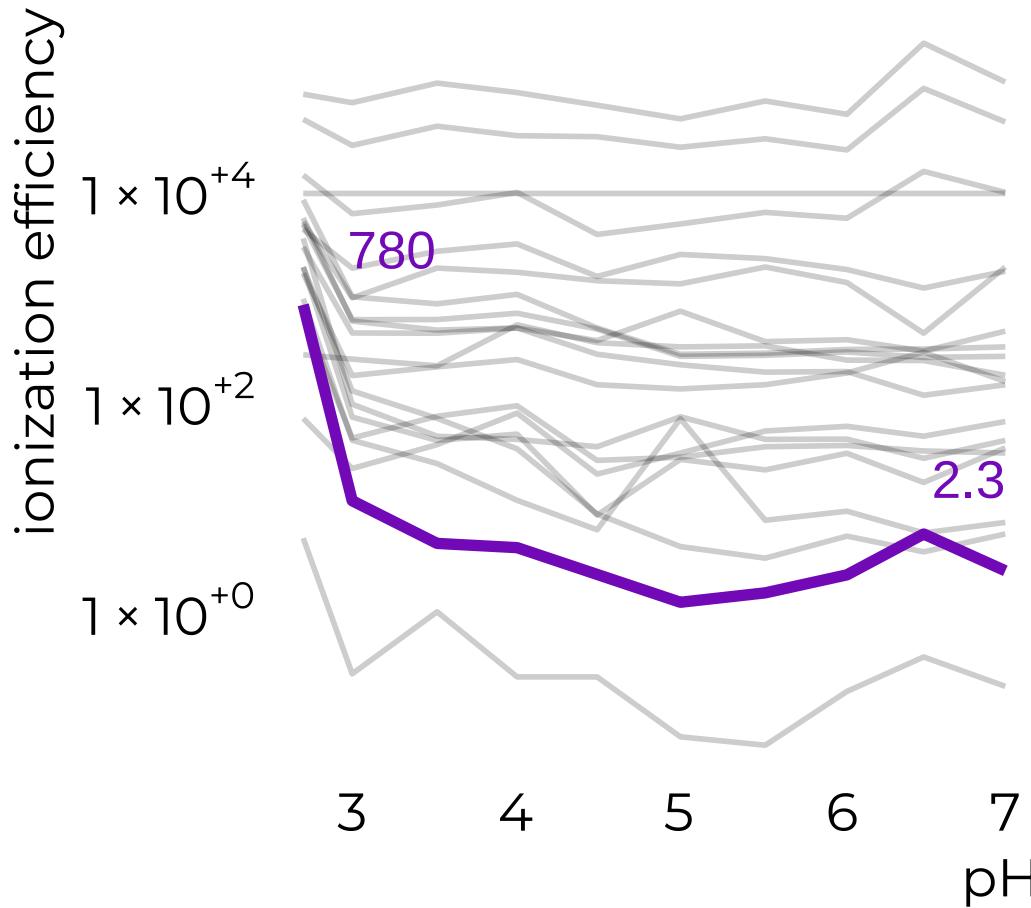
Kruve et al. Anal Chem 2017



# mobile phase: pH

Liigand et al. JASMS 2017

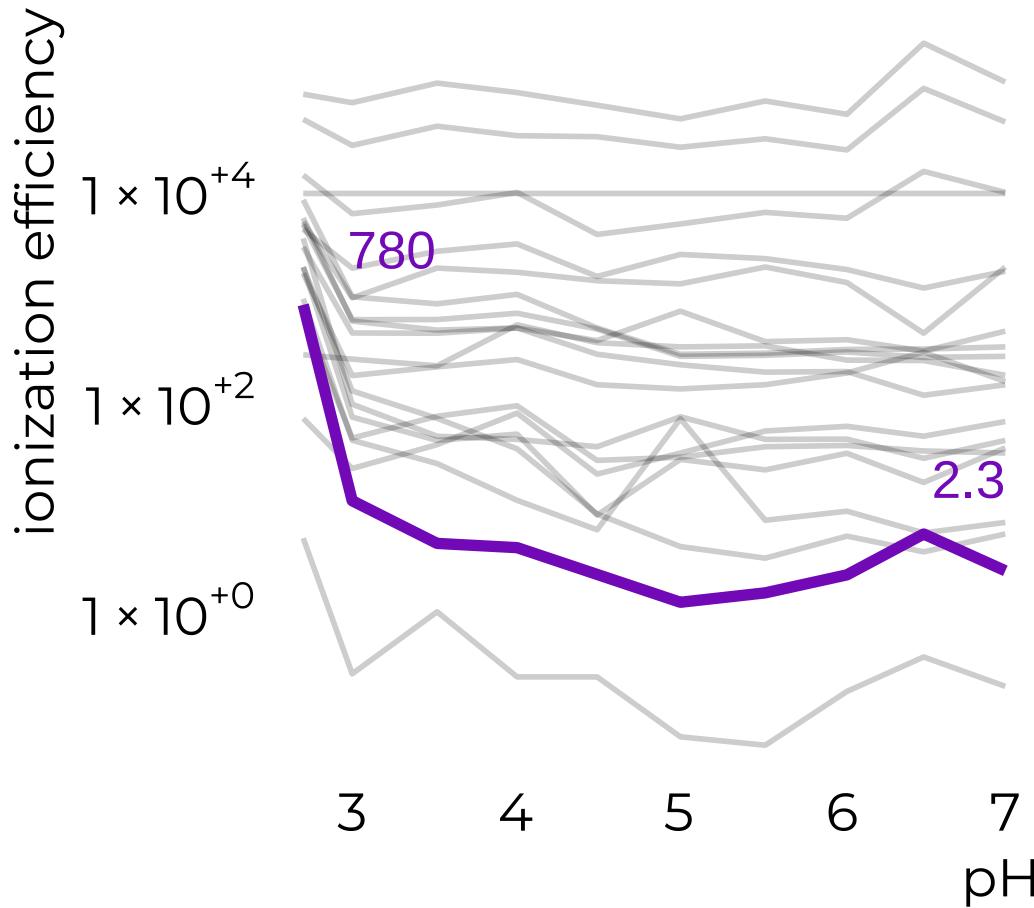
Kruve et al. Anal Chem 2017



# mobile phase: pH

Liigand et al. JASMS 2017

Kruve et al. Anal Chem 2017

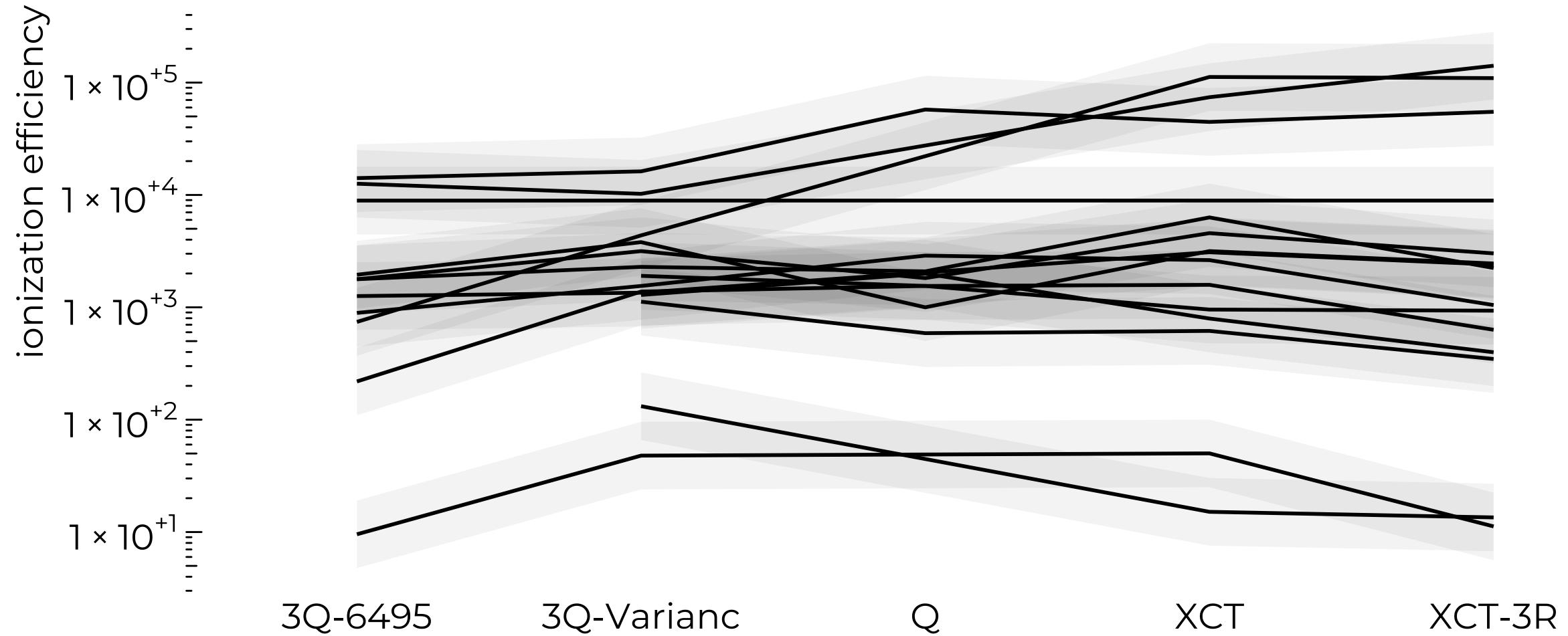


pH & buffer type affect ionization efficiency

# instruments

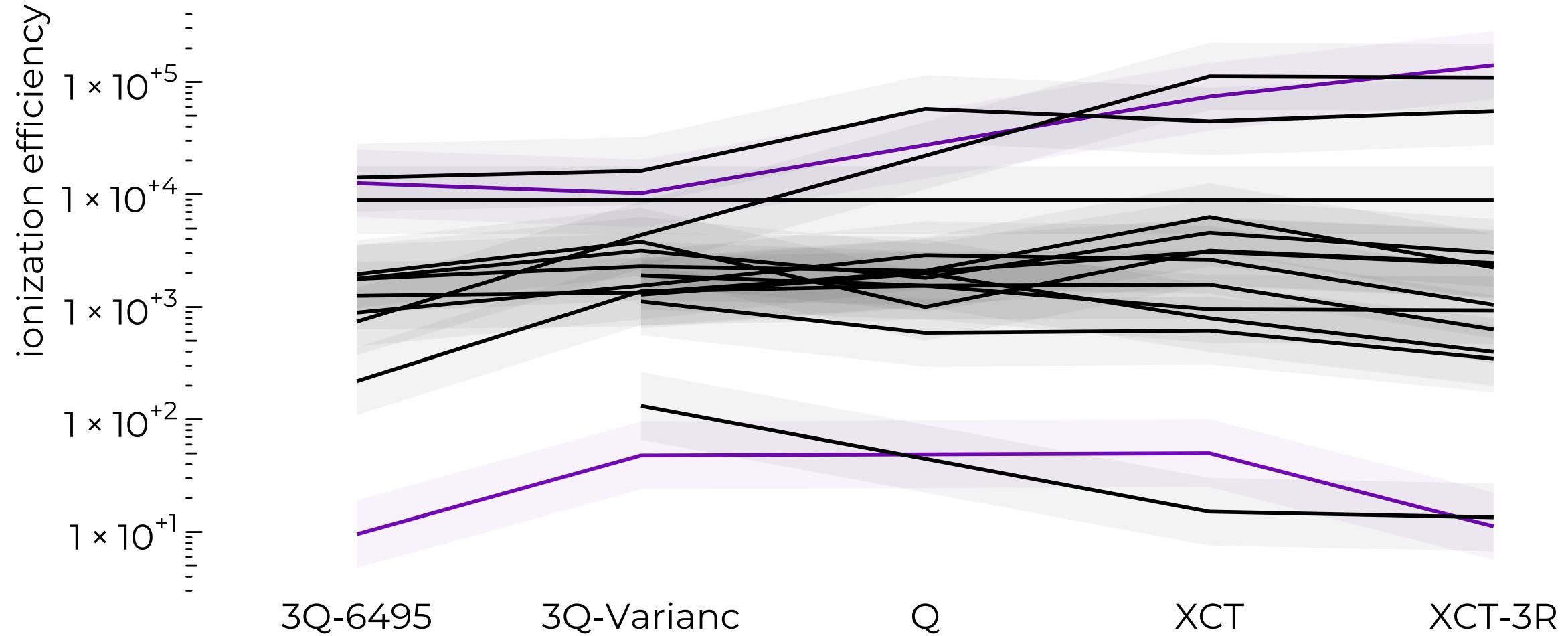
# instruments

Liigand et al. JASMS 2015



# instruments

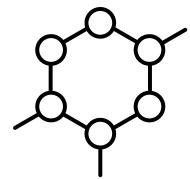
Liigand et al. JASMS 2015



# quantification

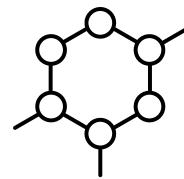
approaches

# three common methods

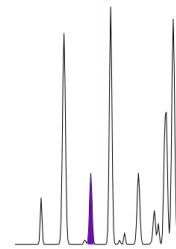


structurally similar chemicals

# three common methods

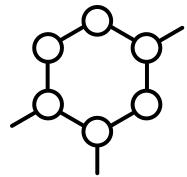


structurally similar chemicals

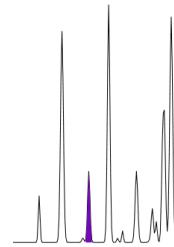


close eluting chemicals

# three common methods



structurally similar chemicals



close eluting chemicals



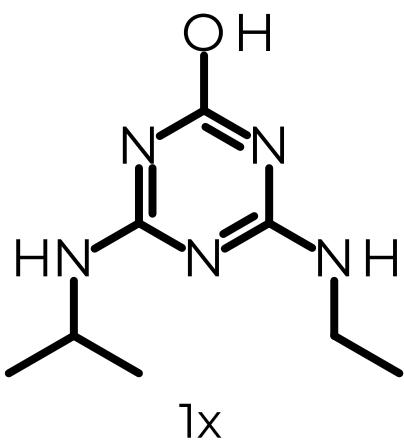
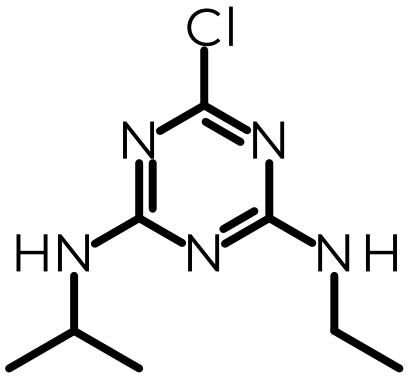
machine learning

# quantification

with structurally similar chemical

# parent – transformation product

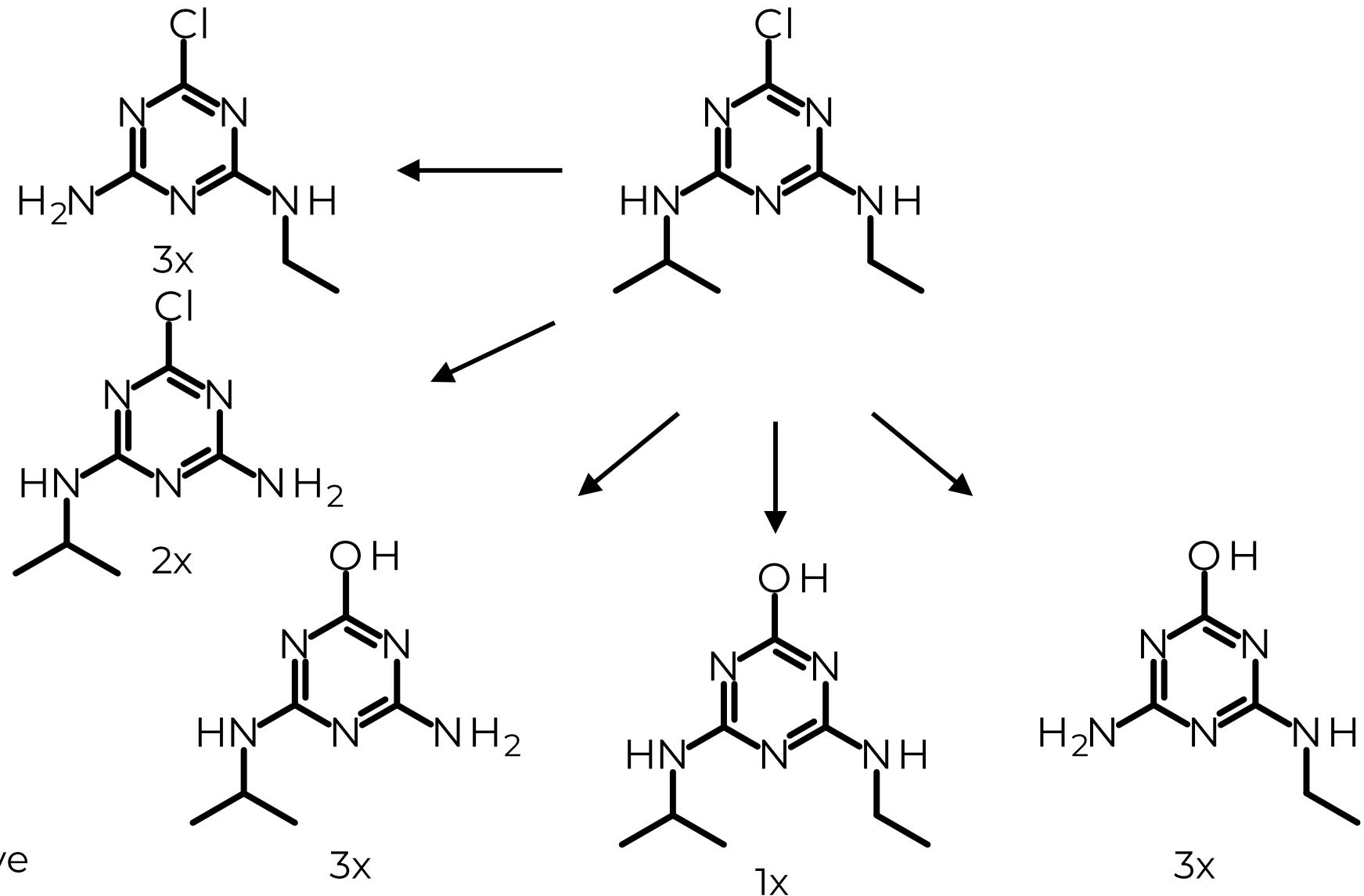
Malm et al. Molecules 2021



1x

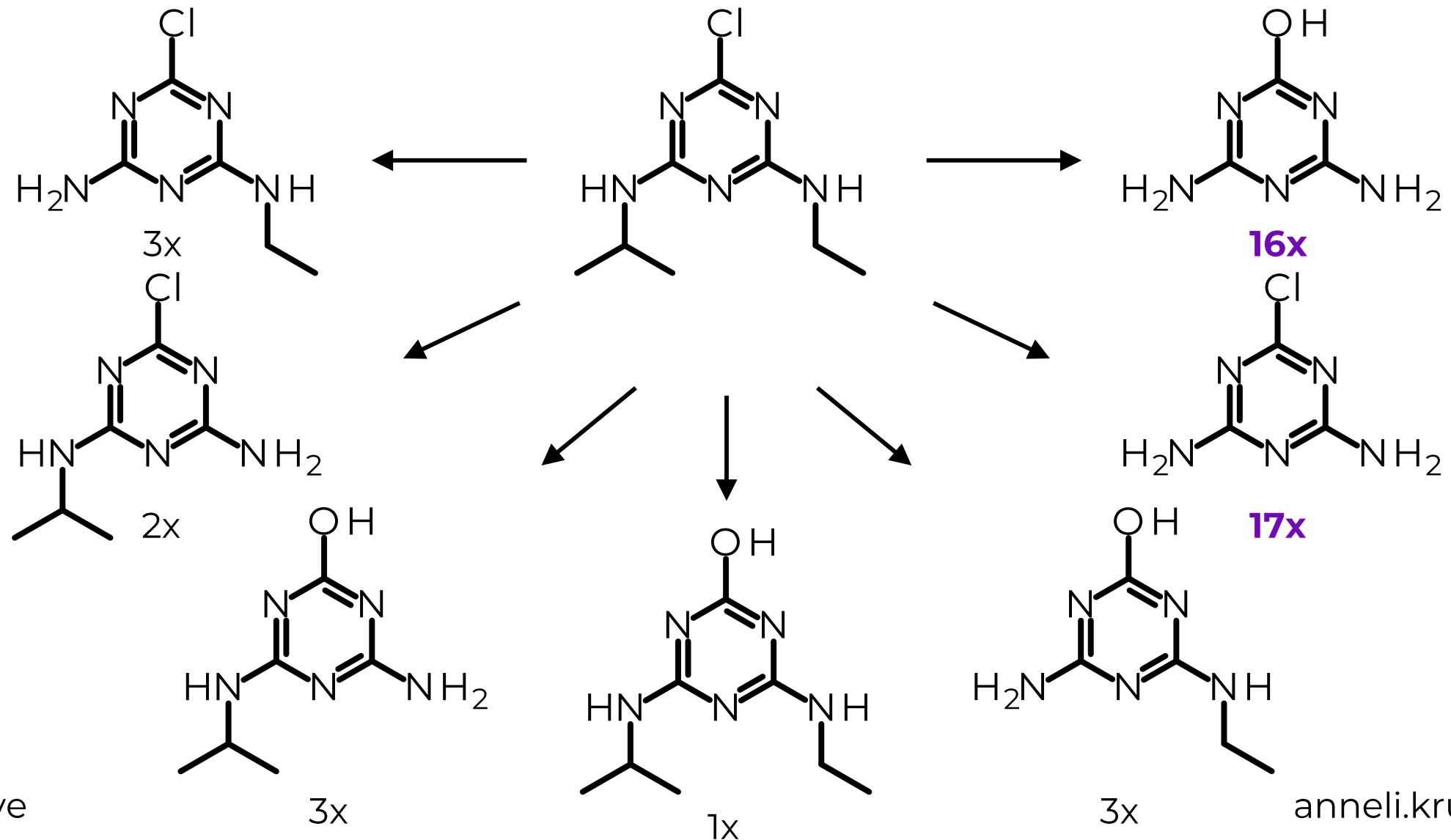
# parent – transformation product

Malm et al. Molecules 2021



# parent – transformation product

Malm et al. Molecules 2021



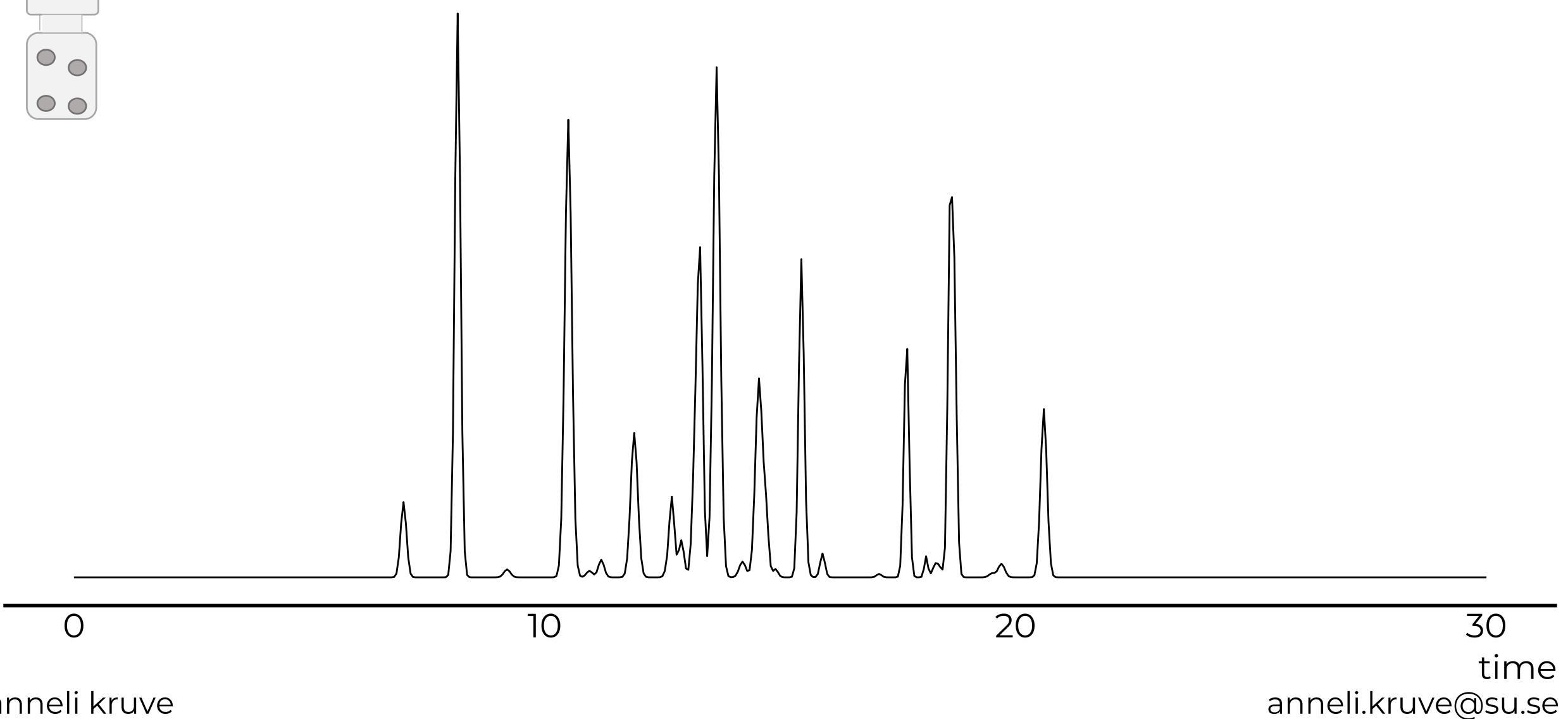
# quantification

with close eluting chemical

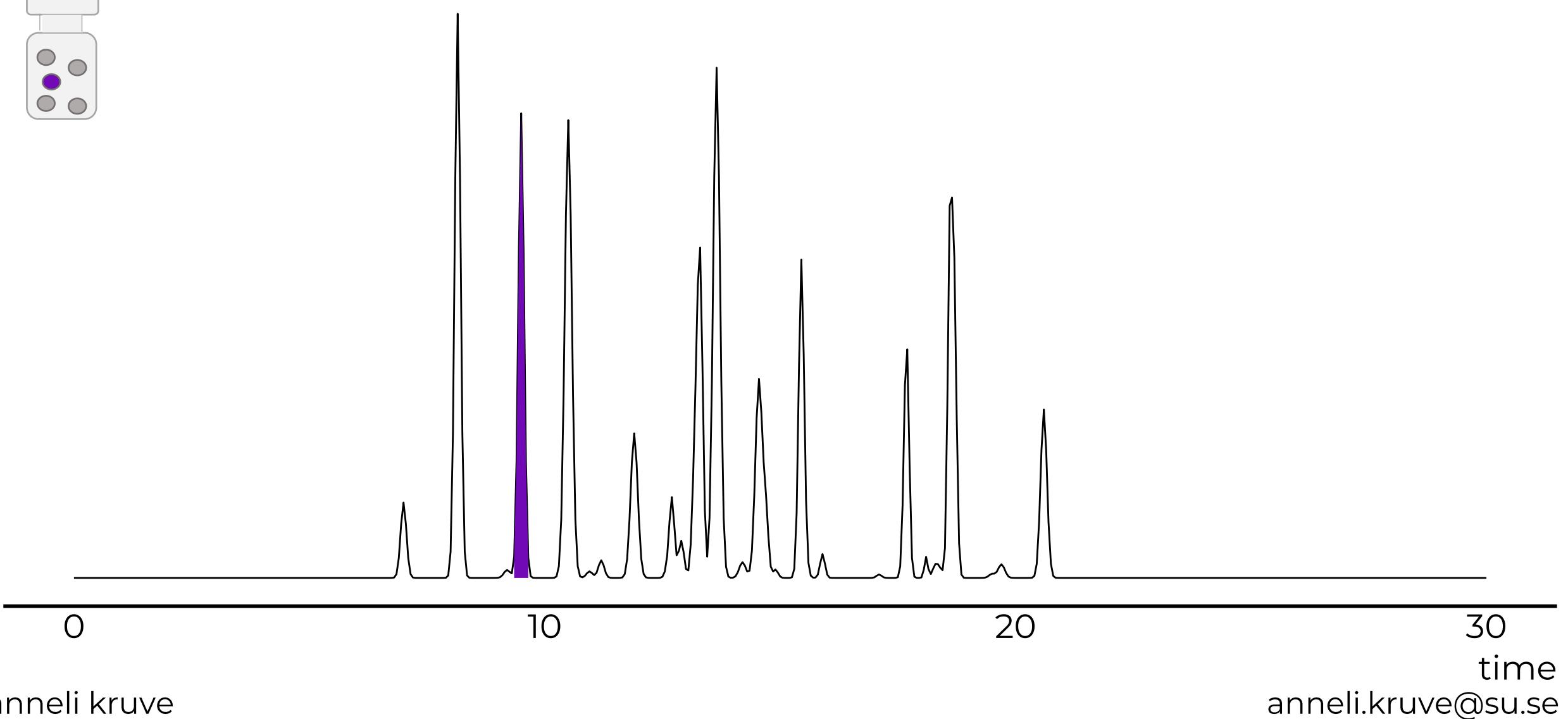
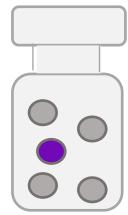
# compound eluting closest



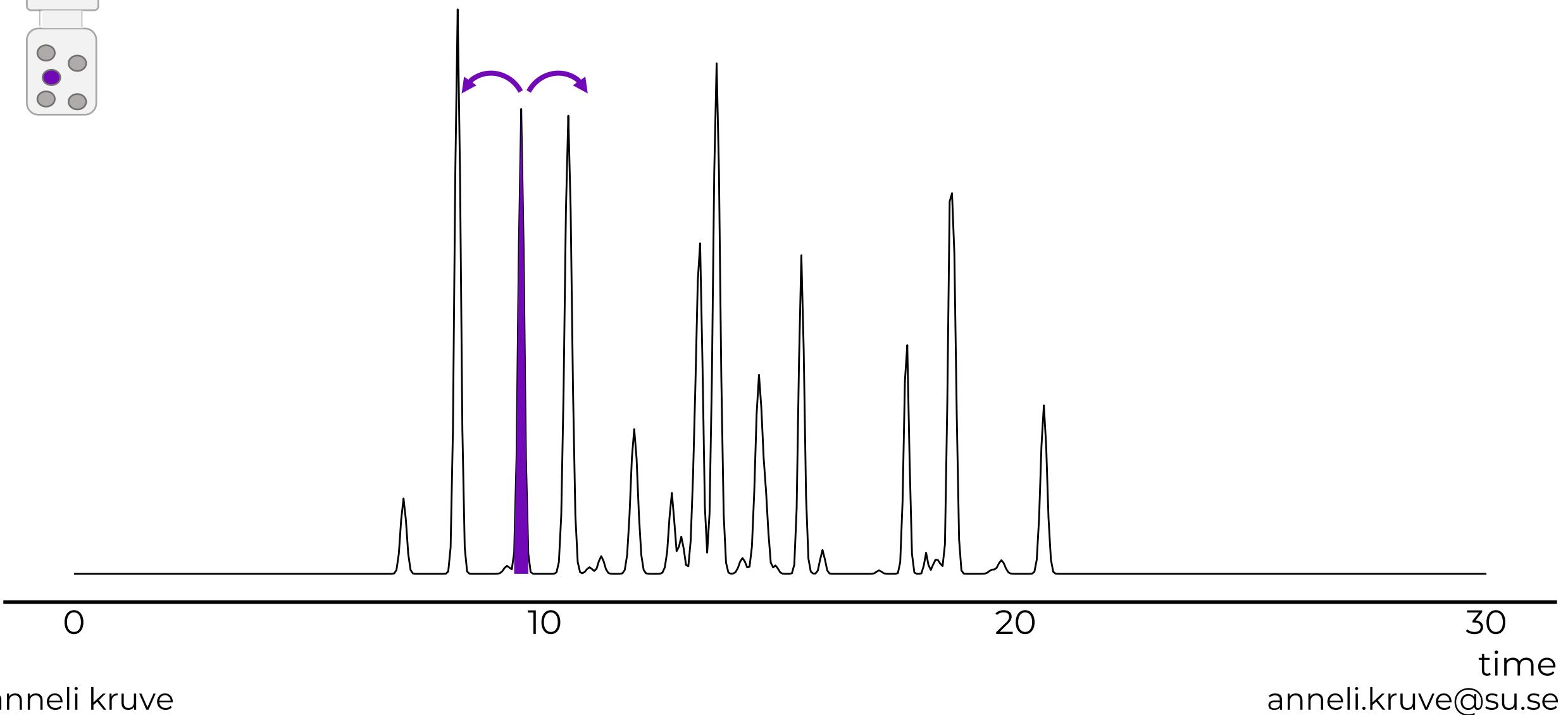
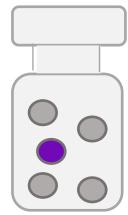
# compound eluting closest



# compound eluting closest



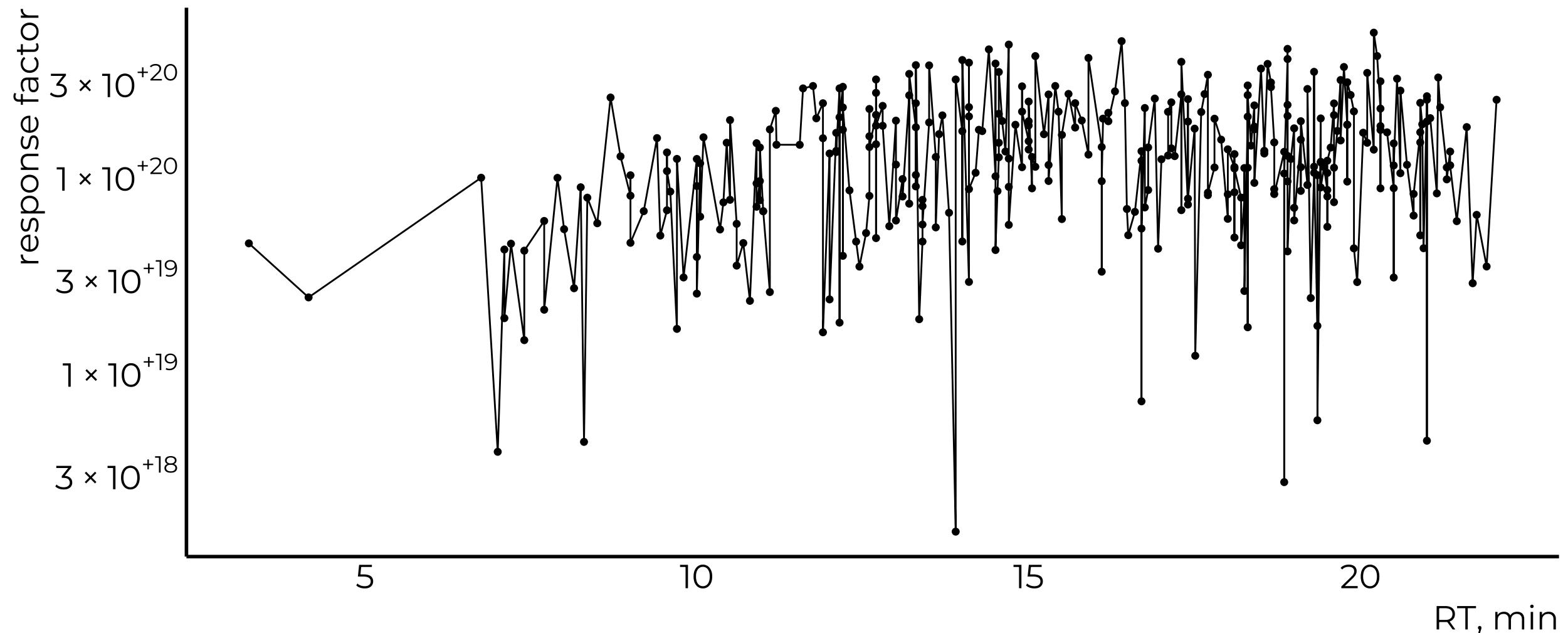
# compound eluting closest



# pesticides and micropollutants

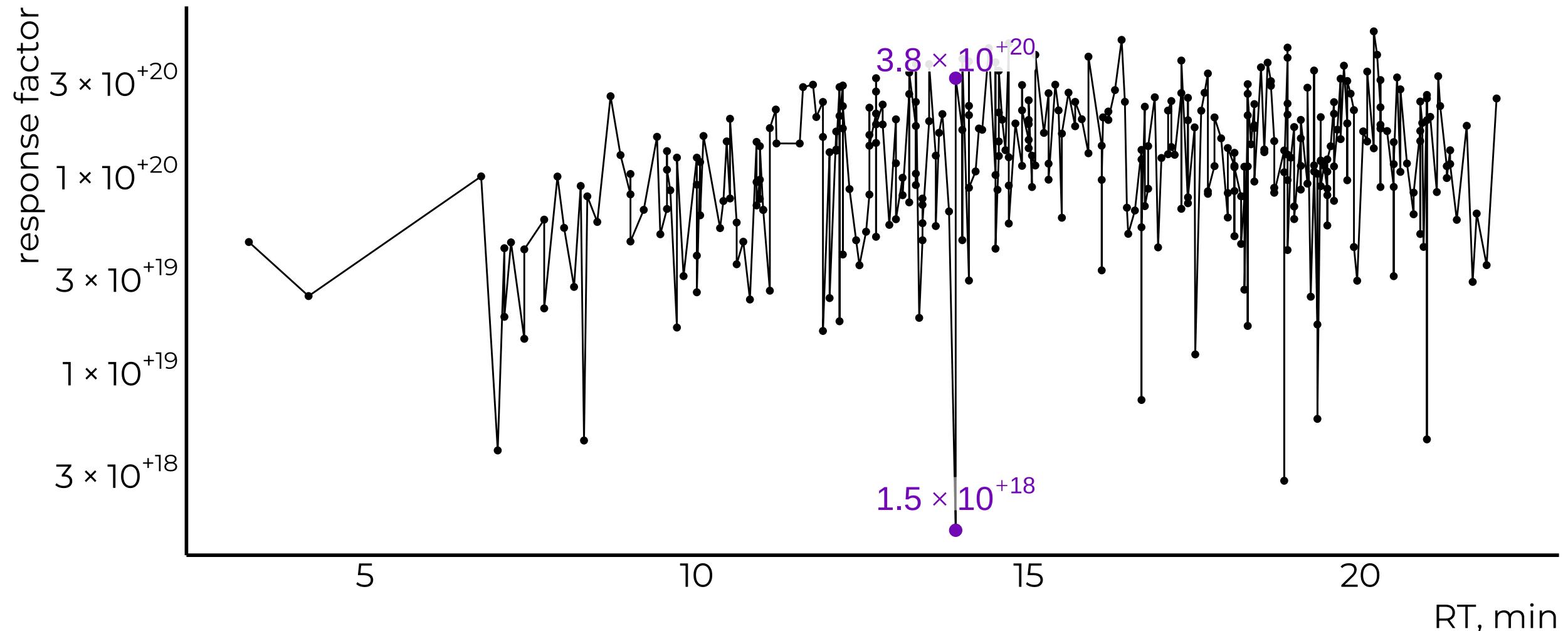
# pesticides and micropollutants

Kruve et al. Anal Bioanal Chem 2021



# pesticides and micropollutants

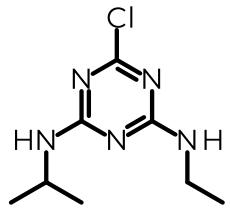
Kruve et al. Anal Bioanal Chem 2021



# quantification

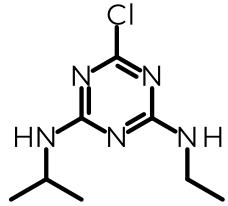
with machine learning

# workflow



SMILES & solvent

# workflow

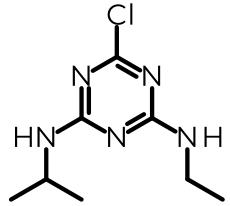


SMILES & solvent



molecular descriptors

# workflow



SMILES & solvent

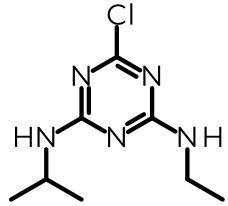


molecular descriptors



training machine learning models

# workflow



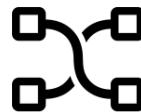
SMILES & solvent



molecular descriptors

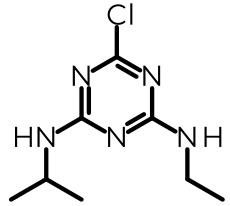


training machine learning models



best model selection

# workflow



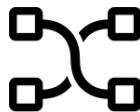
SMILES & solvent



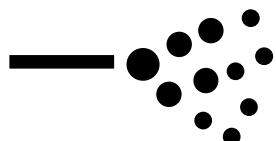
molecular descriptors



training machine learning models



best model selection



predict ionization efficiency

# performance

Liigand et al. Sci Reports 2020  
Sepman et al. under review

1403 chemicals  
 $x\log P$  from -6.6 to 22.5

13 labs/methods  
flow injections  
RP & HILIC methods

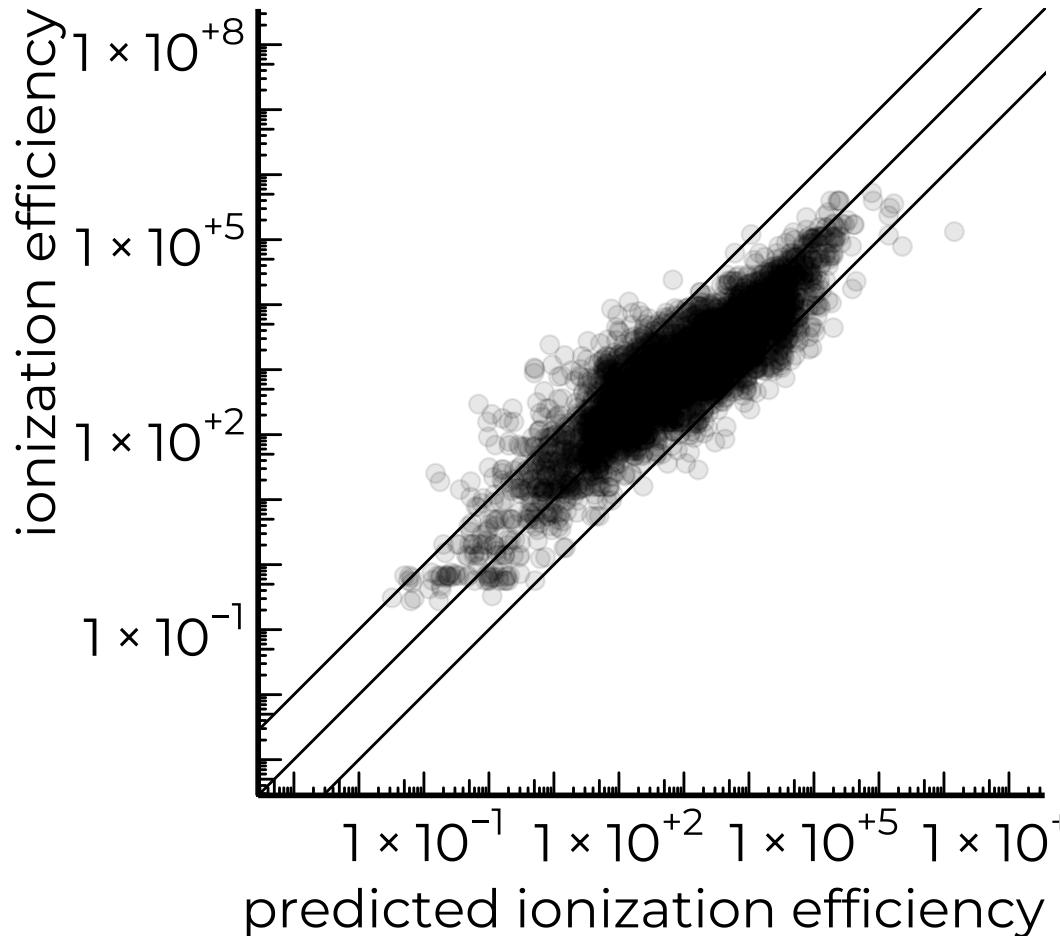
MeCN & MeOH  
0 – 100%

pH 2.1 – 10.8  
different buffers

# performance

Liigand et al. Sci Reports 2020

Sepman et al. under review



IE range

100,000,000

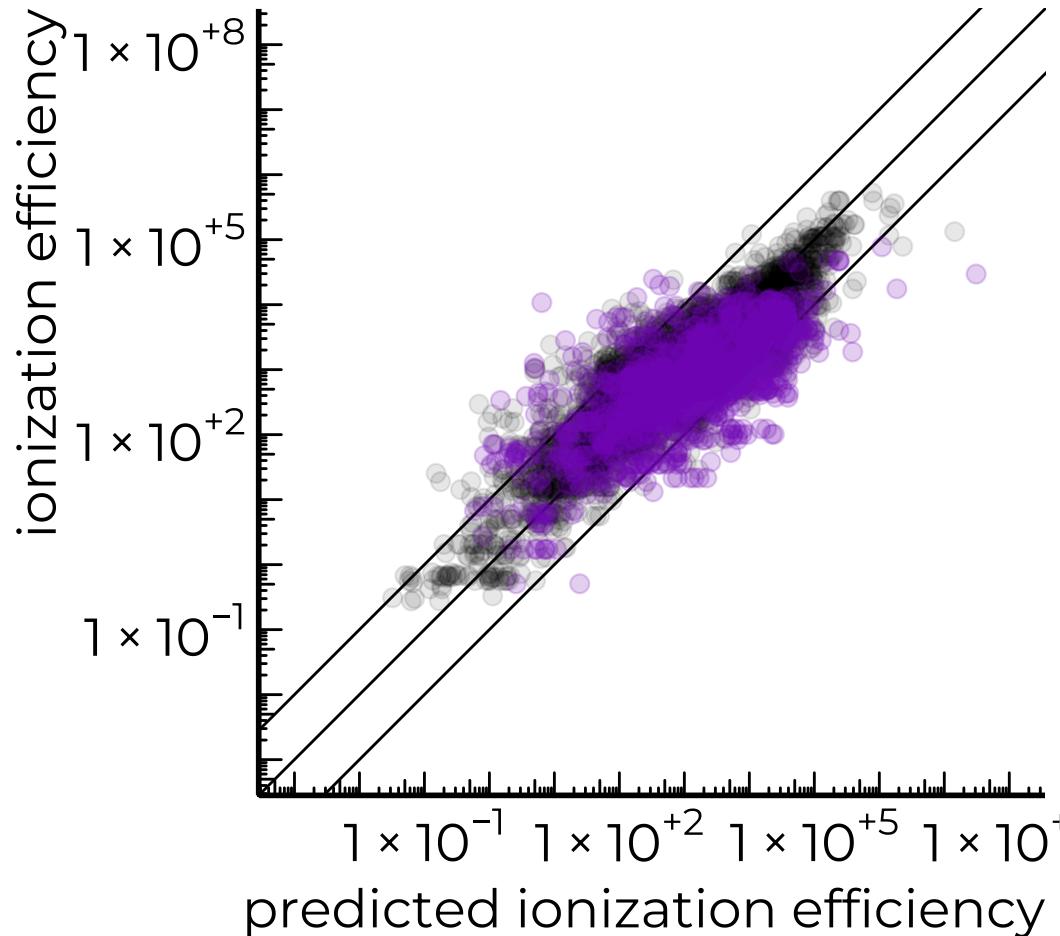
training set

RMSE 3.6x

# performance

Liigand et al. Sci Reports 2020

Sepman et al. under review



IE range

100,000,000

training set

RMSE 3.6x

test set

RMSE 5.6x

# application

<b>compound</b>	<b>peak area</b>
methiocarb sulfoxide	5,300
pyridaben	5,400
aldicarb-sulfone	70,800

# application



predict ionization efficiency

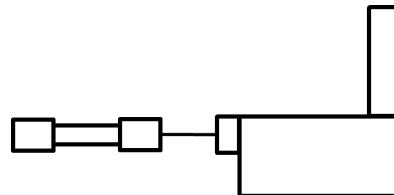
# application

<b>compound</b>	<b>peak area</b>	<b><math>\log E_{\text{pred}}</math></b>
methiocarb sulfoxide	5,300	2.57
pyridaben	5,400	3.78
aldicarb-sulfone	70,800	1.99

# application



predict ionization efficiency



convert to instrument specific values

# application

<b>compound</b>	<b>peak area</b>	<b><math>\log E_{\text{pred}}</math></b>	<b>c (nM)</b>
methiocarb sulfoxide	5,300	2.57	
pyridaben	5,400	3.78	
aldicarb-sulfone	70,800	1.99	
atrazine-D5			4.5
gabapentin-lactam			0.35
sitagliptin			0.23
5-methyl-1H-benzotriazole			0.94
neburon			3.4
caffeine			0.50

# application

<b>compound</b>	<b>peak area</b>	<b><math>\log E_{\text{pred}}</math></b>	<b>c (nM)</b>
methiocarb sulfoxide	5,300	2.57	
pyridaben	5,400	3.78	
aldicarb-sulfone	70,800	1.99	
atrazine-D5	450,000		4.5
gabapentin-lactam	10,400		0.35
sitagliptin	8,100		0.23
5-methyl-1H-benzotriazole	27,000		0.94
neburon	243,000		3.4
caffeine	5,600		0.50

# application

$RF_{\text{measured}} = \text{peak area} / c$

compound	peak area	$\log E_{\text{pred}}$	c (nM)	$RF_{\text{meas}} \cdot 10^{16}$
methiocarb sulfoxide	5,300	2.57		
pyridaben	5,400	3.78		
aldicarb-sulfone	70,800	1.99		
atrazine-D5	450,000		4.5	9.8
gabapentin-lactam	10,400		0.35	3.0
sitagliptin	8,100		0.23	3.5
5-methyl-1H-benzotriazole	27,000		0.94	2.9
neburon	243,000		3.4	7.2
caffeine	5,600		0.50	1.1

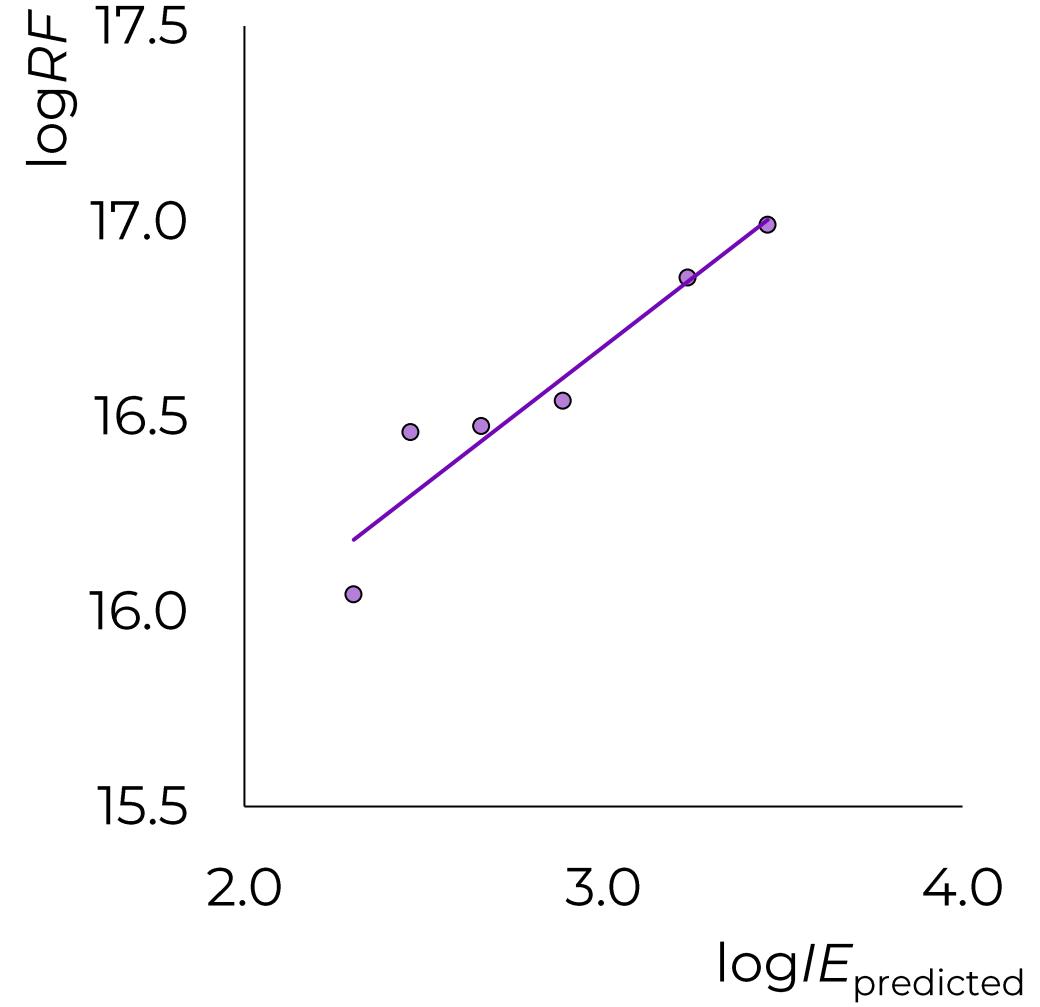
# application

<b>compound</b>	<b>peak area</b>	<b>log<i>E</i><sub>pred</sub></b>	<b>c (nM)</b>	<b><i>RF</i><sub>meas</sub> · 10<sup>16</sup></b>
methiocarb sulfoxide	5,300	2.57		
pyridaben	5,400	3.78		
aldicarb-sulfone	70,800	1.99		
atrazine-D5	450,000	3.46	4.5	9.8
gabapentin-lactam	10,400	2.66	0.35	3.0
sitagliptin	8,100	2.89	0.23	3.5
5-methyl-1H-benzotriazole	27,000	2.46	0.94	2.9
neburon	243,000	3.23	3.4	7.2
caffeine	5,600	2.30	0.50	1.1

# application

compound	peak area	$\log E_{\text{pred}}$
methiocarb sulfoxide	5,300	2.57
pyridaben	5,400	3.78
aldicarb-sulfone	70,800	1.99
atrazine-D5	450,000	3.46
gabapentin-lactam	10,400	2.66
sitagliptin	8,100	2.89
5-methyl-1H-benzotriazole	27,000	2.46
neburon	243,000	3.23
caffeine	5,600	2.30

$$\log RF = \text{slope} \cdot \log E + \text{intercept}$$



# application

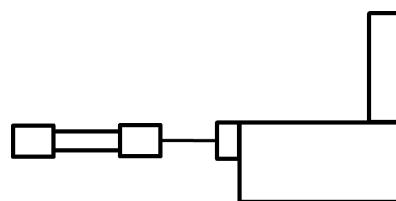
$$\log RF_{\text{predicted}} = \text{slope} \cdot \log I/E_{\text{predicted}} + \text{intercept}$$

compound	peak area	$\log I/E_{\text{pred}}$	c (nM)	$RF_{\text{meas}} \cdot 10^{16}$	$RF_{\text{pred}} \cdot 10^{16}$
methiocarb sulfoxide	5,300	2.57			2.6
pyridaben	5,400	3.78			15.5
aldicarb-sulfone	70,800	1.99			1.1
atrazine-D5	450,000	3.46	4.5	9.8	
gabapentin-lactam	10,400	2.66	0.35	3.0	
sitagliptin	8,100	2.89	0.23	3.5	
5-methyl-1H-benzotriazole	27,000	2.46	0.94	2.9	
neburon	243,000	3.23	3.4	7.2	
caffeine	5,600	2.30	0.50	1.1	

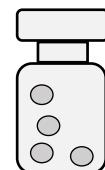
# application



predict ionization efficiency



convert to instrument specific values



estimate concentration

# application

$c = \text{peak area} / RF_{\text{predicted}}$

compound	peak area	$\log E_{\text{pred}}$	c (nM)	$RF_{\text{meas}} \cdot 10^{16}$	$RF_{\text{pred}} \cdot 10^{16}$	$c_{\text{pred}} (\text{nM})$
methiocarb sulfoxide	5,300	2.57			2.6	0.20
pyridaben	5,400	3.78			15.5	0.035
aldicarb-sulfone	70,800	1.99			1.1	6.3
atrazine-D5	450,000	3.46	4.5	9.8		
gabapentin-lactam	10,400	2.66	0.35	3.0		
sitagliptin	8,100	2.89	0.23	3.5		
5-methyl-1H-benzotriazole	27,000	2.46	0.94	2.9		
neburon	243,000	3.23	3.4	7.2		
caffeine	5,600	2.30	0.50	1.1		

# case studies



water

# chemicals in surface water

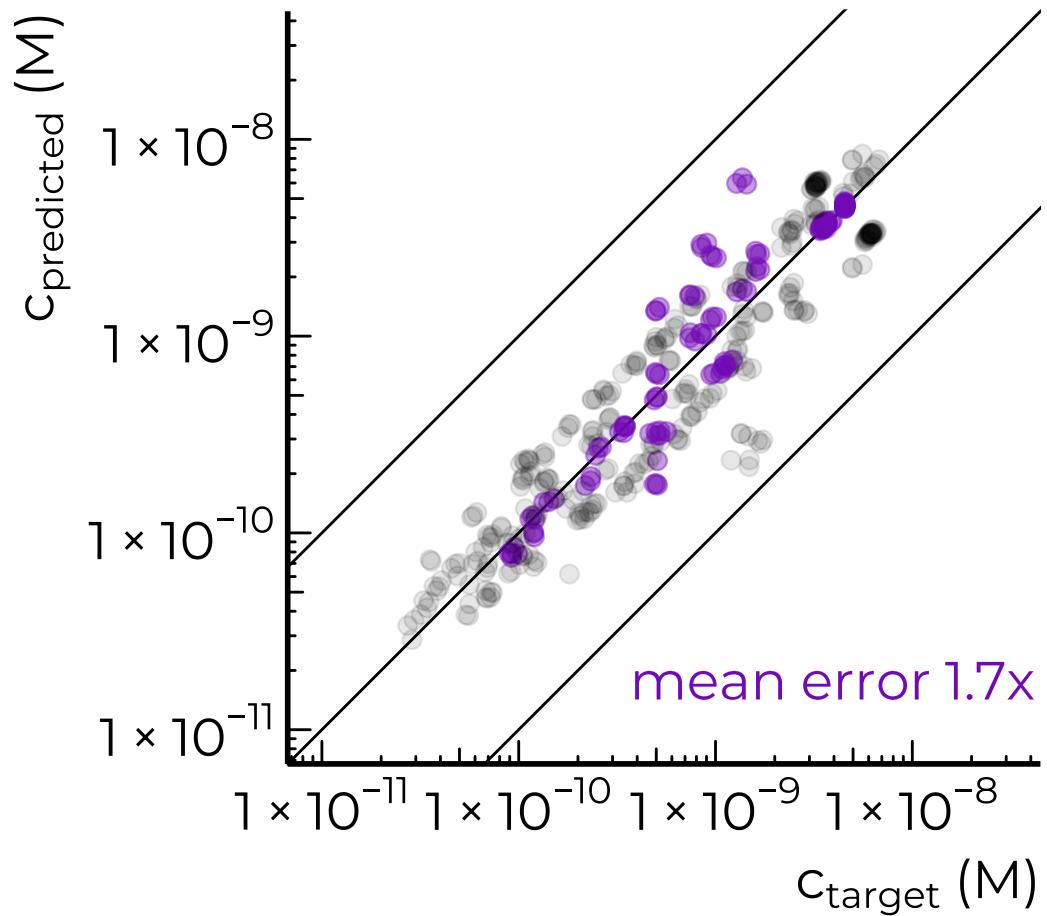
The Netherlands

Been et al. Water Research 2021

# chemicals in surface water

The Netherlands

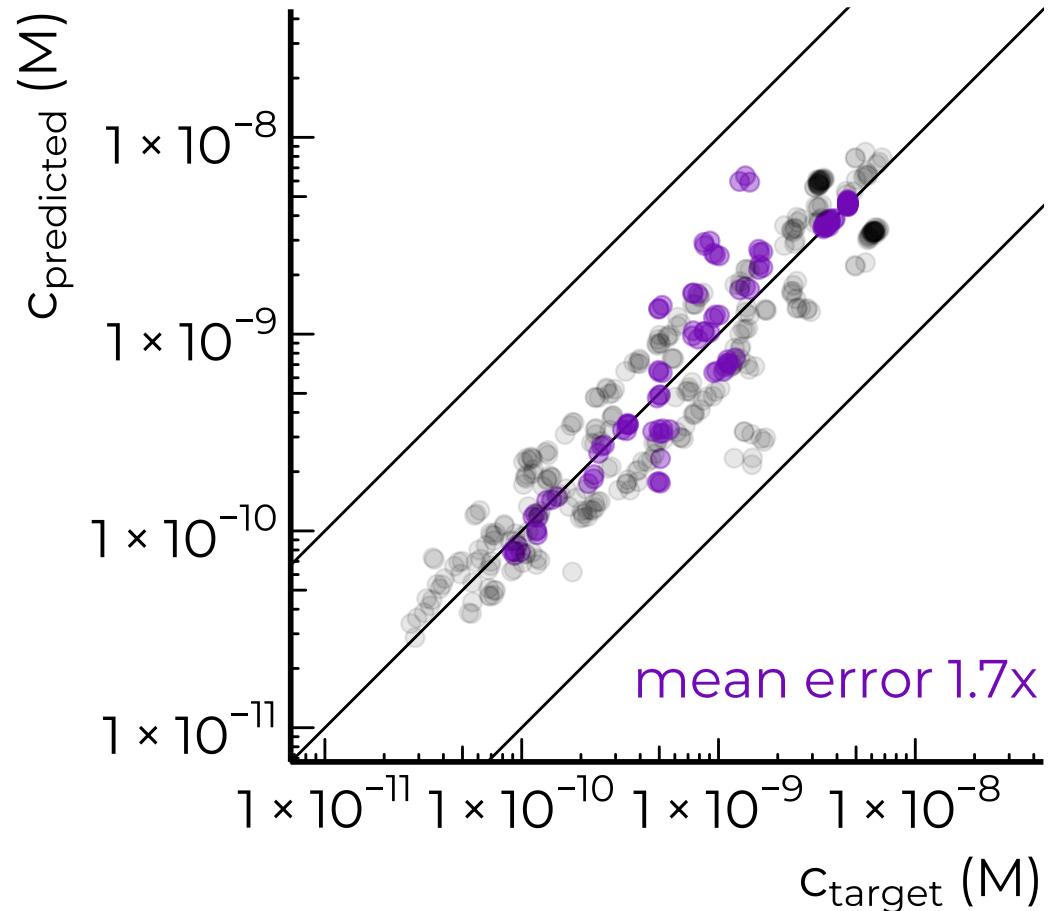
Been et al. Water Research 2021



# chemicals in surface water

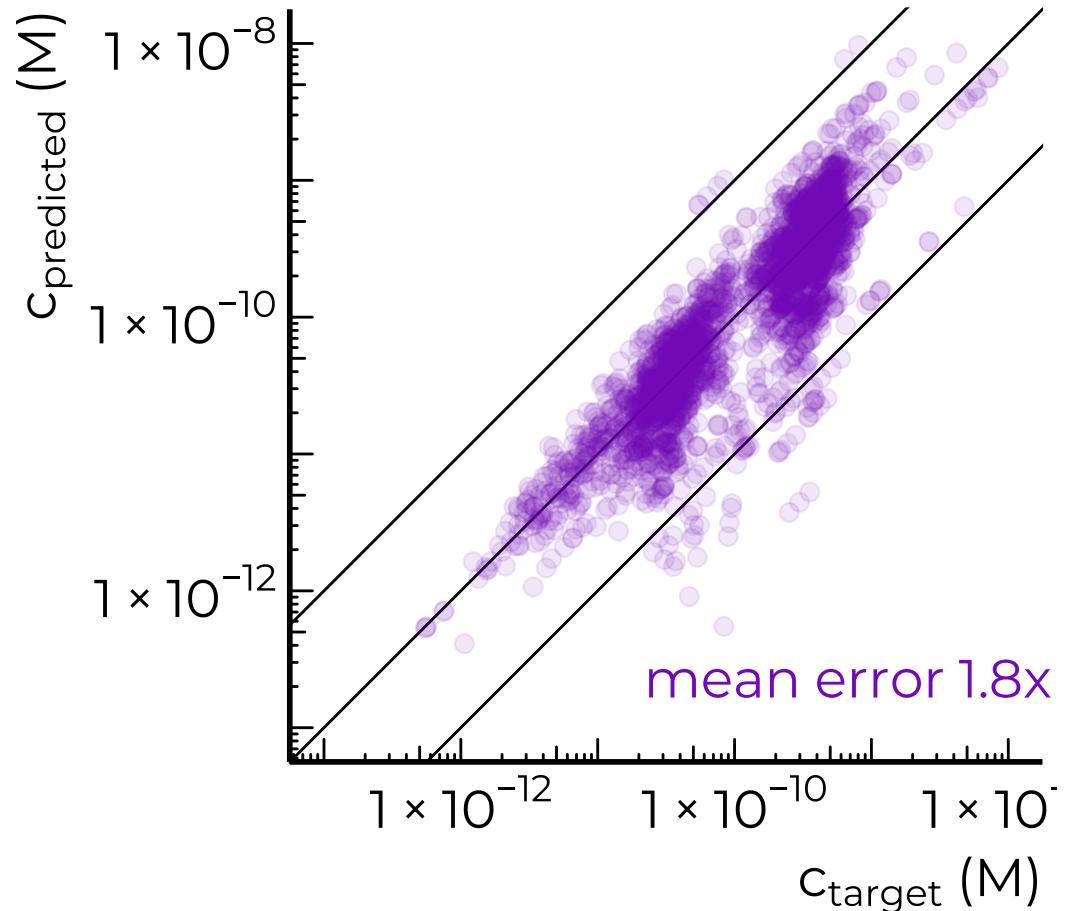
The Netherlands

Been et al. Water Research 2021



Switzerland

Kruve et al. Anal Bioanal Chem 2021



# case studies



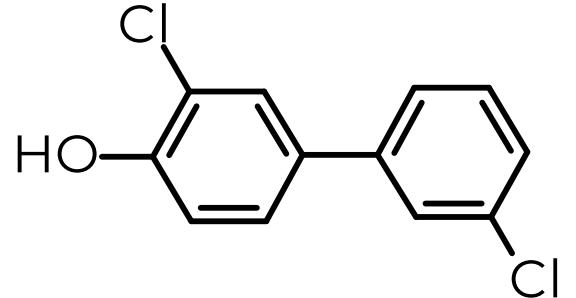
water



blood

# OH-PCBs in blood plasma

Khabazbashi et al. Anal Bioanal Chem 2022

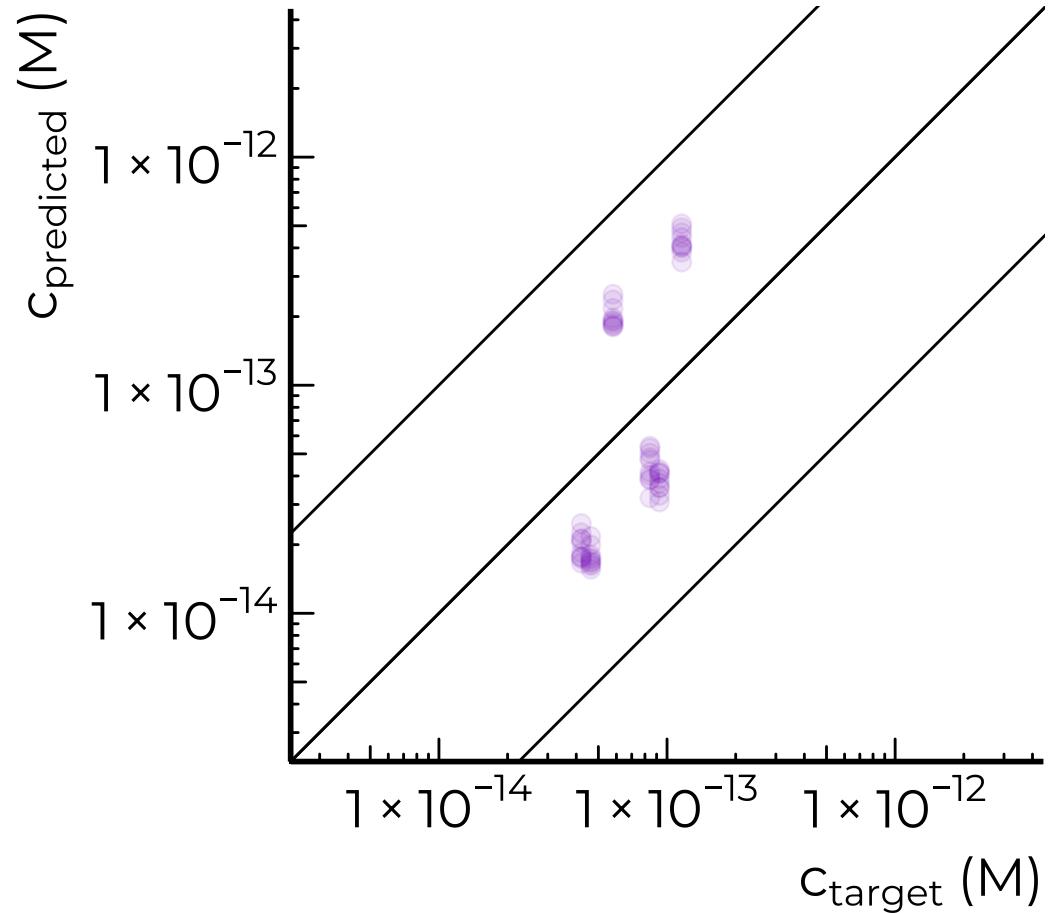
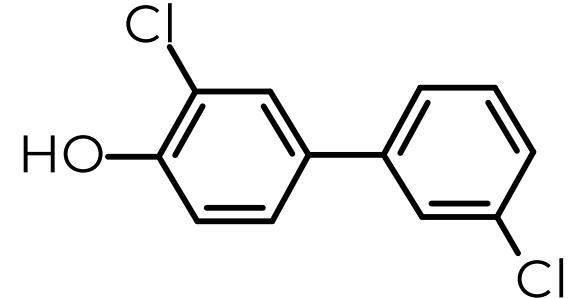


retrained with small set of OH-PCBs

tested on 3 OH-PCBs

# OH-PCBs in blood plasma

Khabazbashi et al. Anal Bioanal Chem 2022



retrained with small set of OH-PCBs

tested on 3 OH-PCBs

different matrixes

mean error 2.0x to 3.6x

# case studies



water



blood



food

# pesticides in food

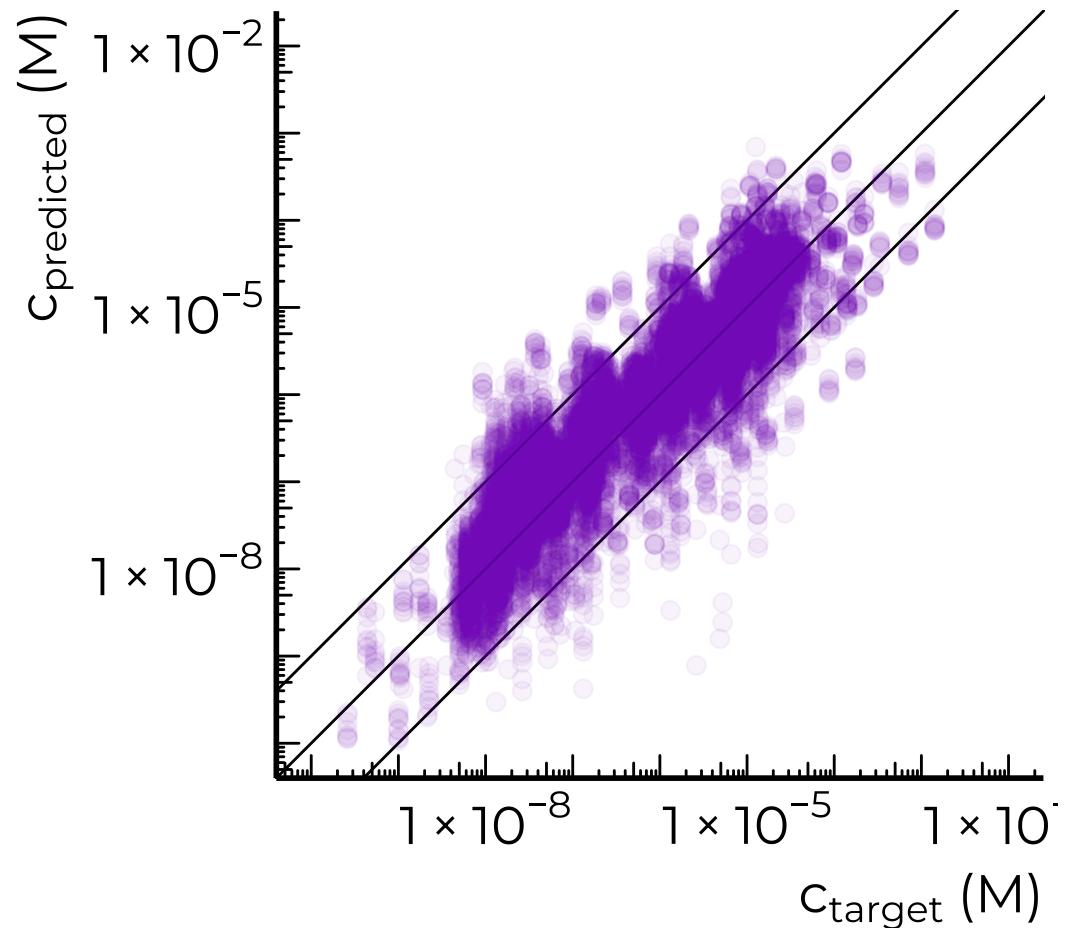
Wang et al. Food Chem 2020

195 pesticides in 5 cereal matrices

11,500 data points

# pesticides in food

Wang et al. Food Chem 2020



195 pesticides in 5 cereal matrices

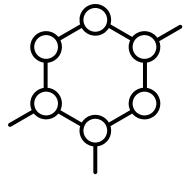
11,500 data points

mean error 5.3x

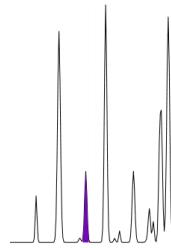
89% of points with error <10x

norman interlab

# tested methods



structurally similar chemicals  
transformation product - parent  
Tanimoto similarity

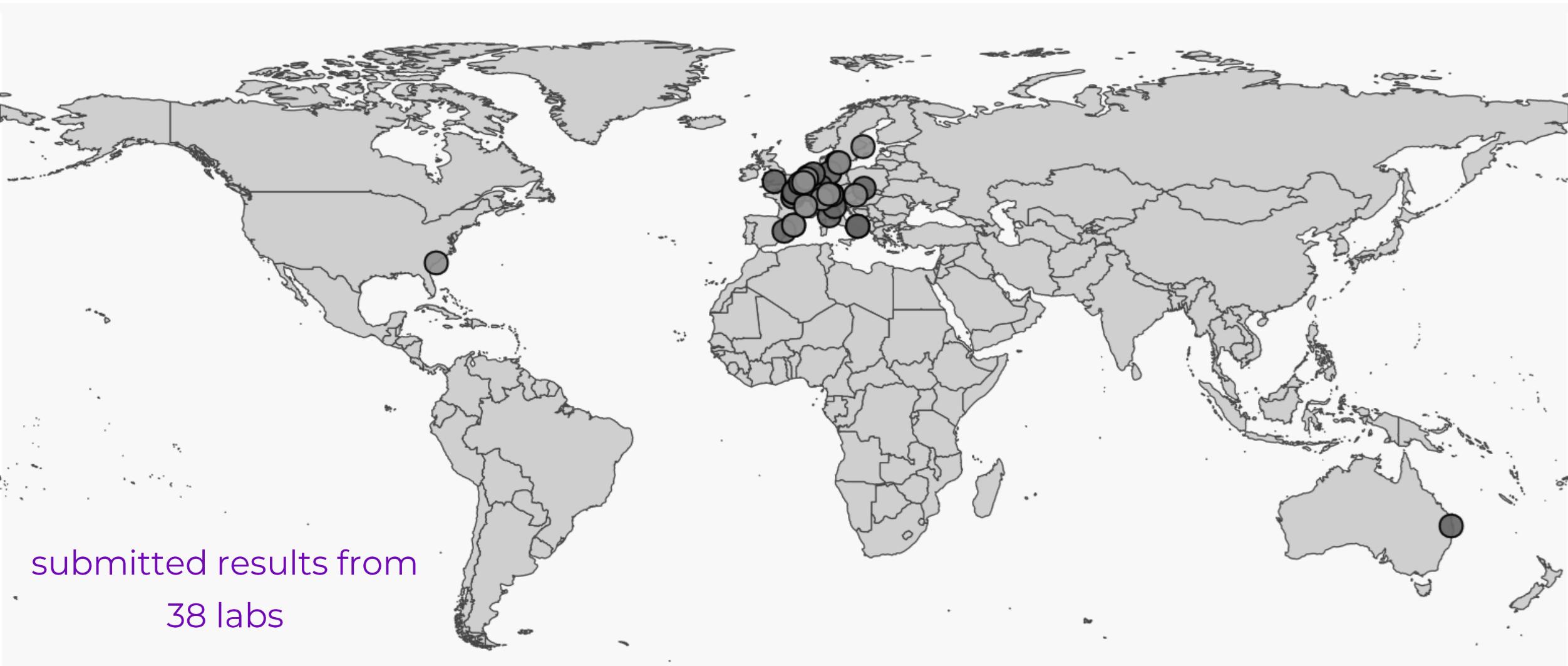


close eluting chemicals



machine learning  
Liigand et al.  
Aalizadeh et al.

# participants



# sample



HPLC water

# sample



HPLC water



drinking water

# sample



HPLC water

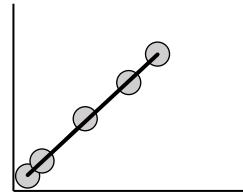


drinking water



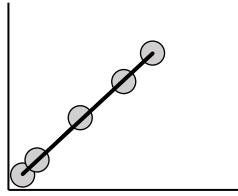
surface water

# chemicals

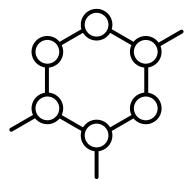


36 calibration chemicals  
known concentration

# chemicals

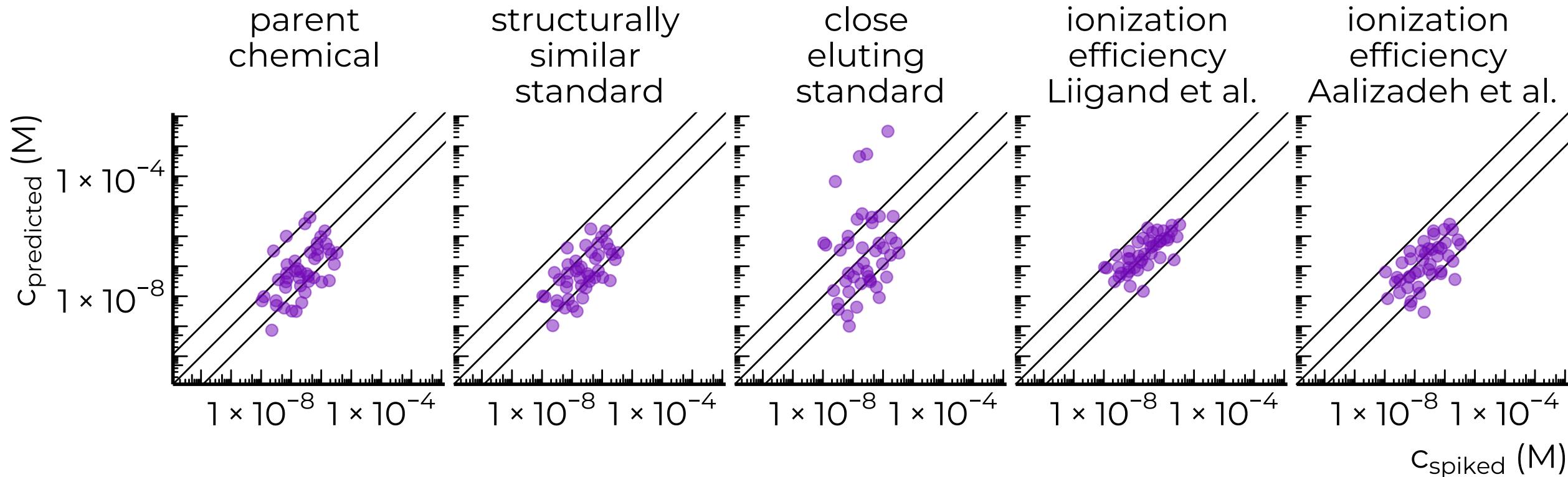


36 calibration chemicals  
known concentration

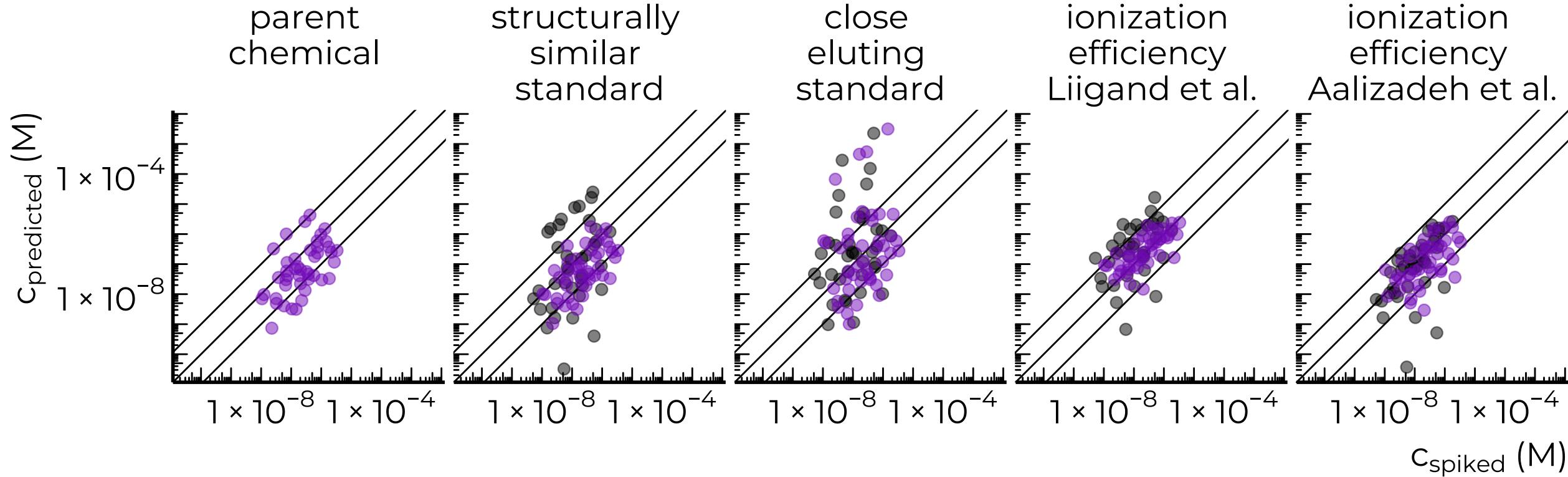


38 suspect chemicals  
low and high concentration spike

# quantification accuracy TPs

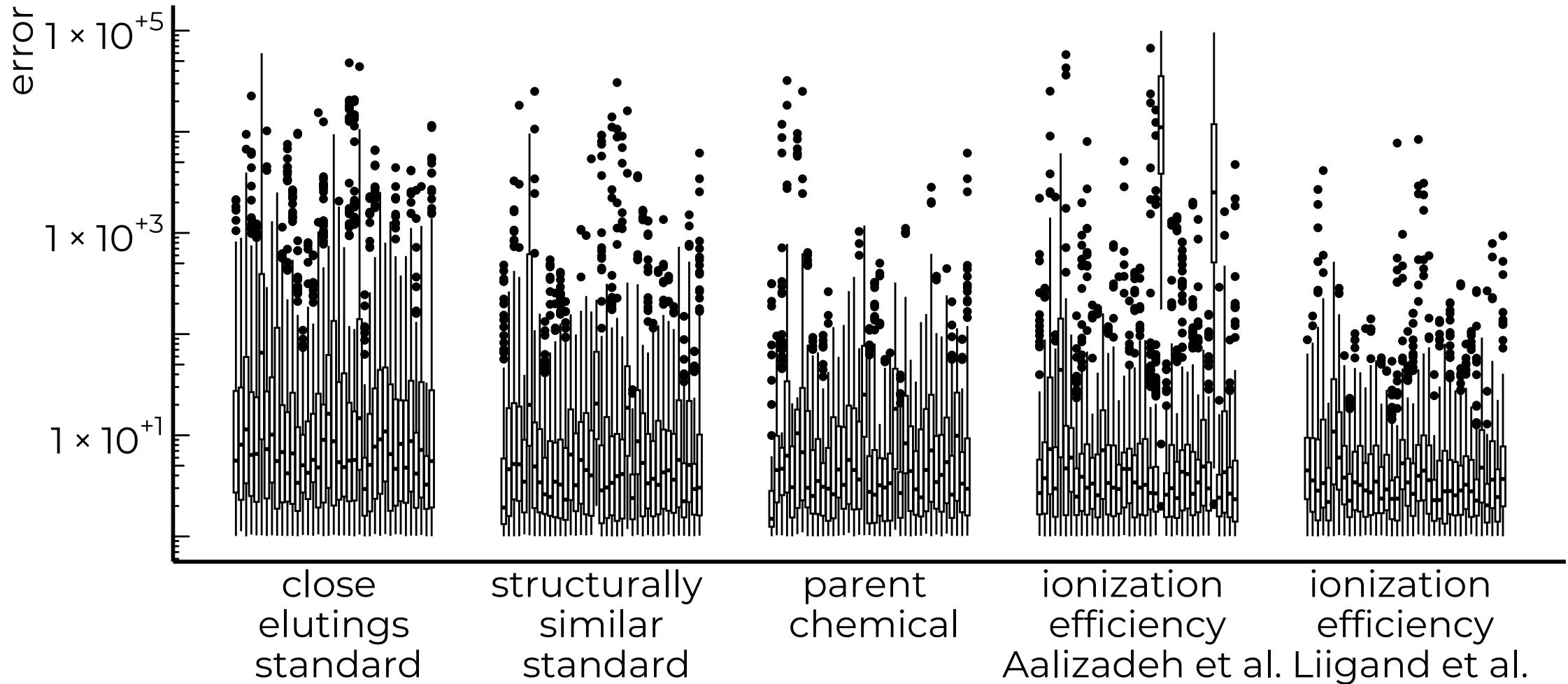


# quantification accuracy

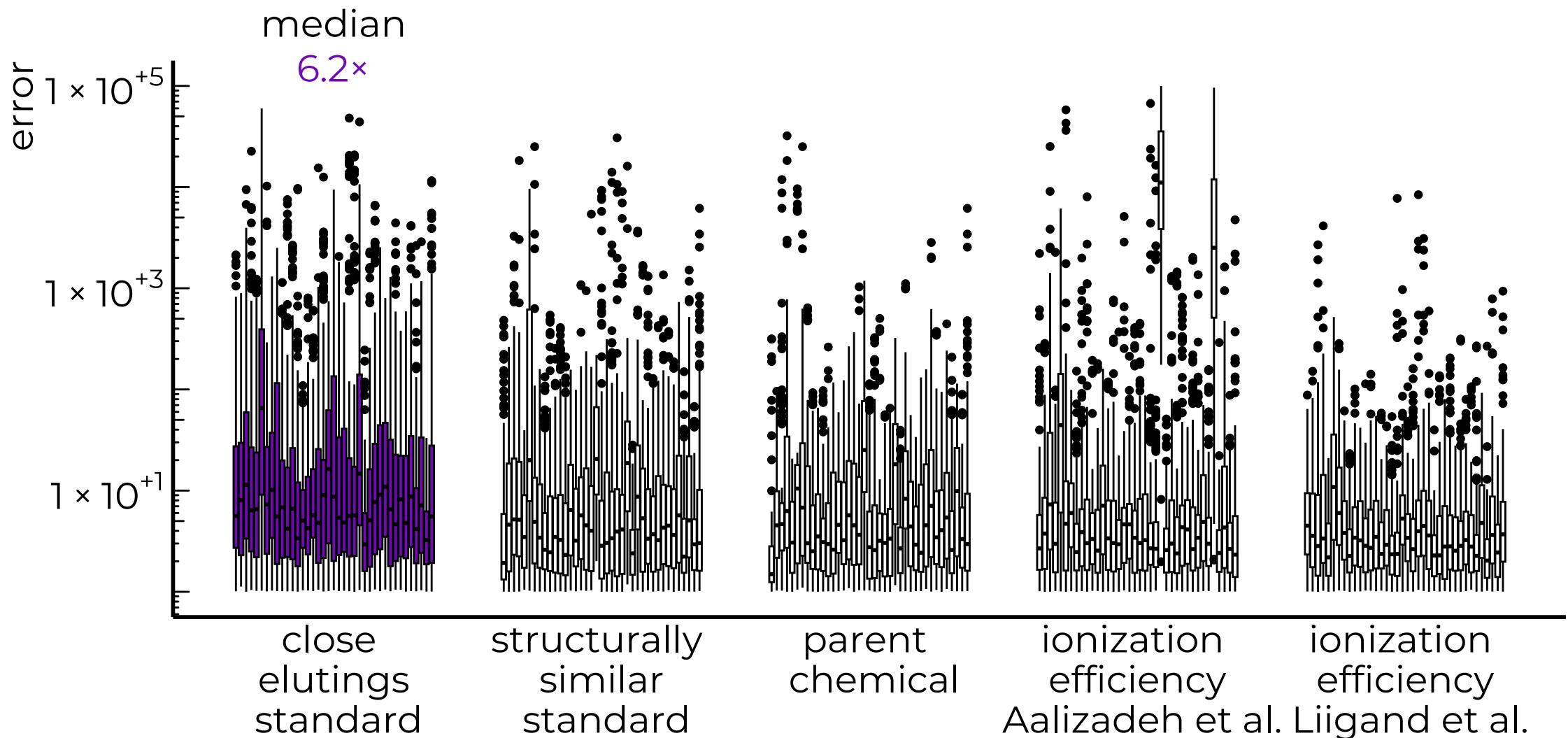


# prediction error

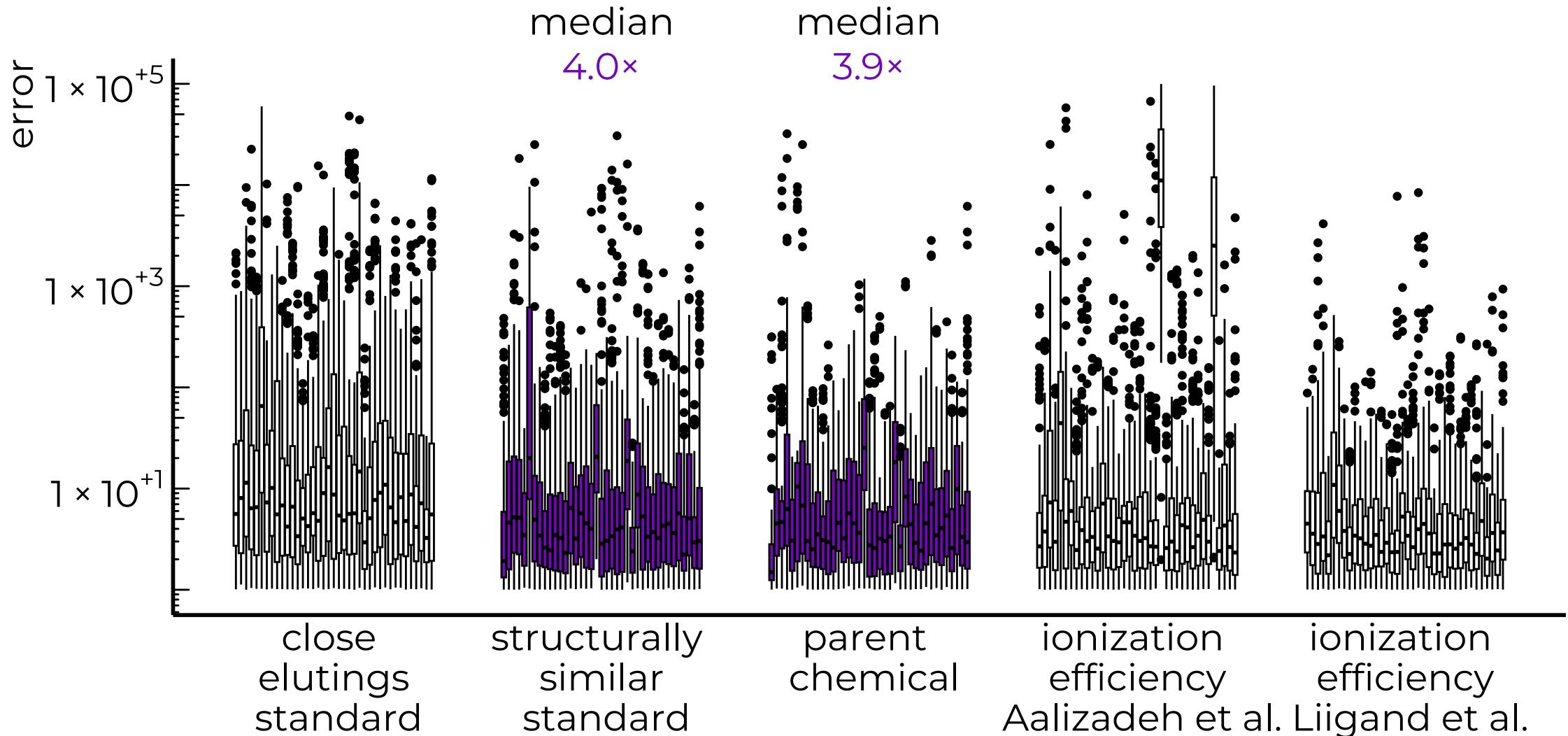
# prediction error



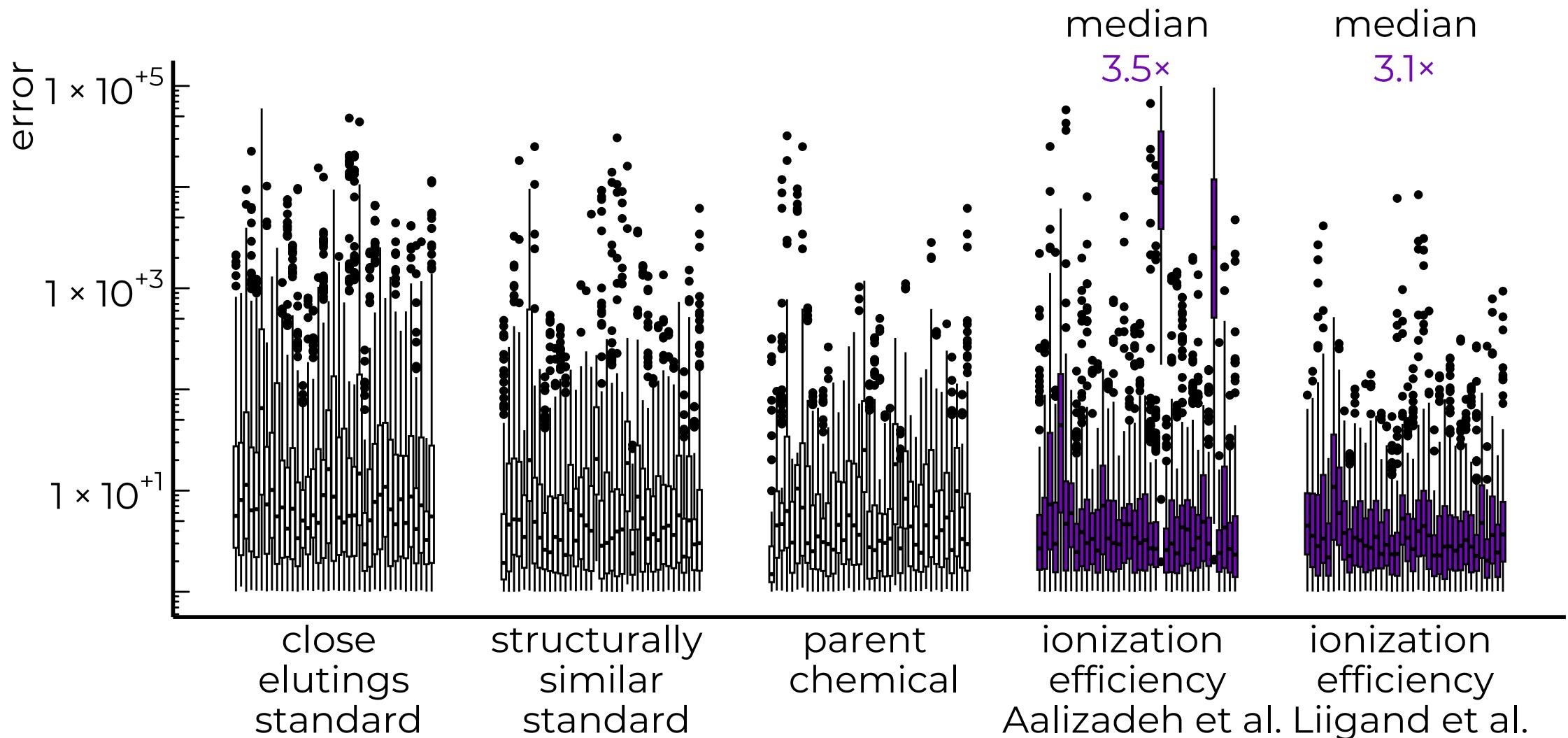
# prediction error



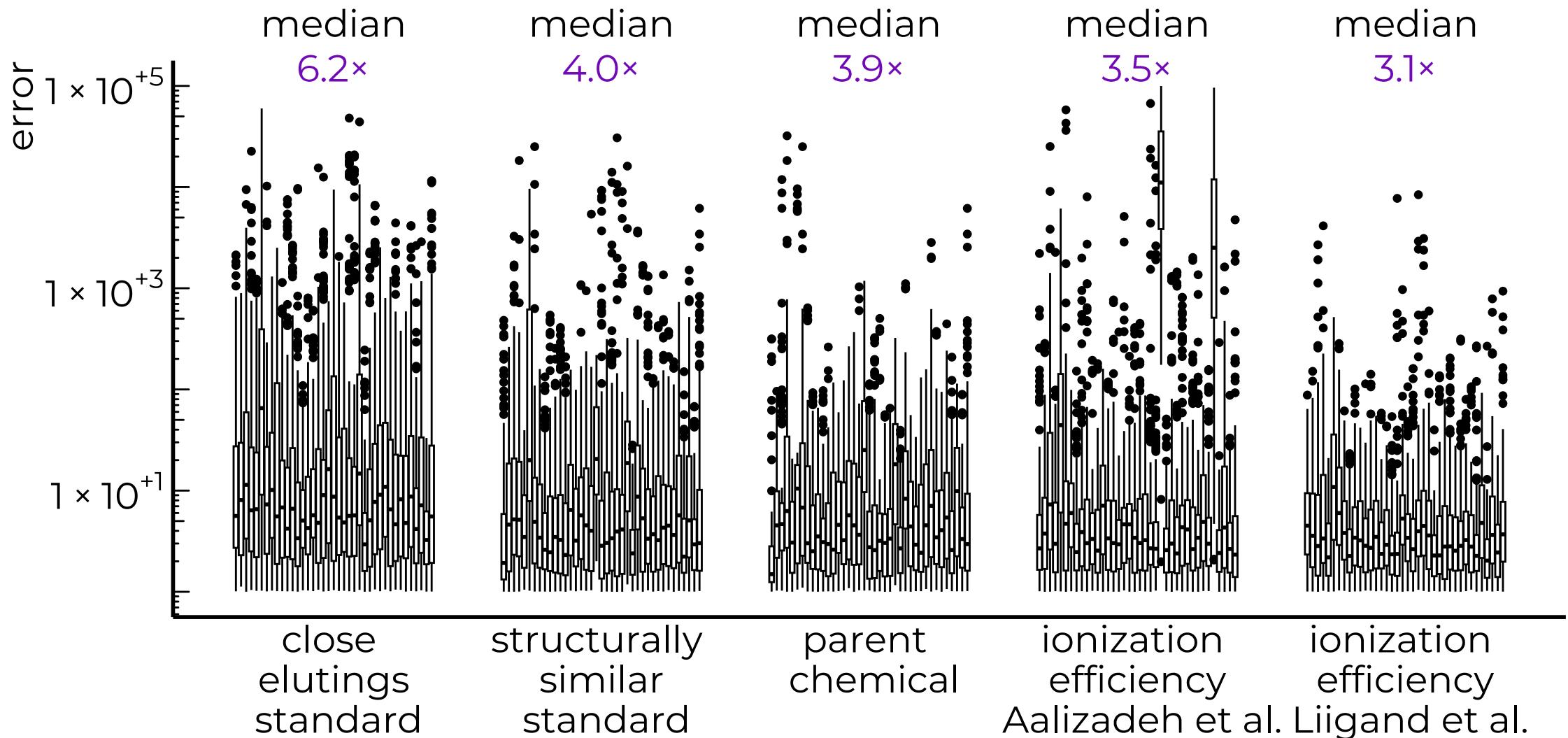
# prediction error



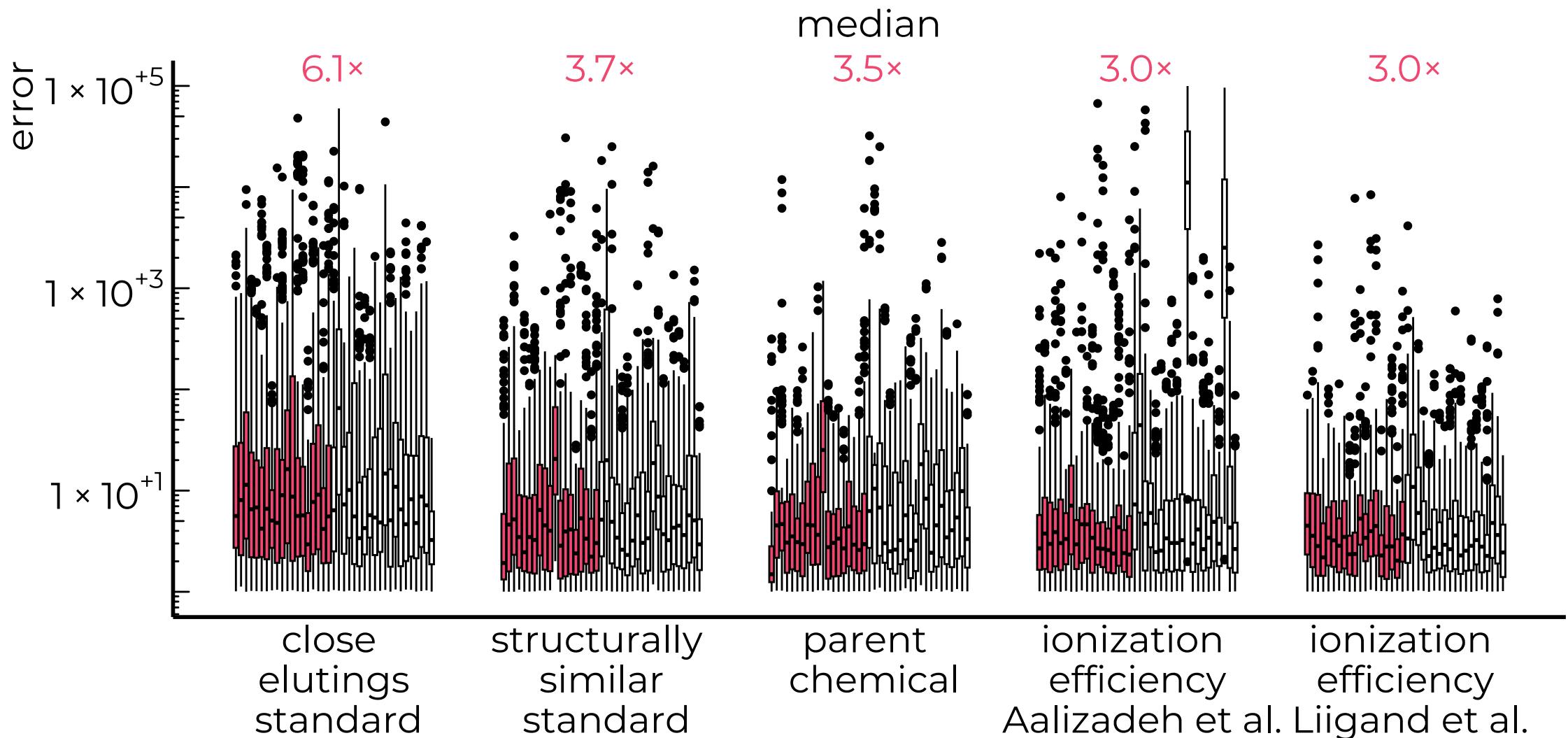
# prediction error



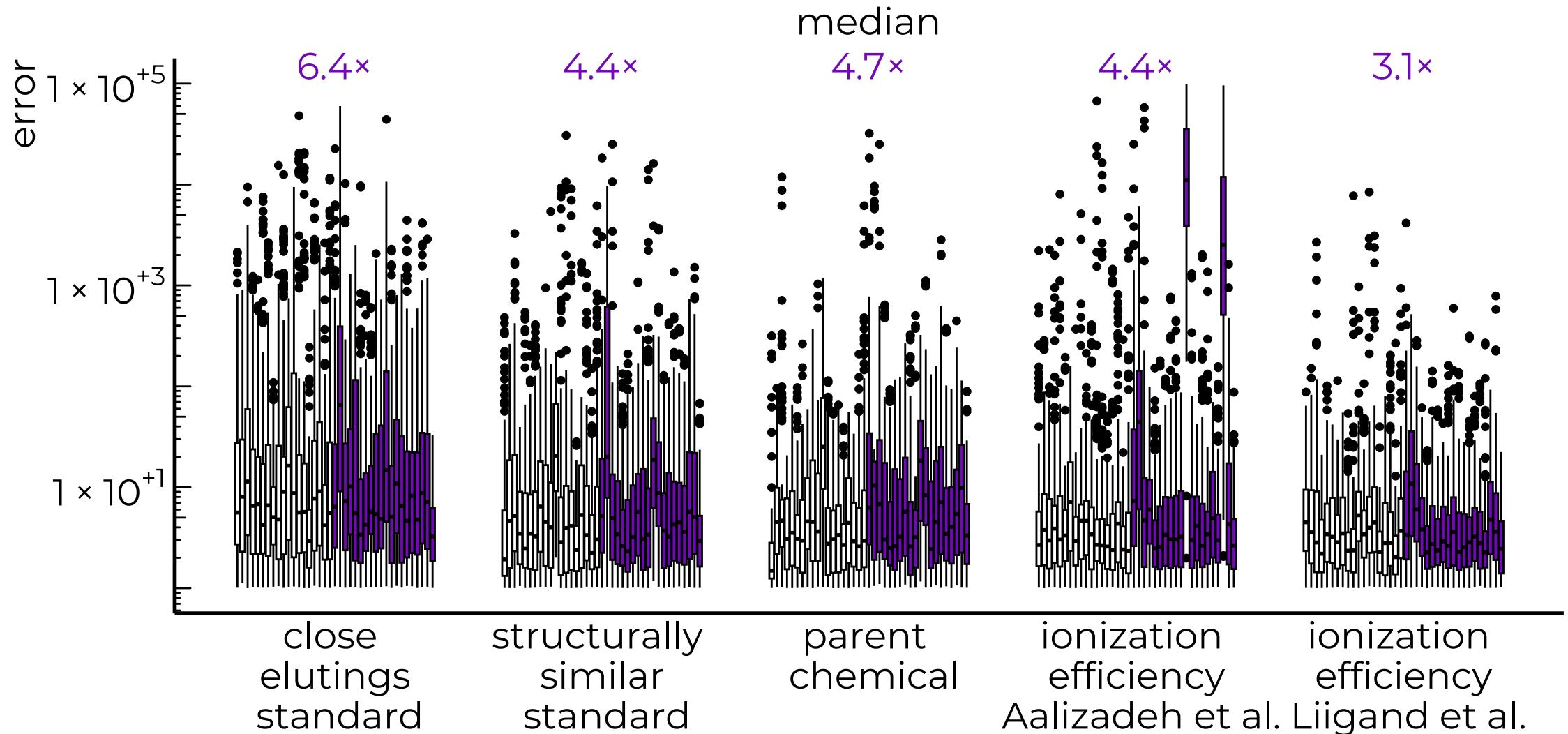
# prediction error



# prediction error orbitrap

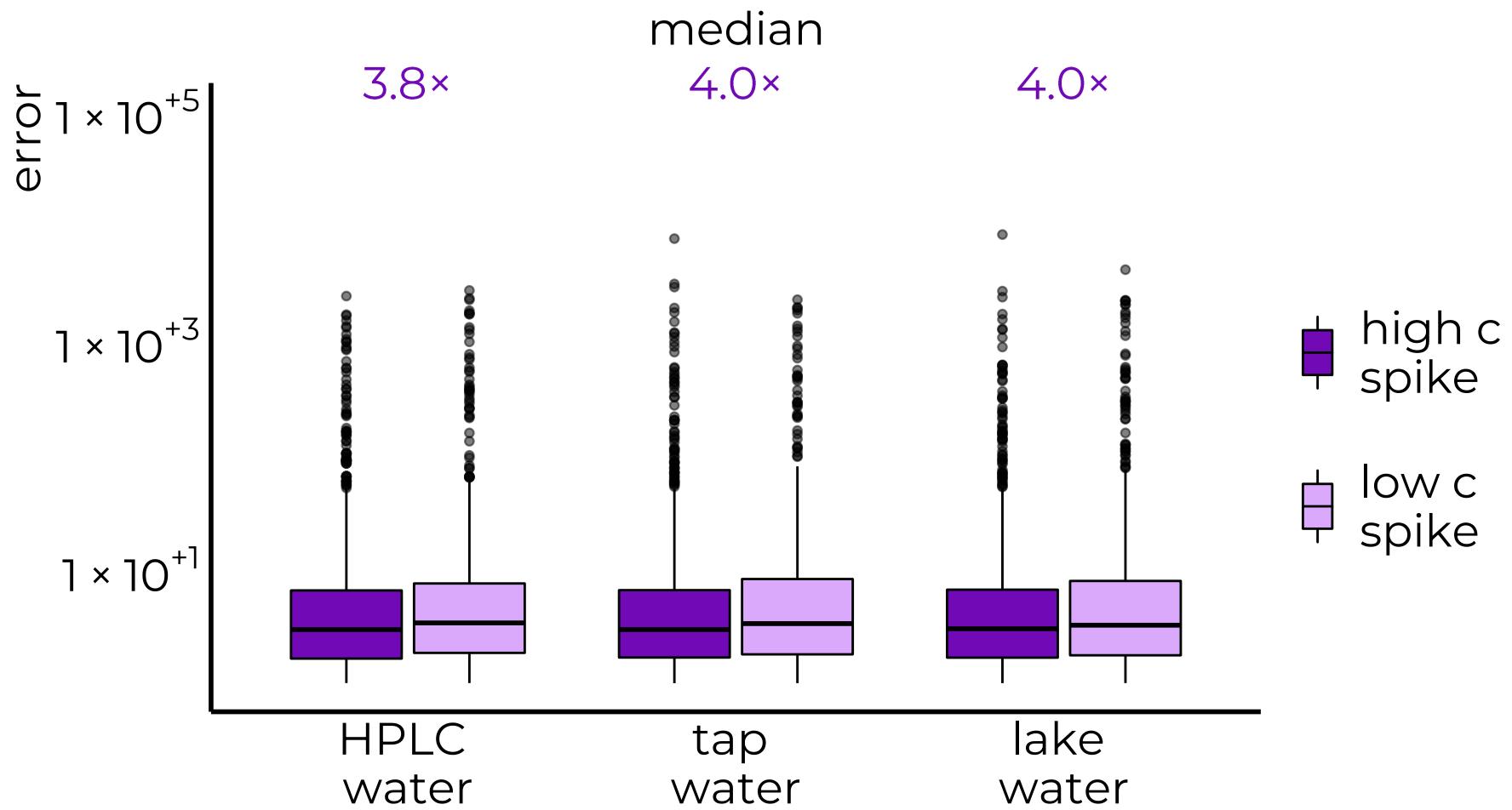


# prediction error time-of-flight



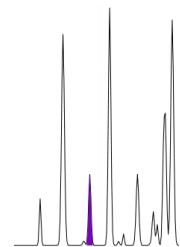
# across samples

# across samples



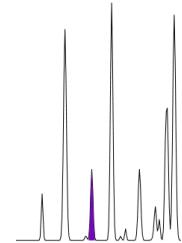
# conclusions

close eluting

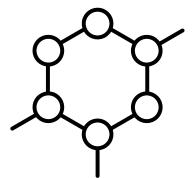


# conclusions

close eluting



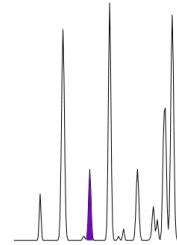
structurally similar



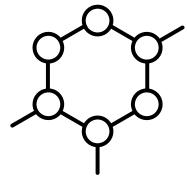
better

# conclusions

close eluting



structurally similar



better

ionization efficiency



best

# unidentified chemicals

from MS<sup>2</sup> spectra

# unidentified peaks

US EPA household dust study

# unidentified peaks

US EPA household dust study

3000 & 2000 features detected in ESI+ &ESI-

# unidentified peaks

US EPA household dust study

3000 & 2000 features detected in ESI+ &ESI-

978 formulas

# unidentified peaks

US EPA household dust study

3000 & 2000 features detected in ESI+ &ESI-

978 formulas → 3228 possible structures

# unidentified peaks

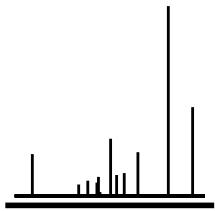
US EPA household dust study

3000 & 2000 features detected in ESI+ &ESI-

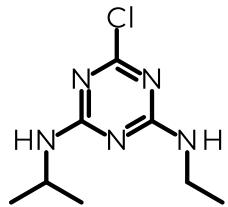
978 formulas → 3228 possible structures

33 identified

# workflow



MS<sup>2</sup> spectra



structure as SMILES

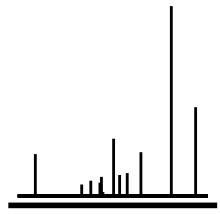


molecular descriptors



predict toxicity and ionization efficiency

# workflow



MS<sup>2</sup> spectra

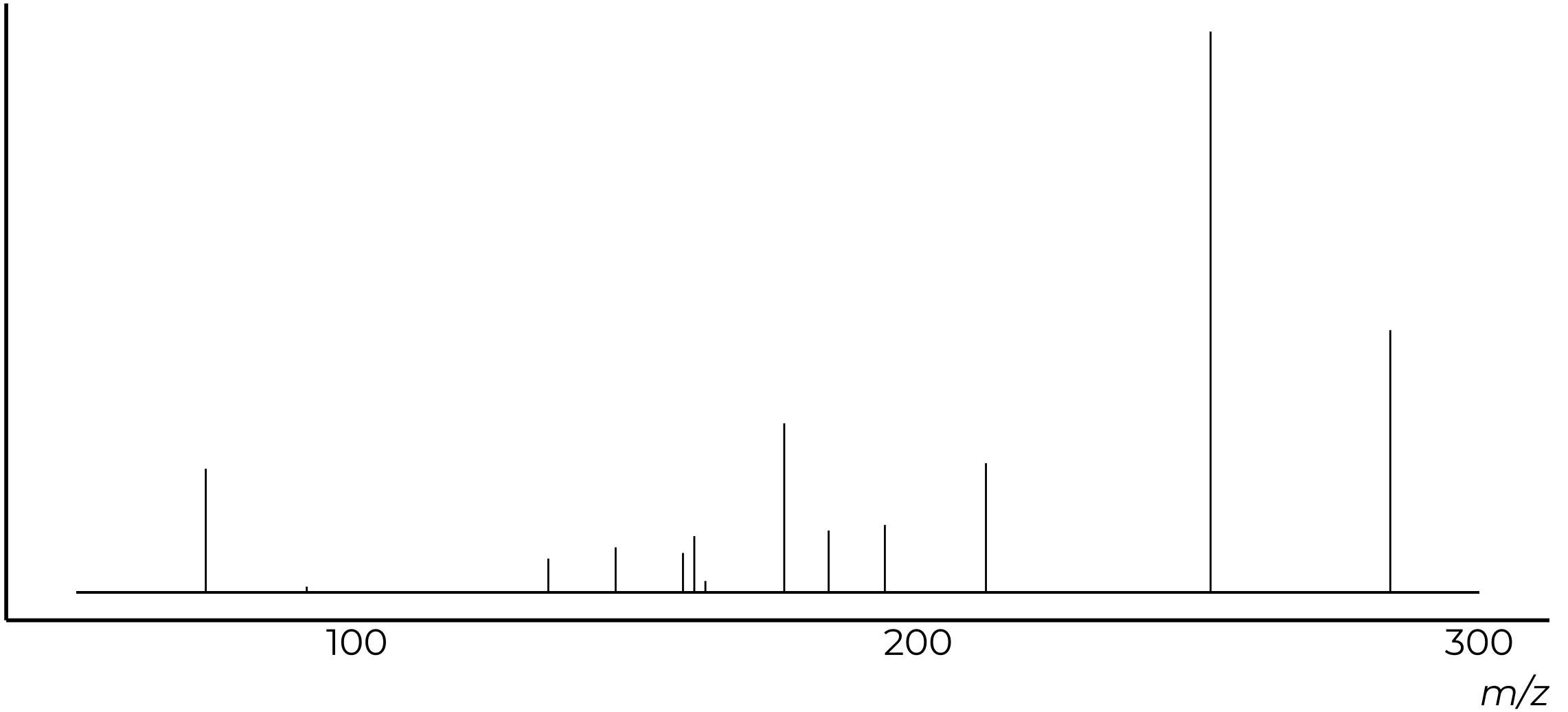


molecular fingerprints

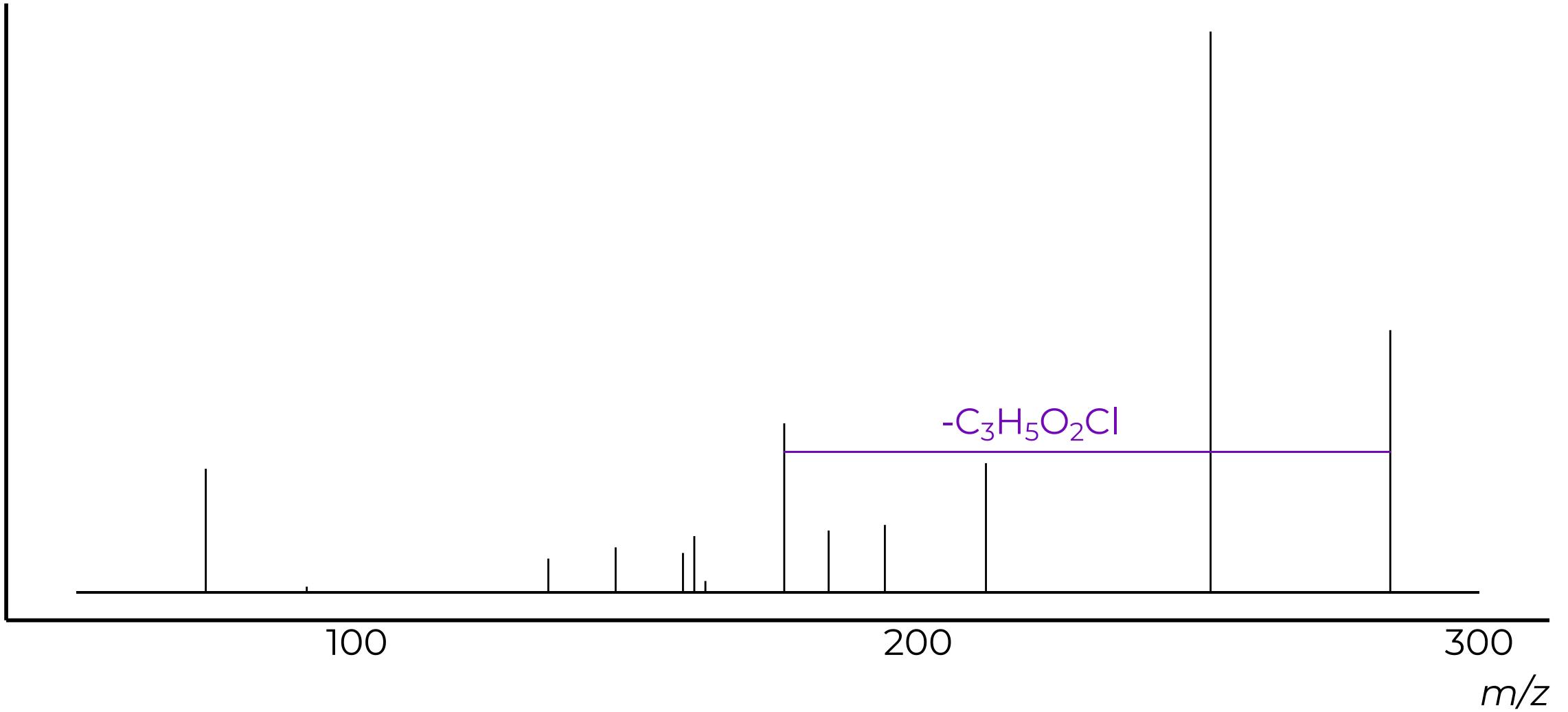


predict toxicity and ionization efficiency

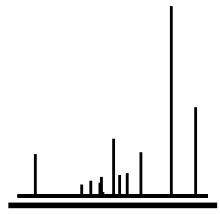
# MS<sup>2</sup> spectra



# MS<sup>2</sup> spectra



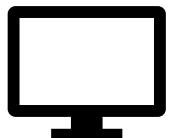
# workflow



MS<sup>2</sup> spectra



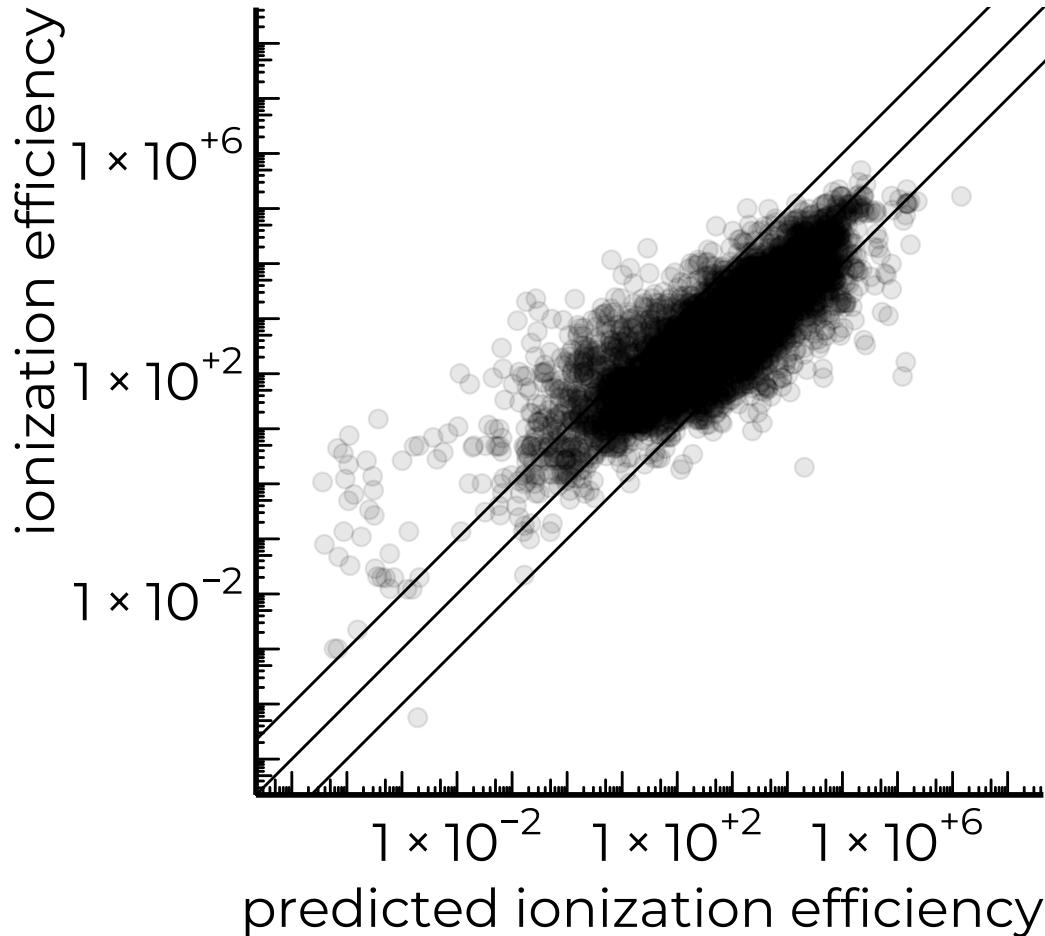
molecular fingerprints with SIRIUS



predict toxicity and ionization efficiency

# ionization efficiency

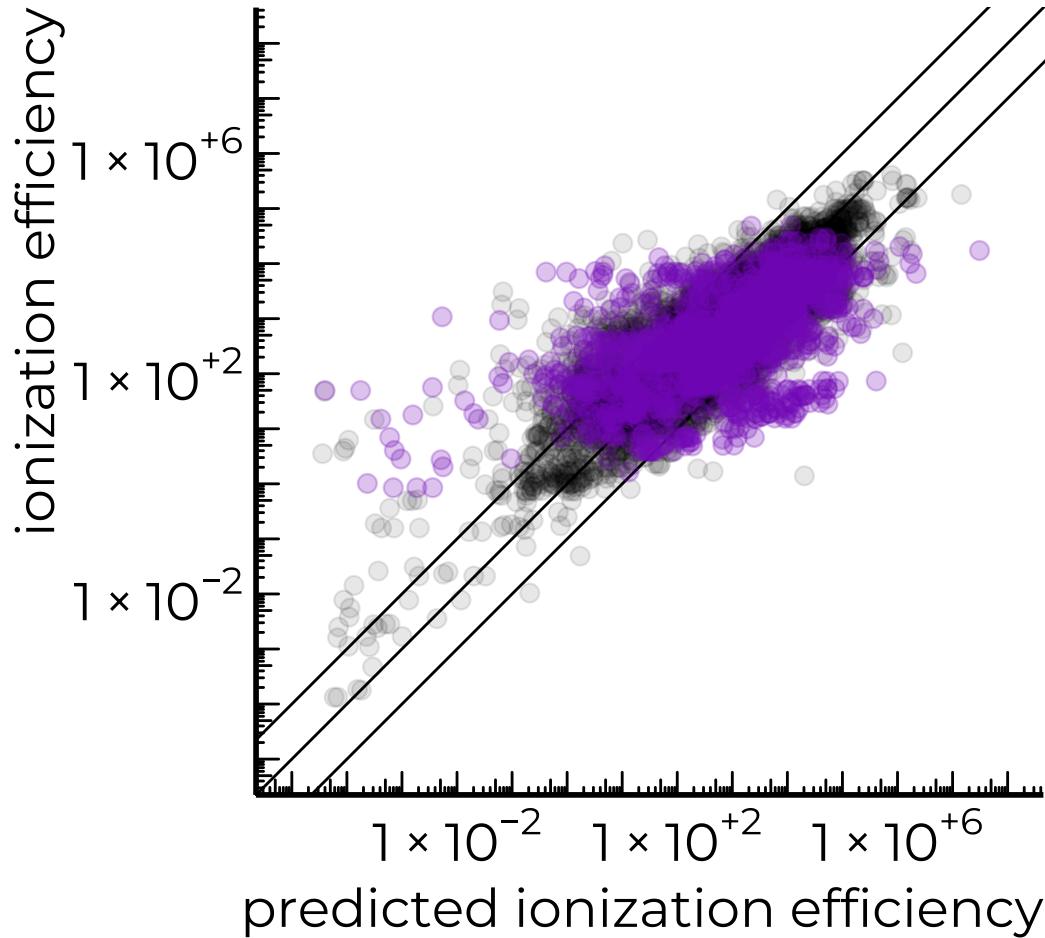
Sepman et al. in review



IE range  
100,000,000  
training set  
RMSE 3.5x

# ionization efficiency

Sepman et al. in review



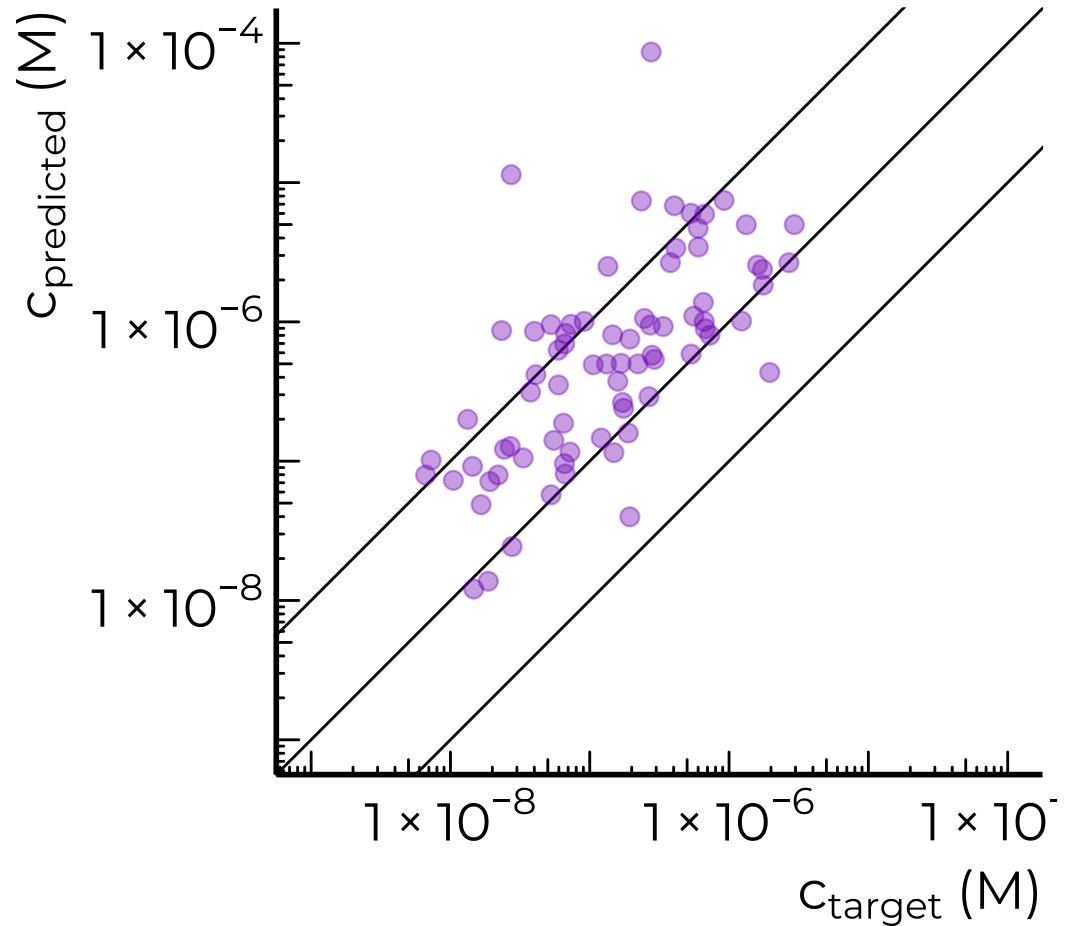
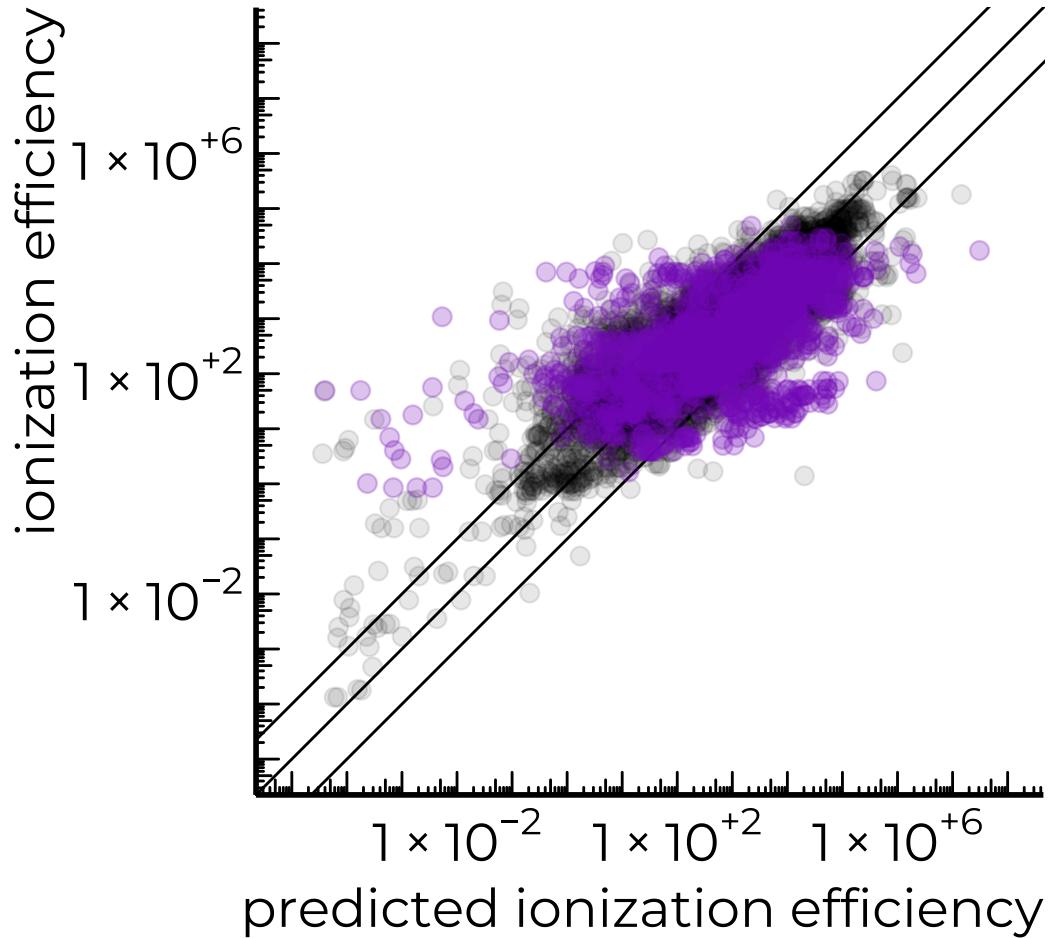
IE range  
100,000,000

training set  
RMSE 3.5x

test set  
RMSE 6.3x

# ionization efficiency

Sepman et al. in review



# ionization efficiency

Sepman et al. in review

mean prediction error

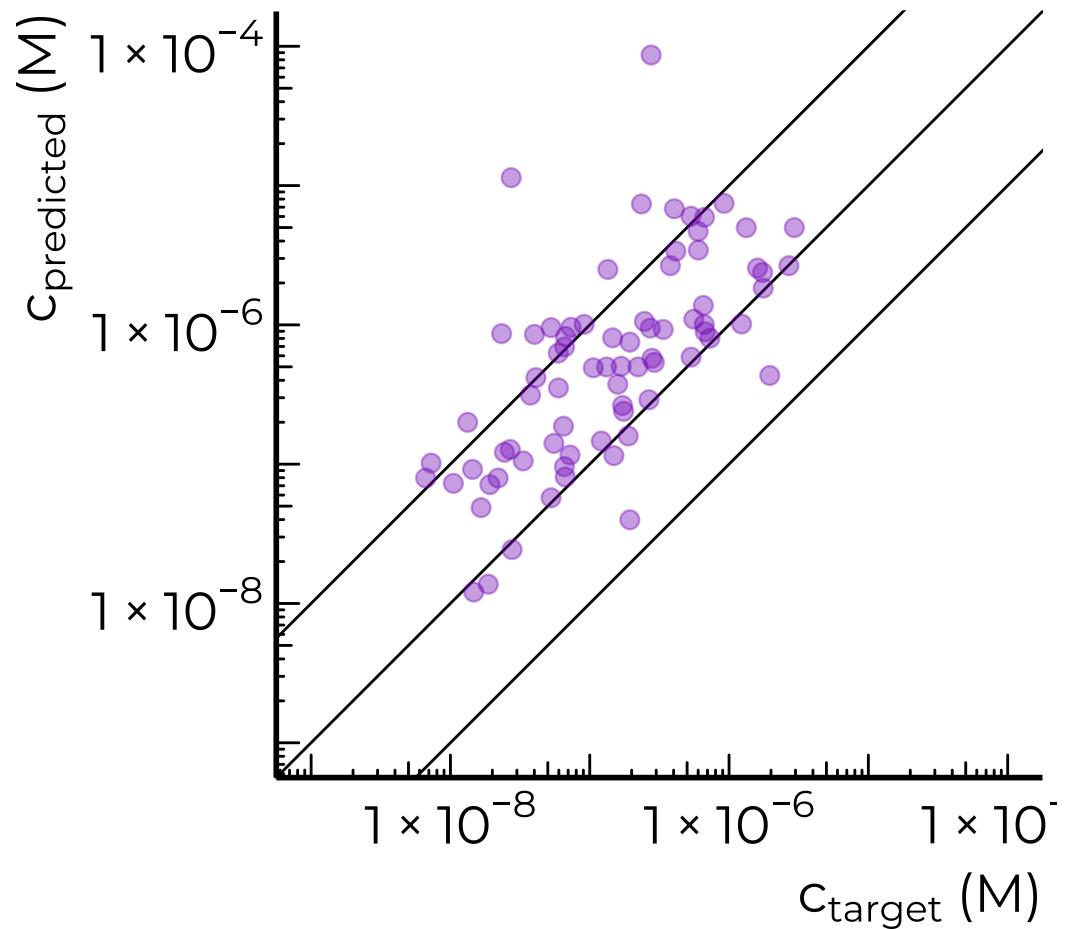
7.4x

geometric mean prediction error

4.5x

median prediction error

4.0x



summary

# prioritization in NTS

toxicity



concentration



risk



# prioritization in NTS

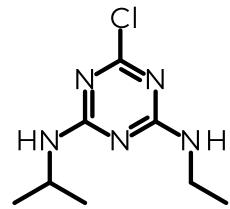
toxicity



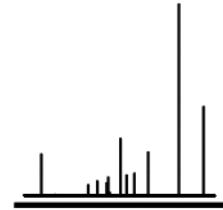
concentration



risk



structure



MS<sup>2</sup> spectrum



kruvelab.com

anneli.kruve@su.se