A Cross-Entropy Method that Optimizes Partially Decomposable Problems:

A New Way to Interpret NMR Spectra

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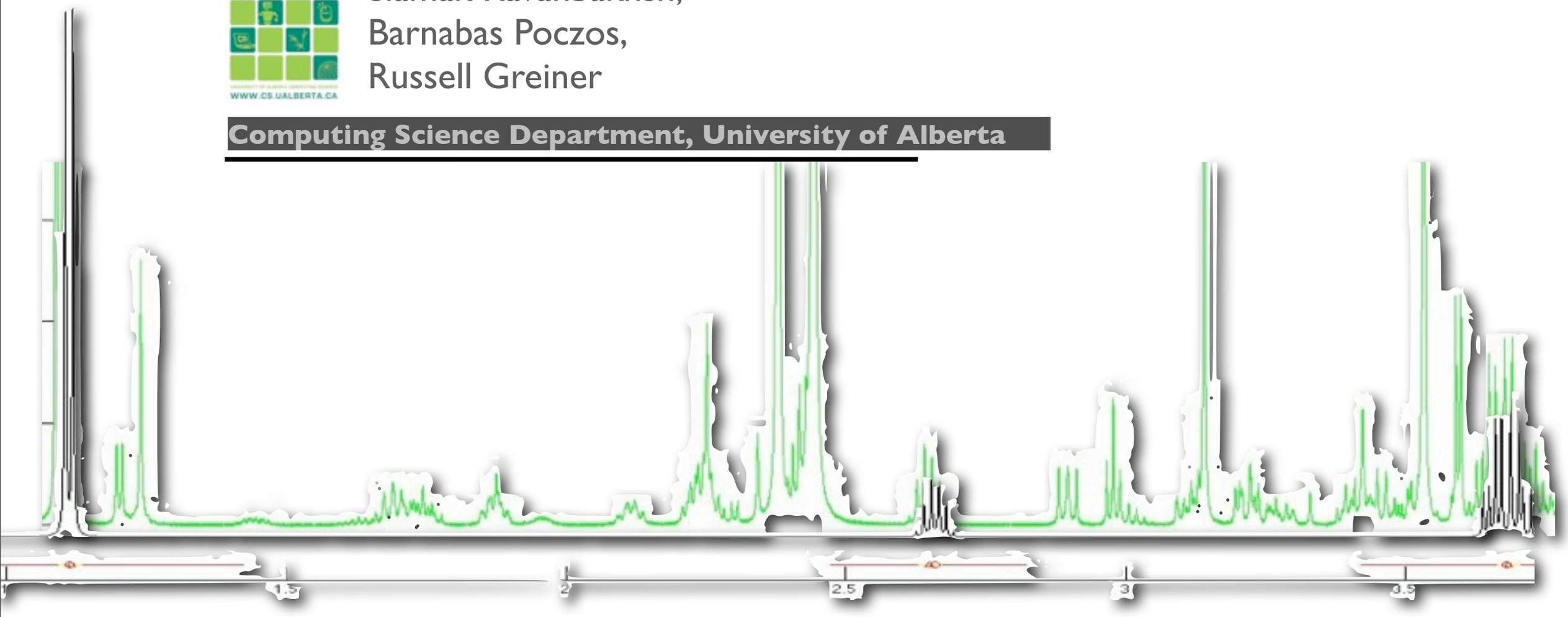

Metabolomics & NMR Spectroscopy

Metabolomics study of chemical fingerprints that cellular processes leave behind

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Metabolites: *end products of gene expression*

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Metabolites: *end products of gene expression*

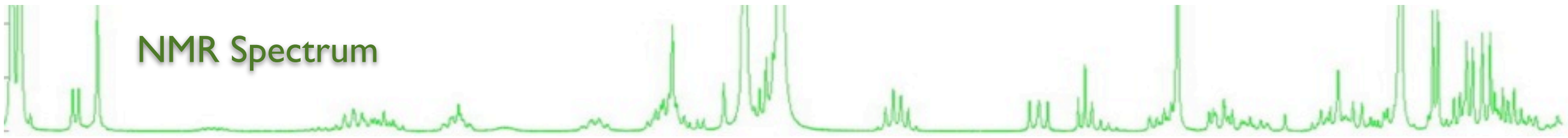
Toolbox:

- Gas chromatography
- High performance liquid chromatography
- Capillary electrophoresis
- Mass Spectrometry
- **Nuclear Magnetic Resonance (NMR) Spectroscopy**

source: wikipedia

Nuclear Magnetic Resonance (NMR) Spectroscopy

Nuclear Magnetic Resonance (NMR) Spectroscopy



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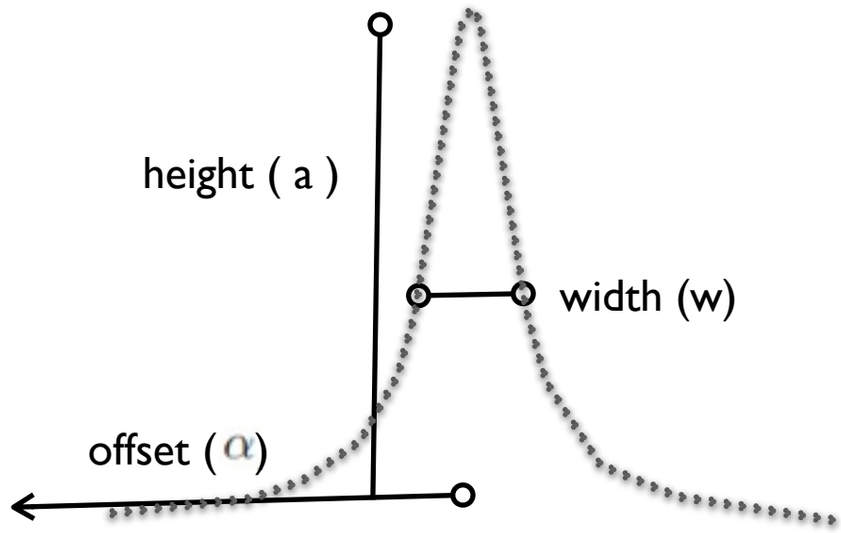
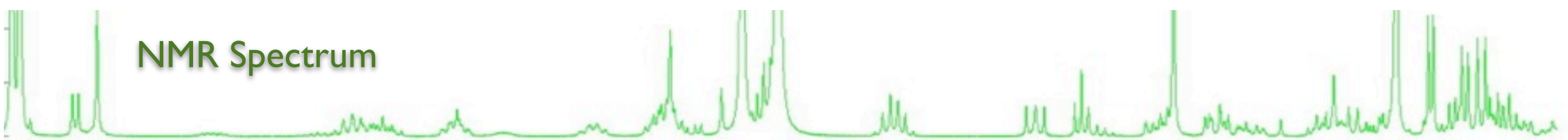
Nuclear Magnetic Resonance (NMR) Spectroscopy

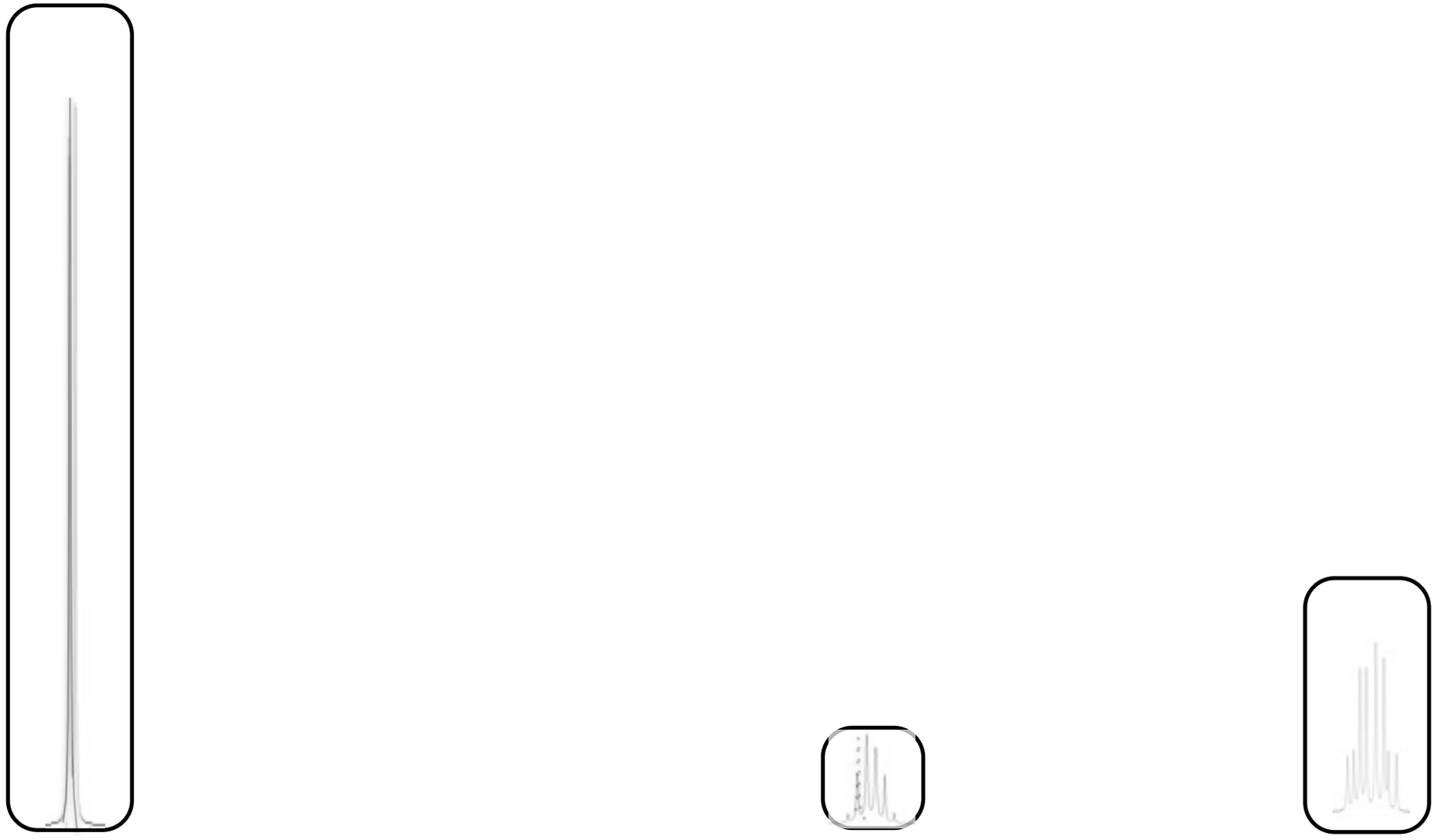
Spectrum is made of many

Lorentzian peaks

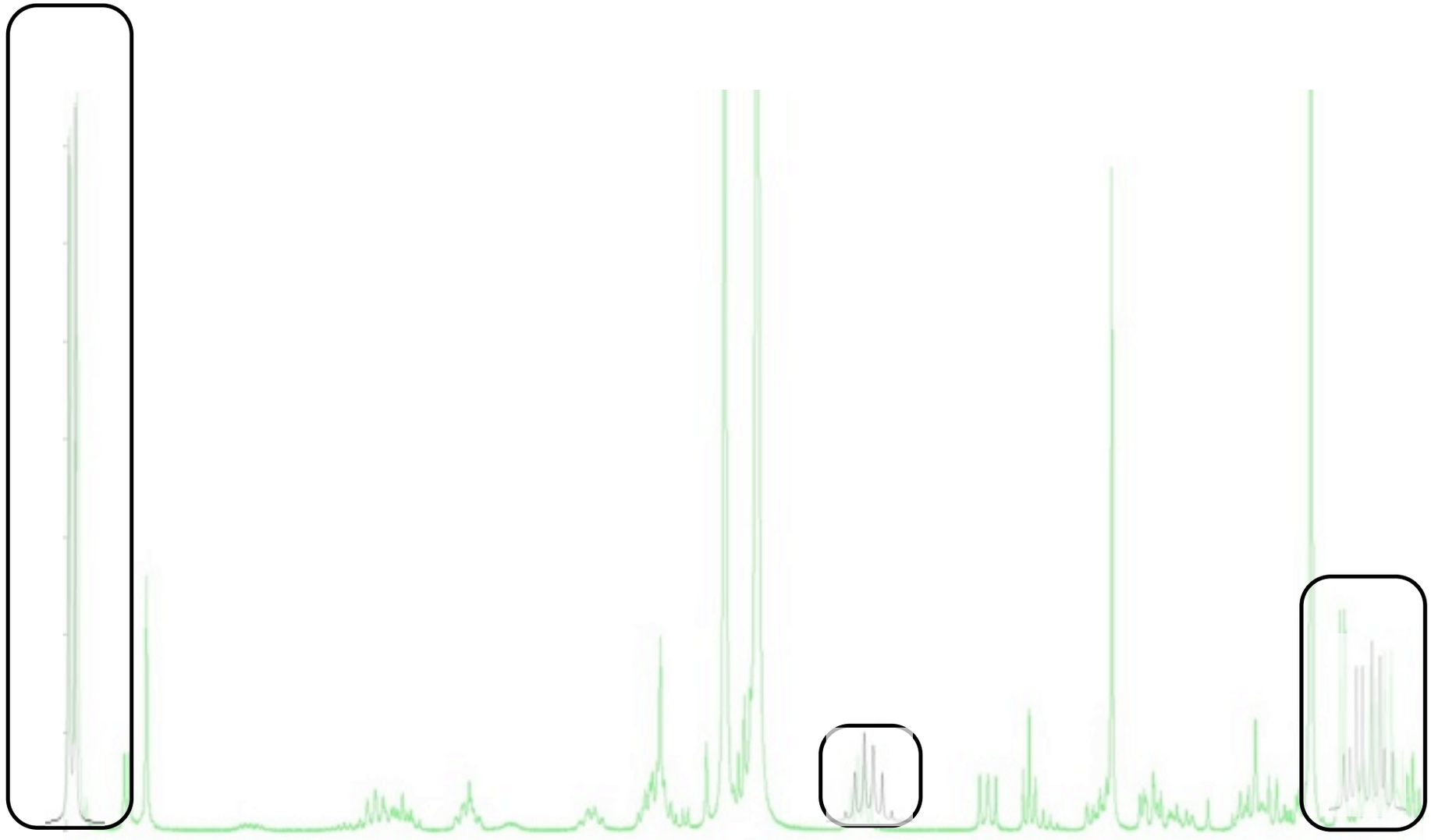


$$S(y) = \frac{aw^2}{w^2 + 4(y-\alpha)^2}$$

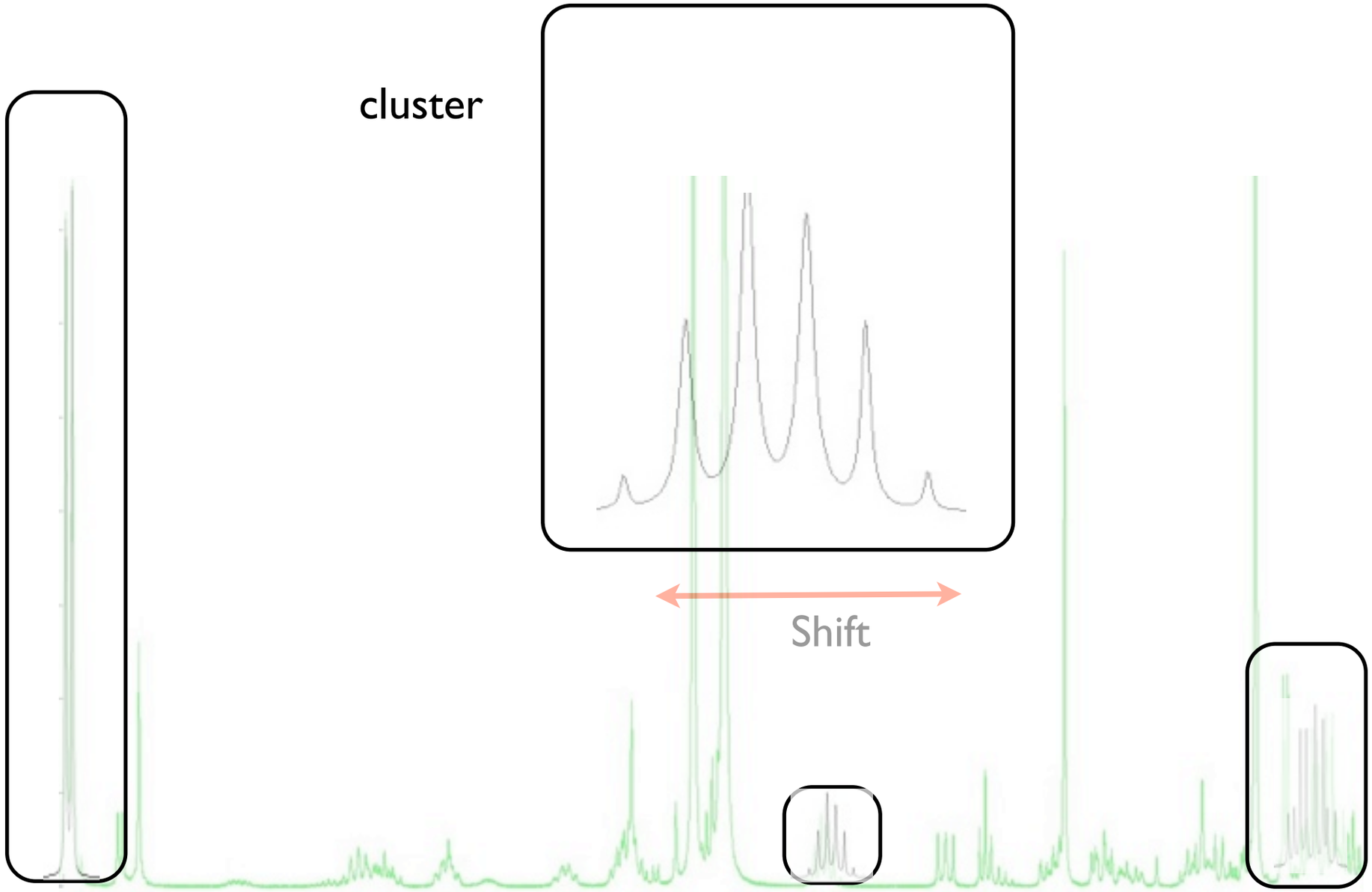




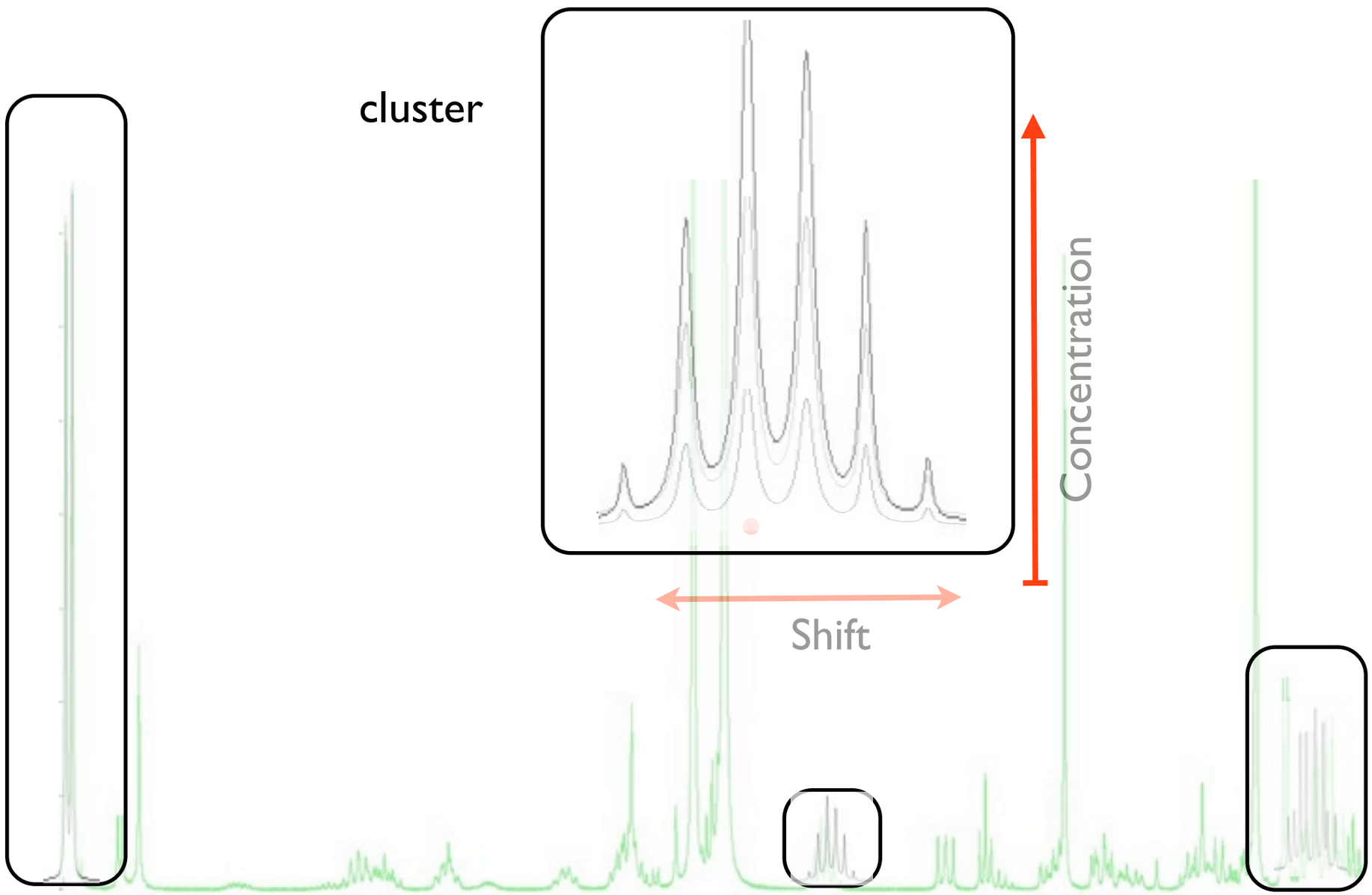
3-Hydroxyisobutyric acid



3-Hydroxyisobutyric acid



3-Hydroxyisobutyric acid



3-Hydroxyisobutyric acid

The Goal is:

Given The library of metabolite Signatures
Find Corresponding Concentration

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Optimization variables:

- Metabolite Concentrations
- Chemical Shifts

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Given The library of metabolite Signatures
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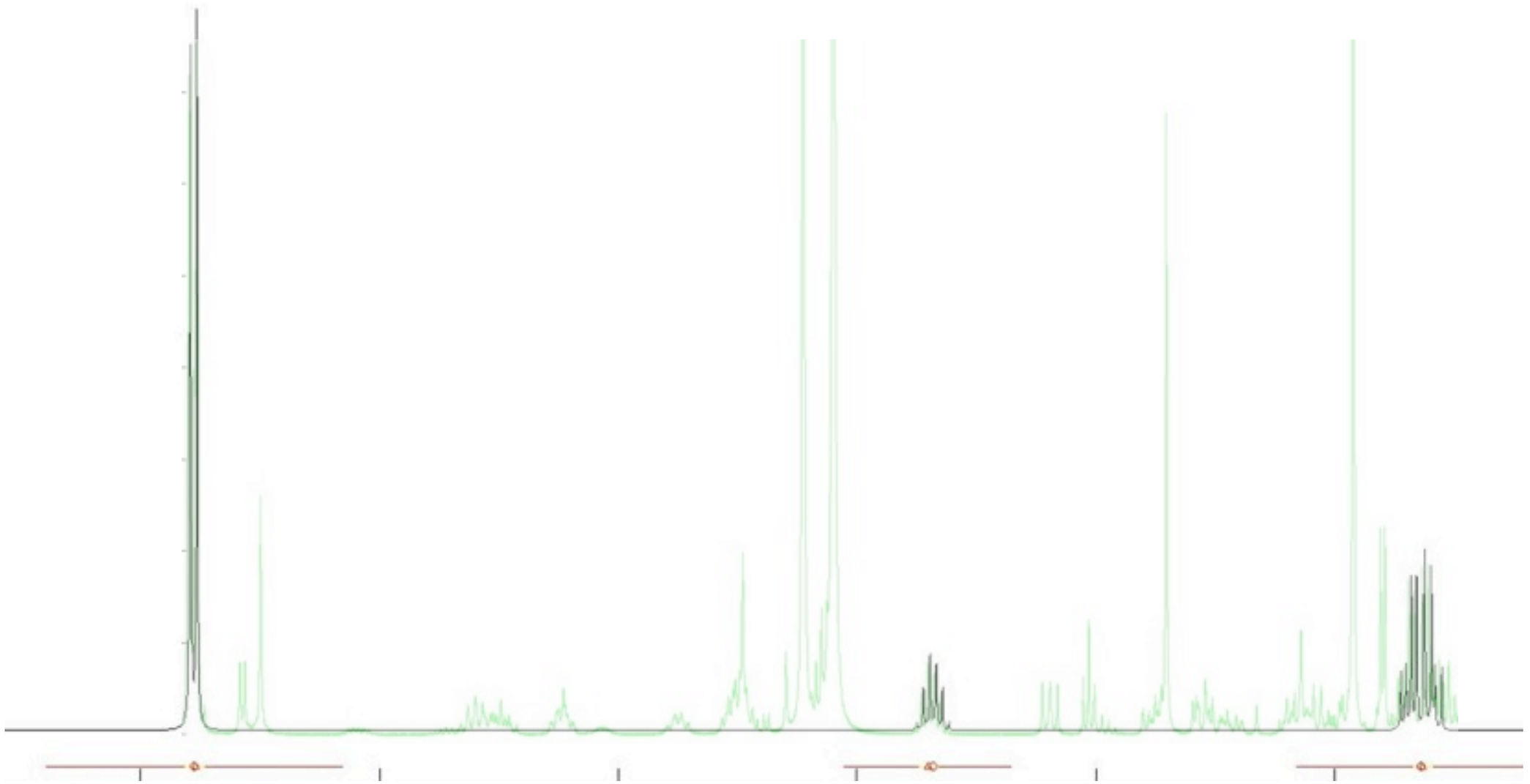
Optimization variables:

- Metabolite Concentrations
- Chemical Shifts

It's a difficult Problem

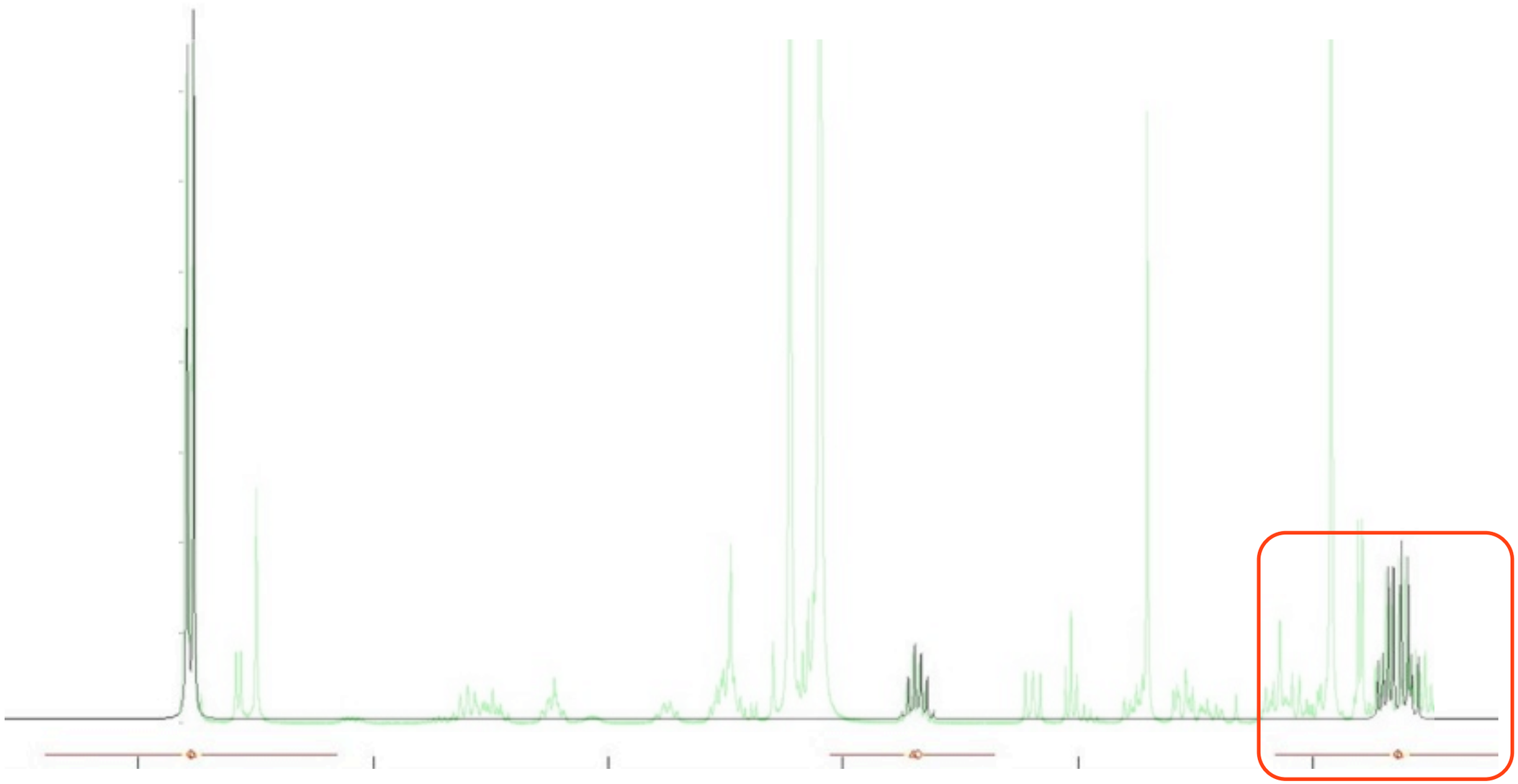
- Nonlinear in shift variables
- Involves hundreds of variables
- Loss is very expensive to evaluate (100K points)
- Non-Convex even around local optima
- Incomplete library results in over-fitting

Exploiting the structure



Exploiting the structure

Sub-problems are easier to solve but they share variables



Exploiting the structure

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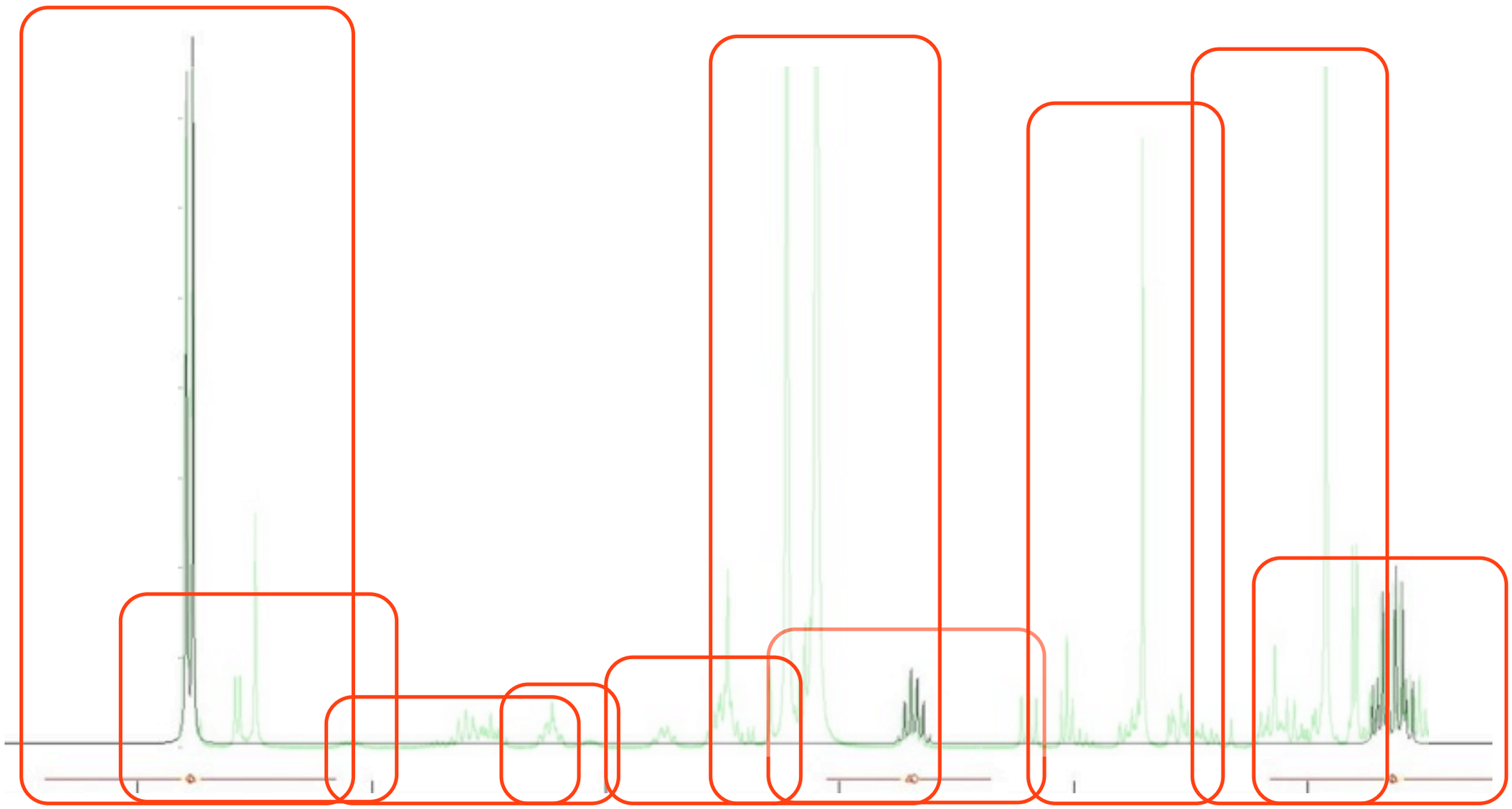
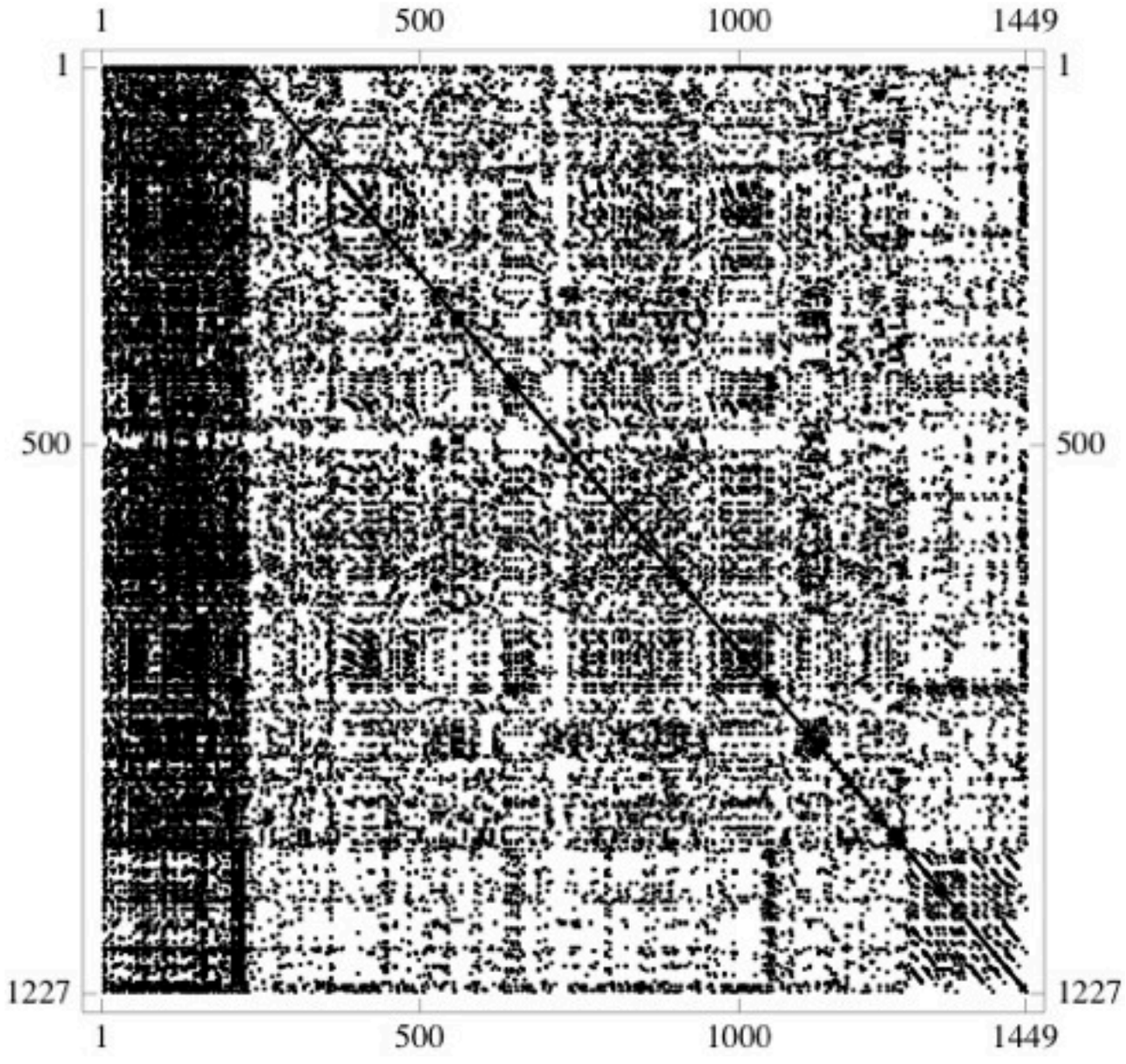


Coupling Matrix represents the structure



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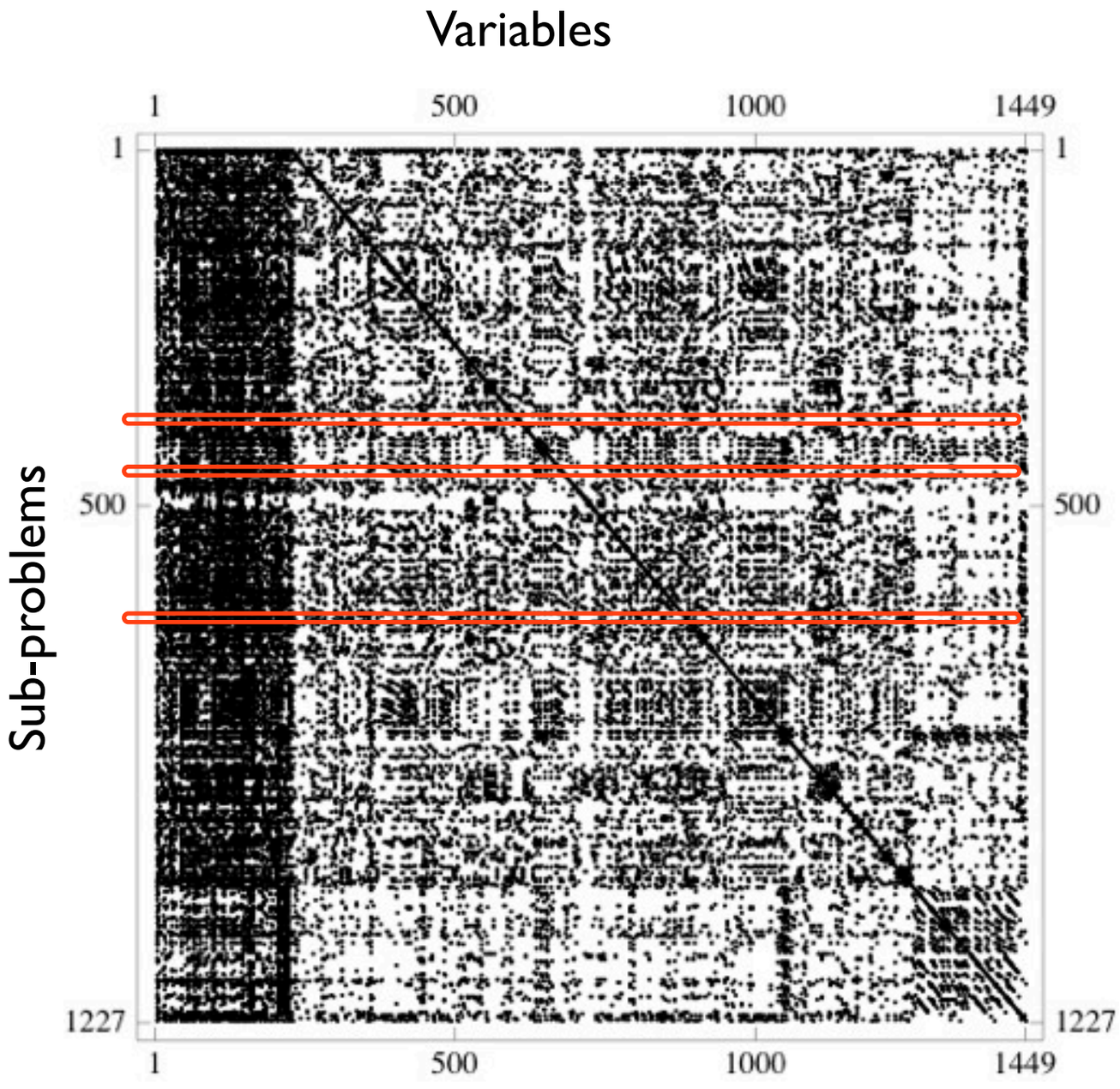
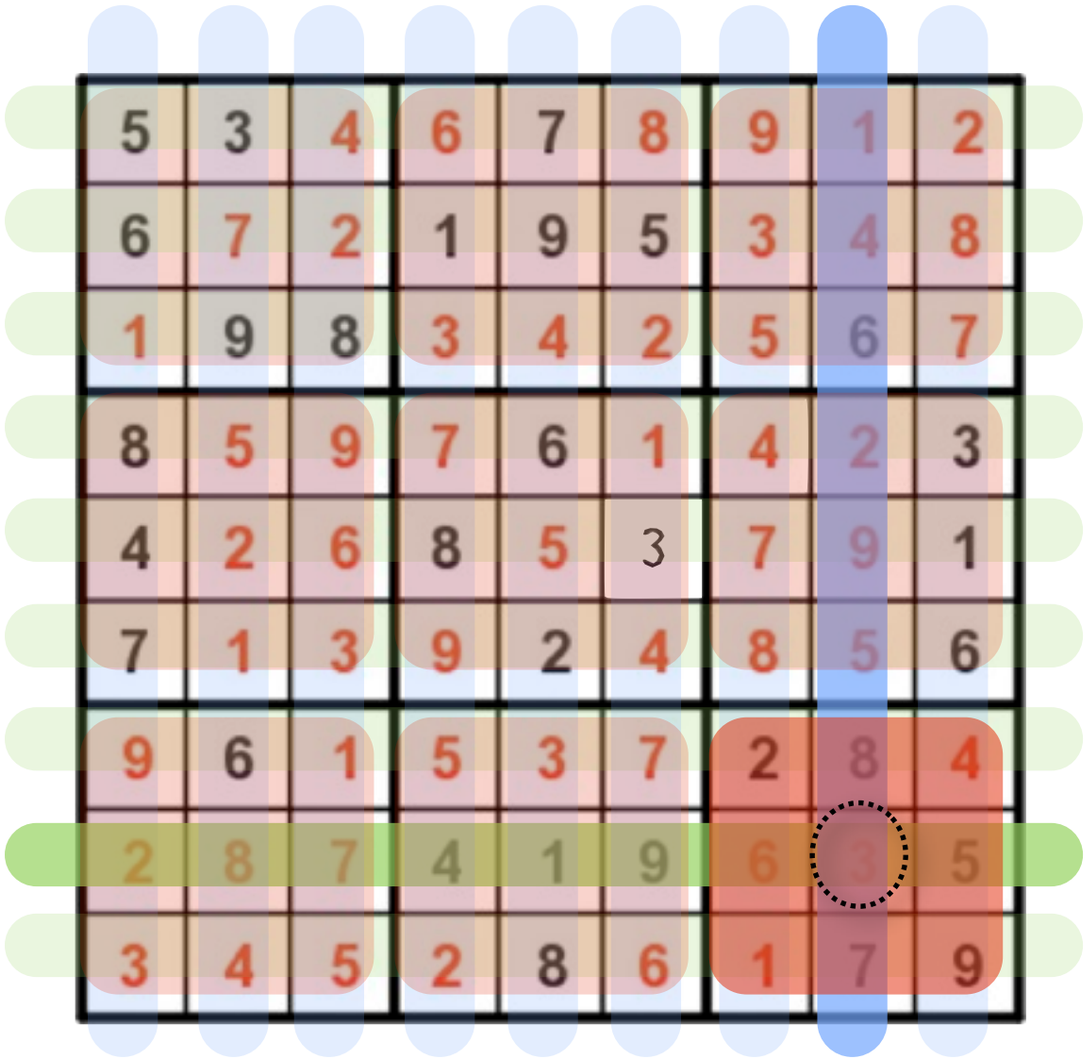
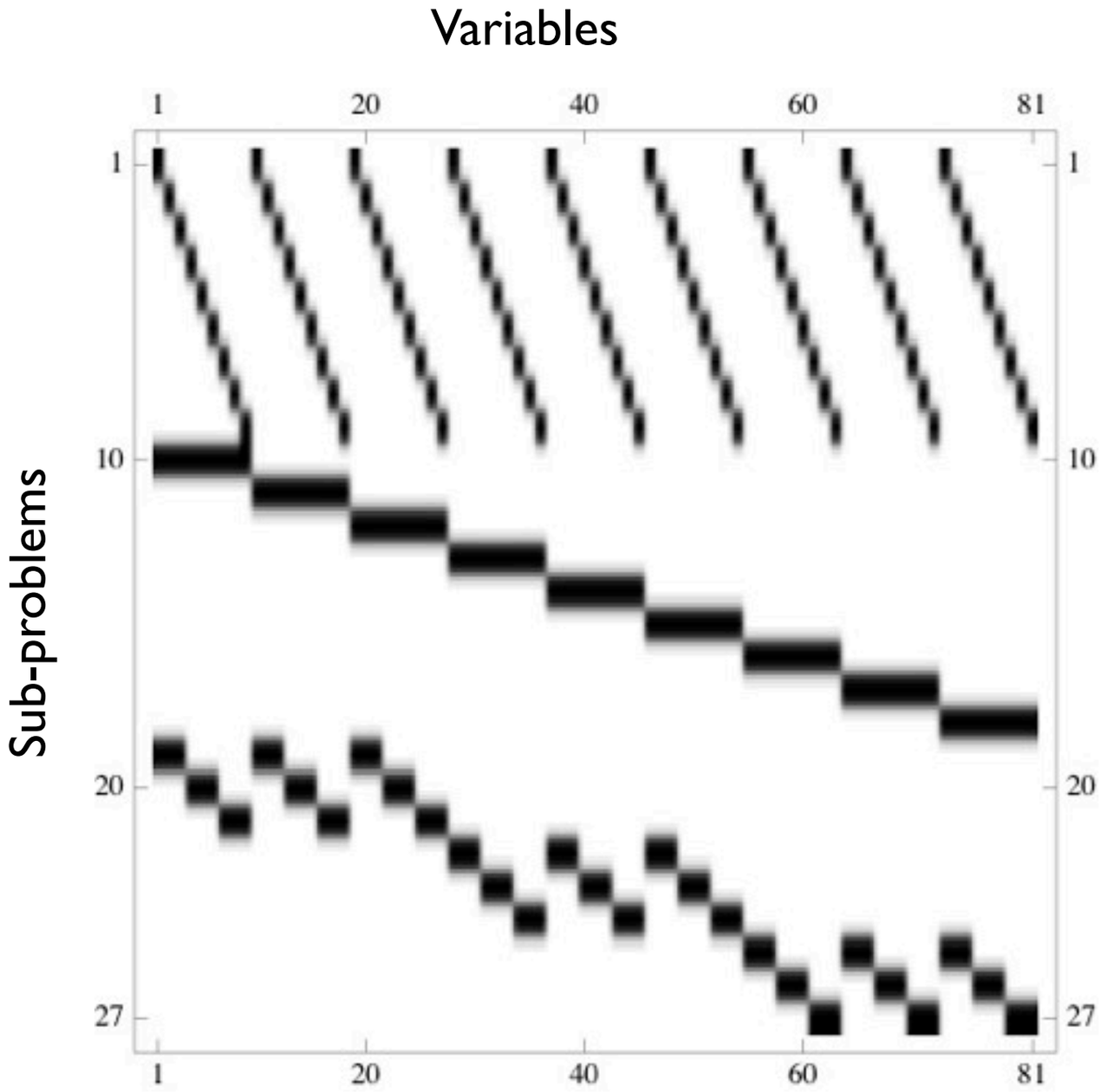


Coupling Matrix represents the structure

Sudoku Puzzle

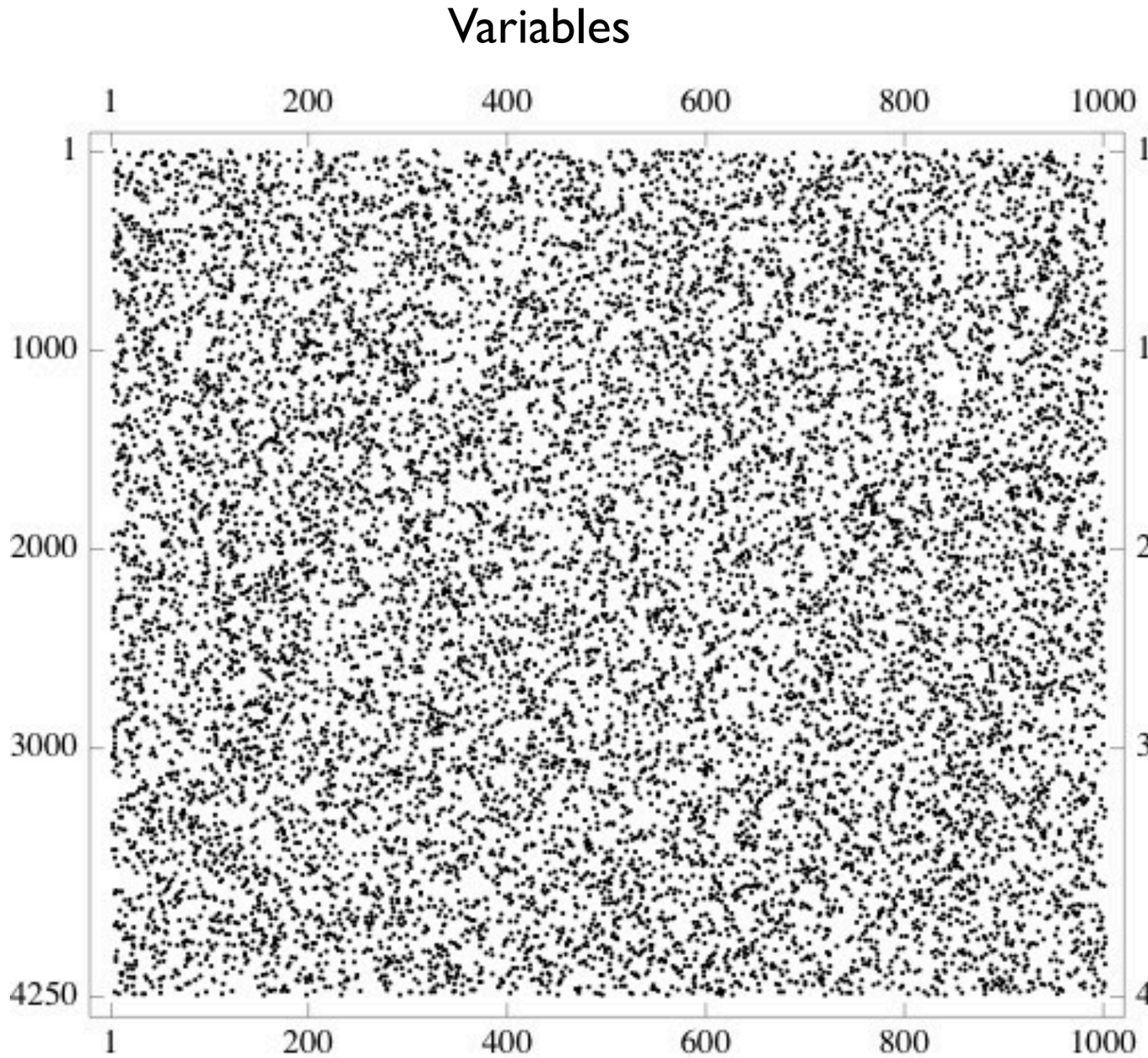




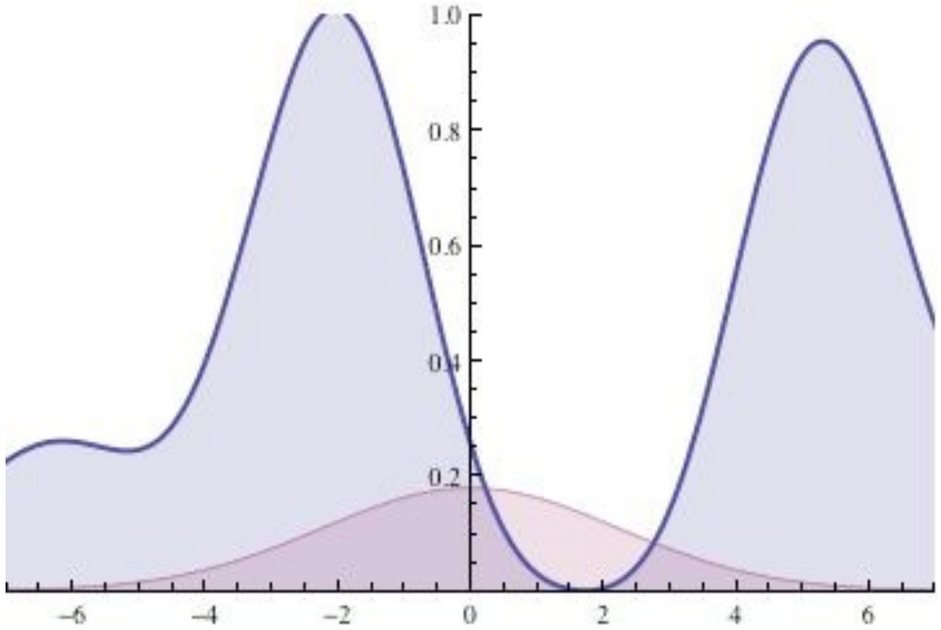
Coupling Matrix represents the structure

SATisfiability Problem

$$(A \vee B) \wedge (\neg B \vee C \vee \neg D) \wedge (D \vee \neg E)$$



Cross Entropy (CE) Method for Optimization

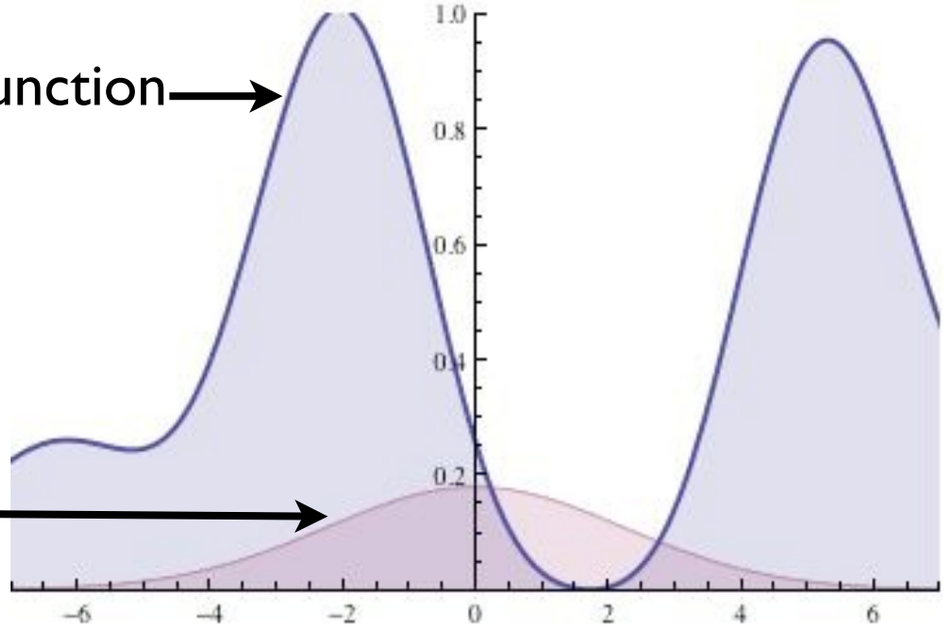


Cross Entropy (CE) Method for Optimization

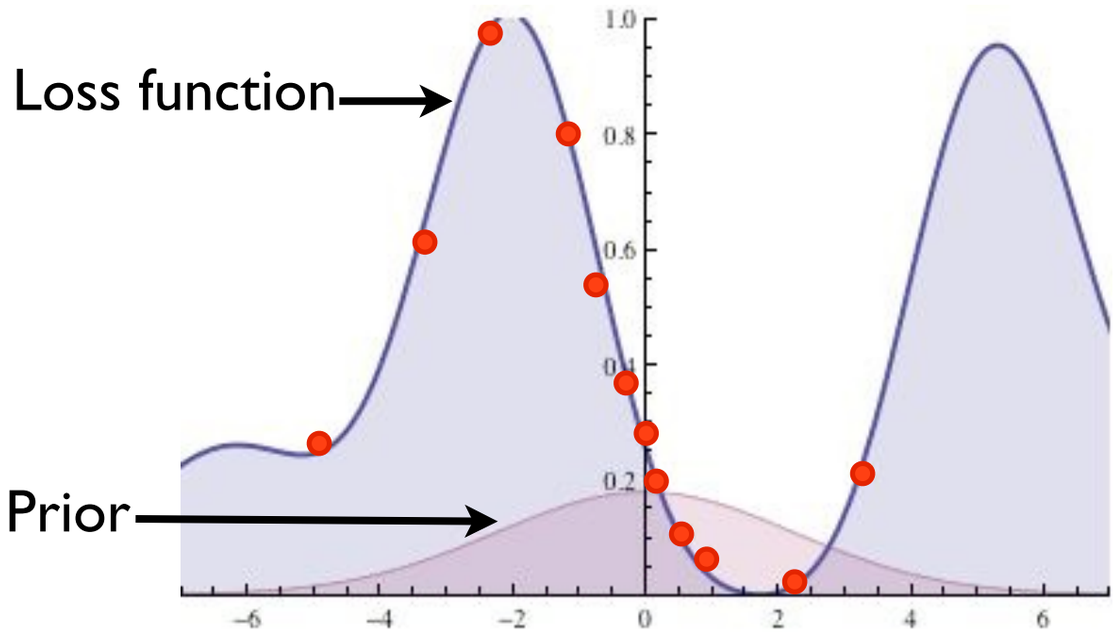
Start from a prior

Loss function

Prior



Cross Entropy (CE) Method for Optimization



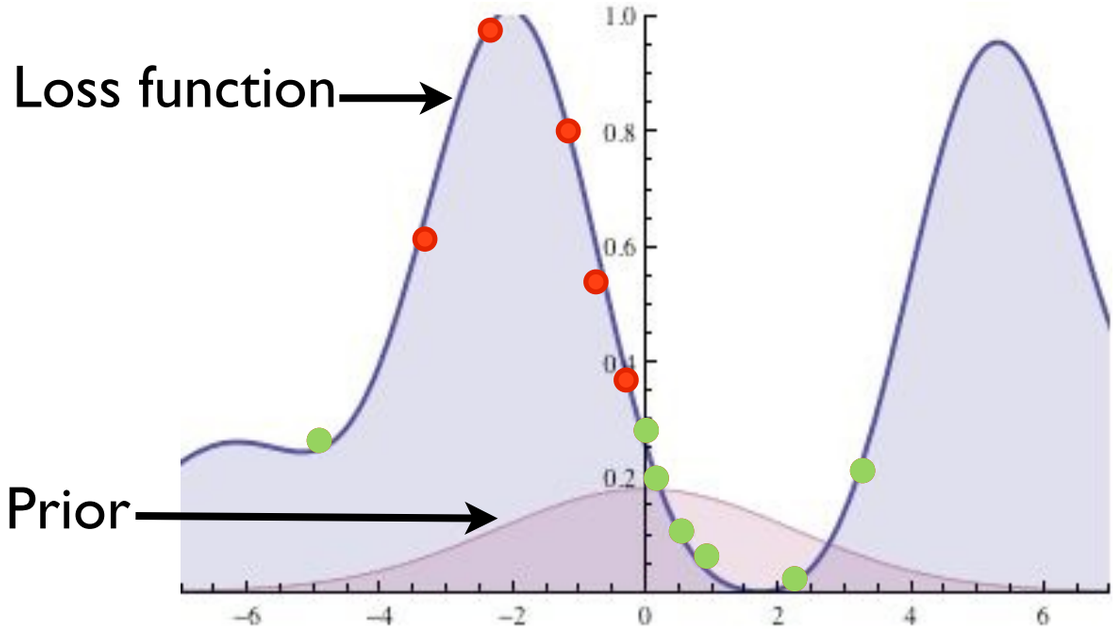
Start from a prior

Repeat until convergence

Take samples from current dist.

Calculate the loss for samples

Cross Entropy (CE) Method for Optimization



Start from a prior

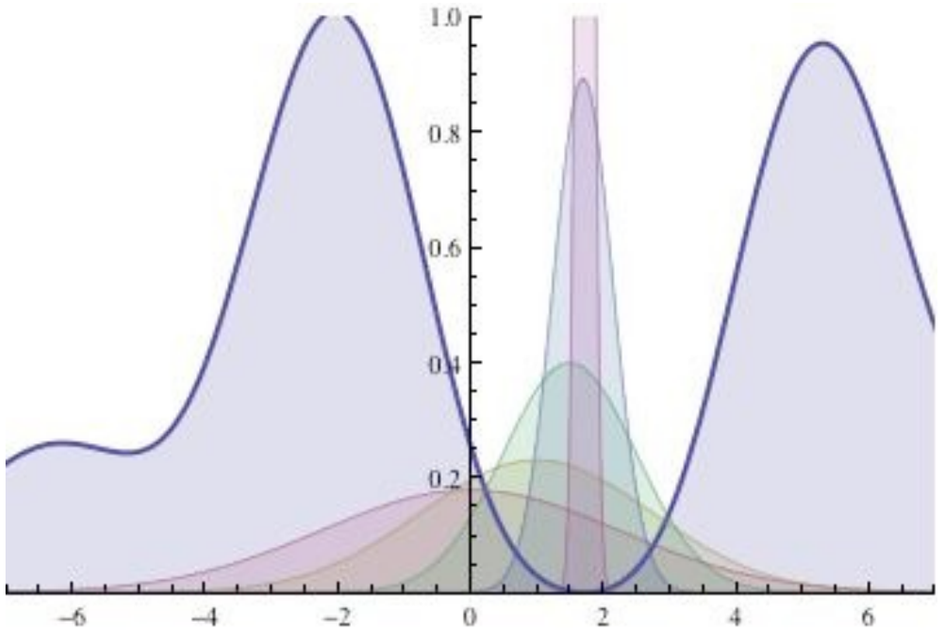
Repeat until convergence

Take samples from current dist.

Calculate the loss for samples

Select Elite samples

Cross Entropy (CE) Method for Optimization



Start from a prior

Repeat until convergence

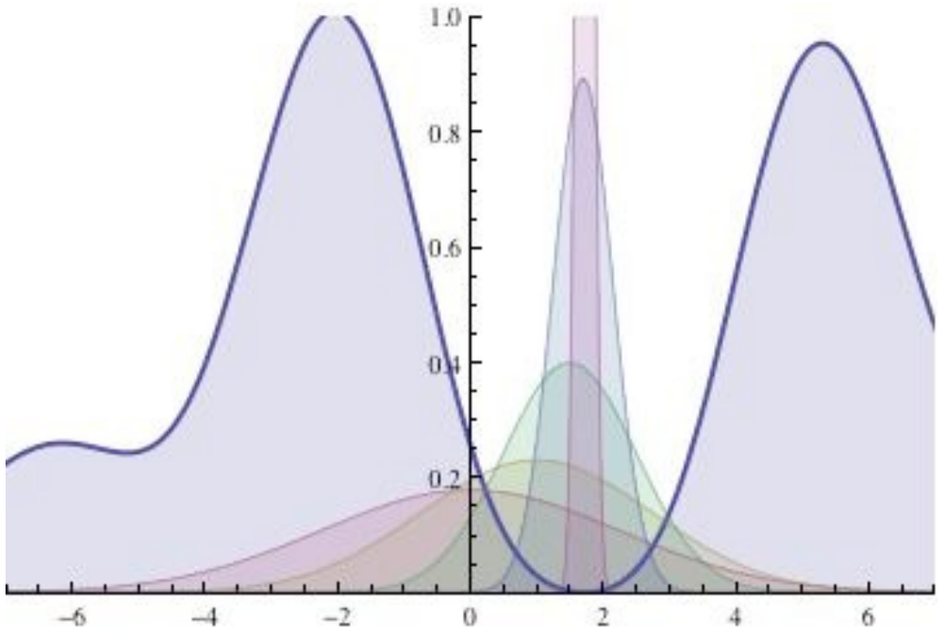
Take samples from current dist.

Calculate the loss for samples

Select Elite samples

Find maximum likelihood dist. for Elites

Cross Entropy (CE) Method for Optimization



Start from a prior

Repeat until convergence

Take samples from current dist.

Calculate the loss for samples

Select Elite samples

Find maximum likelihood dist. for Elites

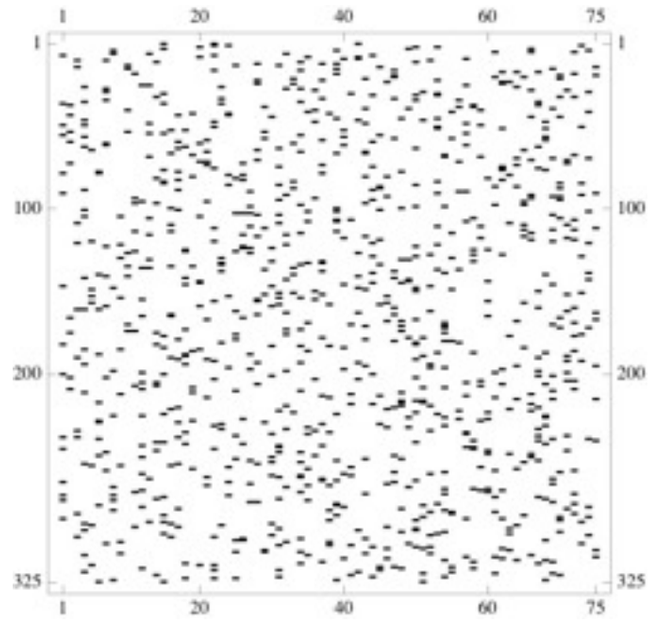
A subroutine to be used again

Variation of CE

that Exploits Decomposability (**CEED**)

Variables (i)

Sub-problems (k)



coupling matrix

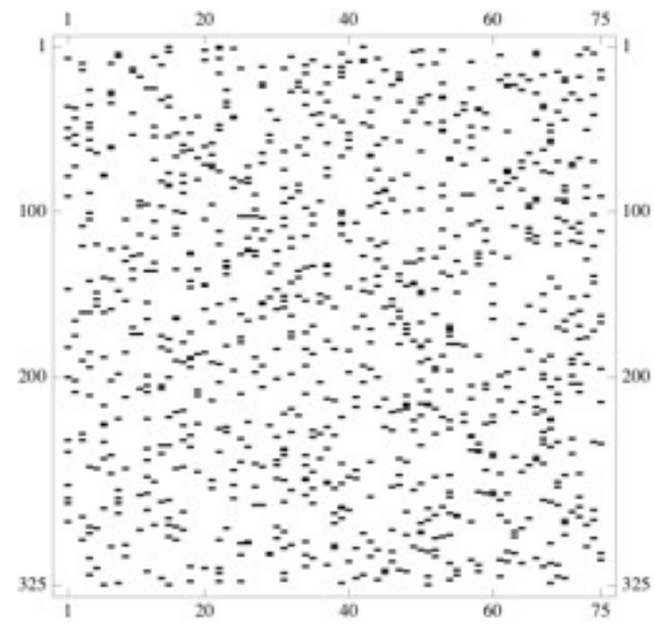
Variation of CE

that Exploits Decomposability (**CEED**)

Start from a prior

Repeat,

Until convergence



coupling matrix

Variation of CE

that Exploits Decomposability (**CEED**)

Start from a prior

Repeat,

For each sub-problem **k**

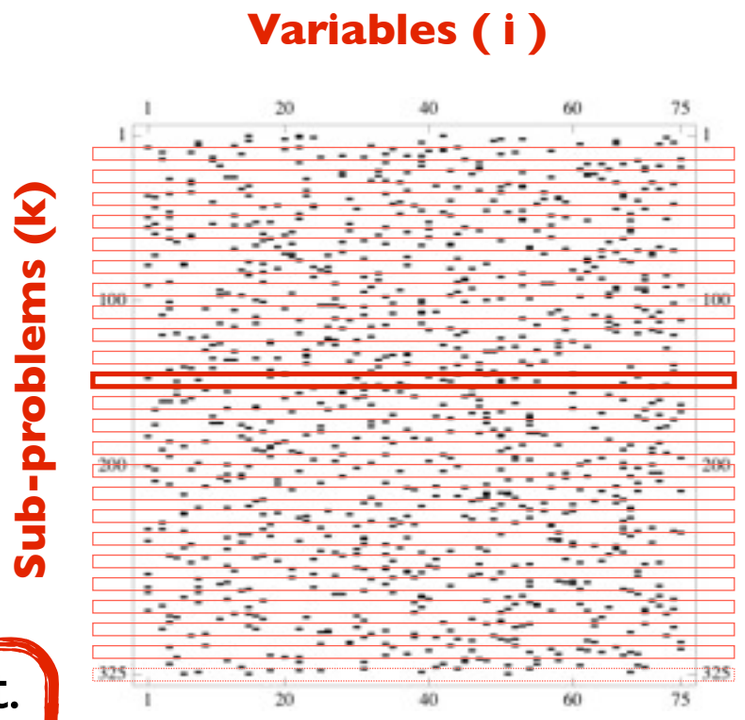
Draw samples from marginal of current dist.

Calculate the loss for samples

Select Elite samples

Find maximum likelihood dist. for Elites

Until convergence



coupling matrix

Variation of CE

that Exploits Decomposability (**CEED**)

Start from a prior

Repeat,

For each sub-problem **k**

Draw samples from marginal of current dist.

Calculate the loss for samples

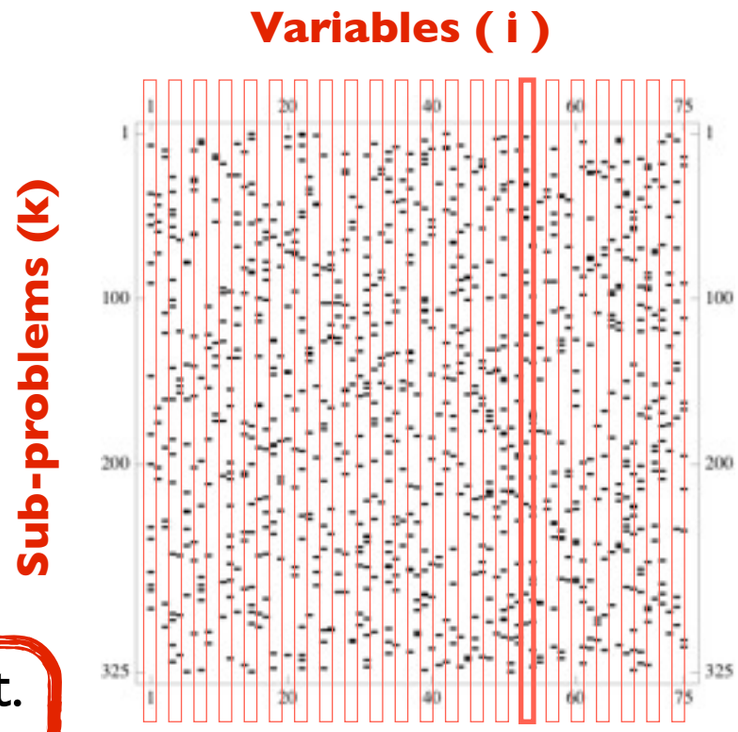
Select Elite samples

Find maximum likelihood dist. for Elites

For each variable **i**

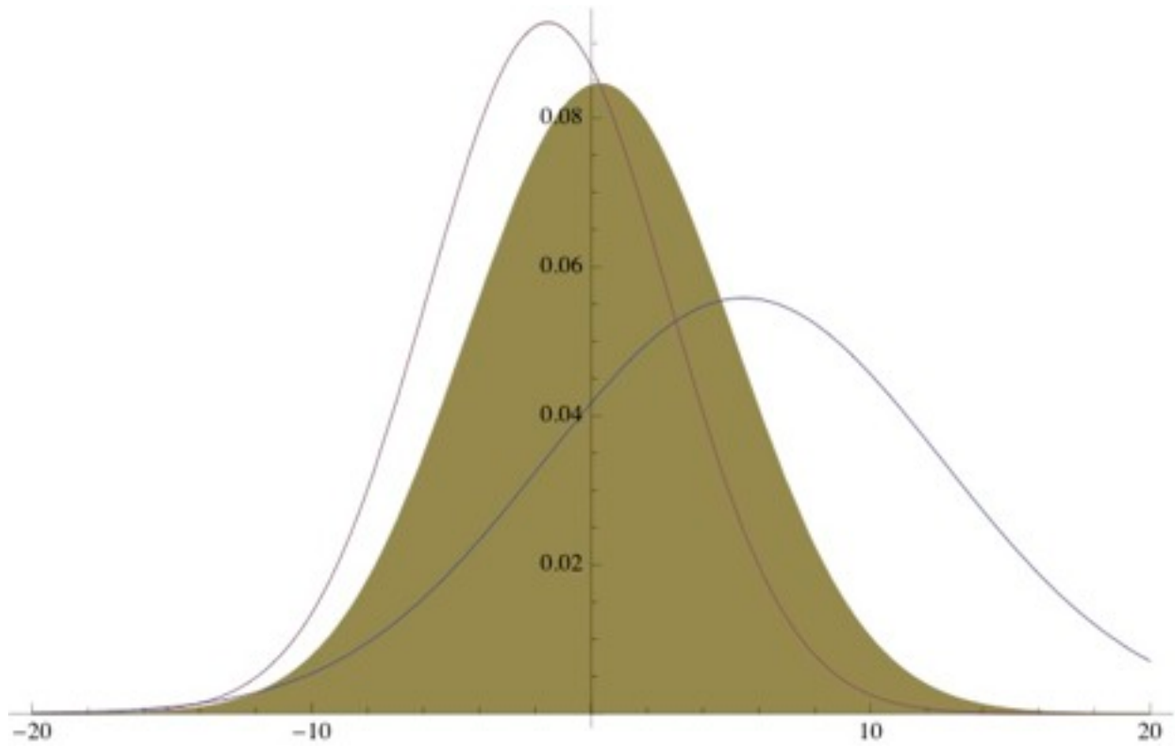
Combine dist's from related sub-problems

Until convergence



coupling matrix

How to Combine ML Distributions



How to Combine ML Distributions

We use linear combination by Fisher Information

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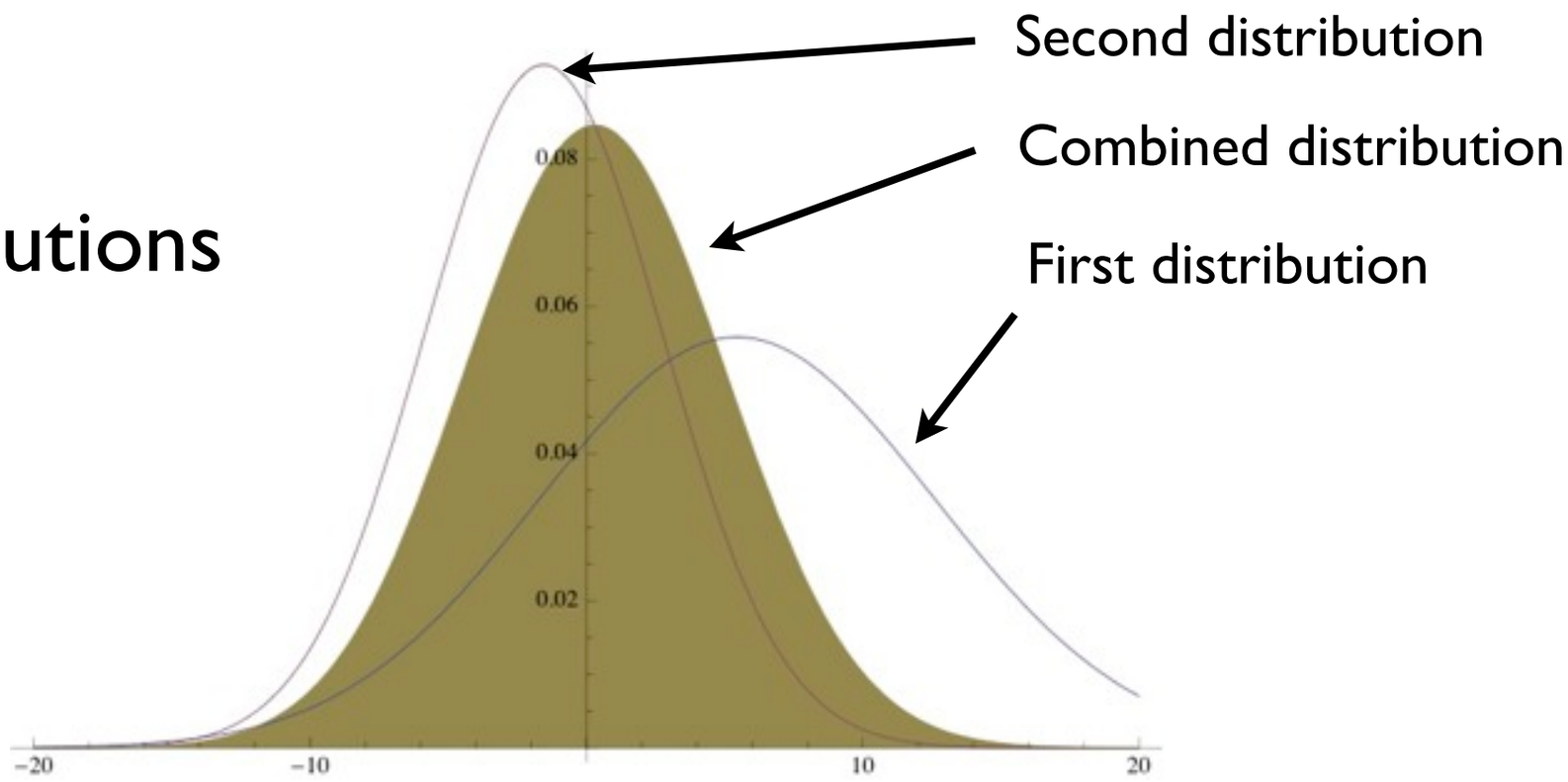
Gaussian Distributions



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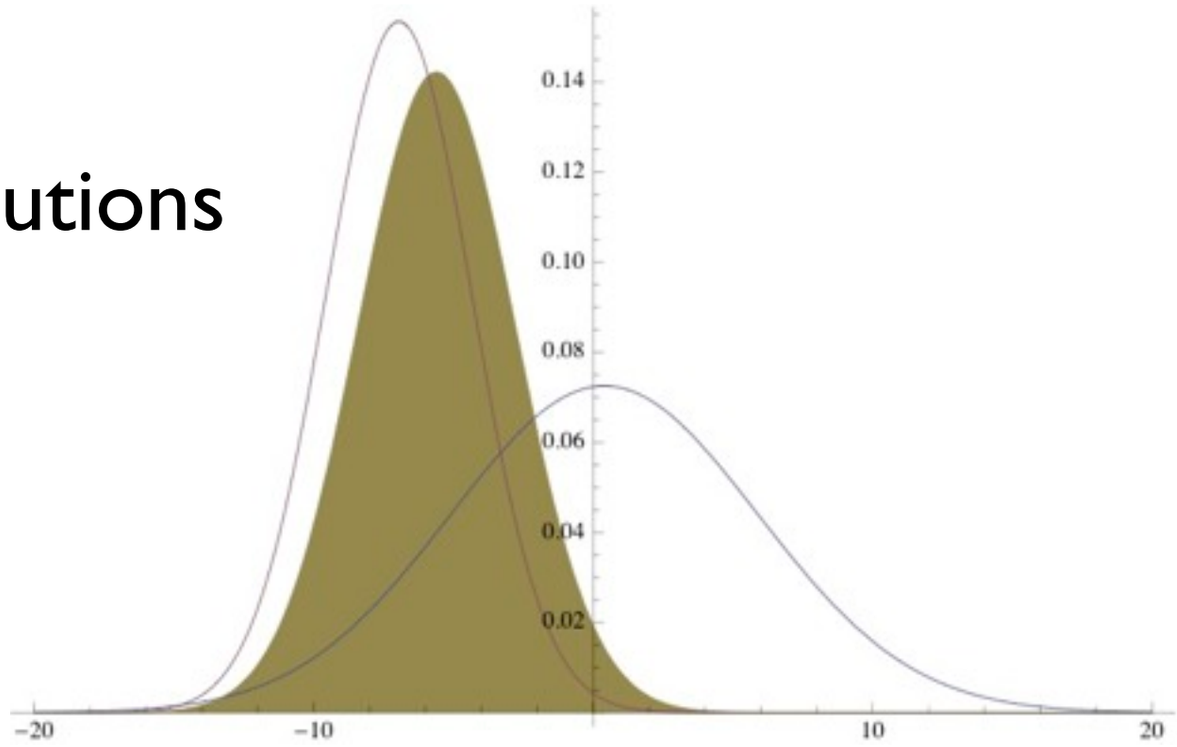

How to Combine ML Distributions

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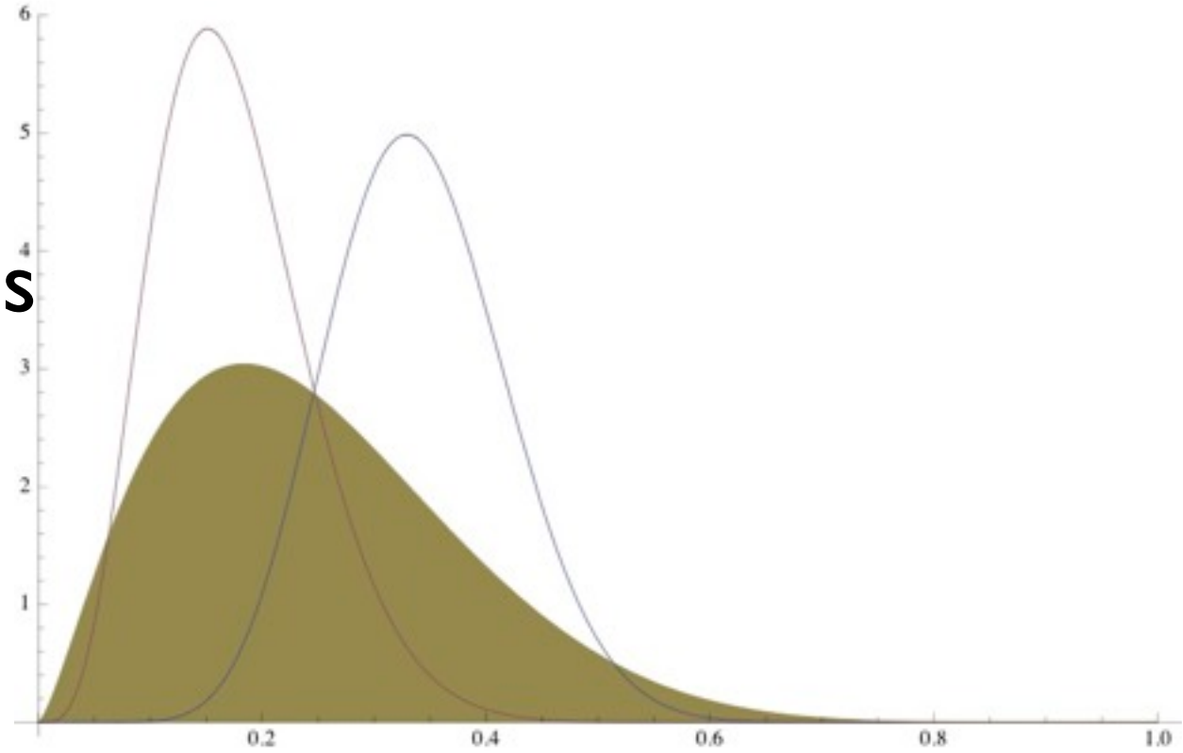
Beta Distributions



How to Combine ML Distributions

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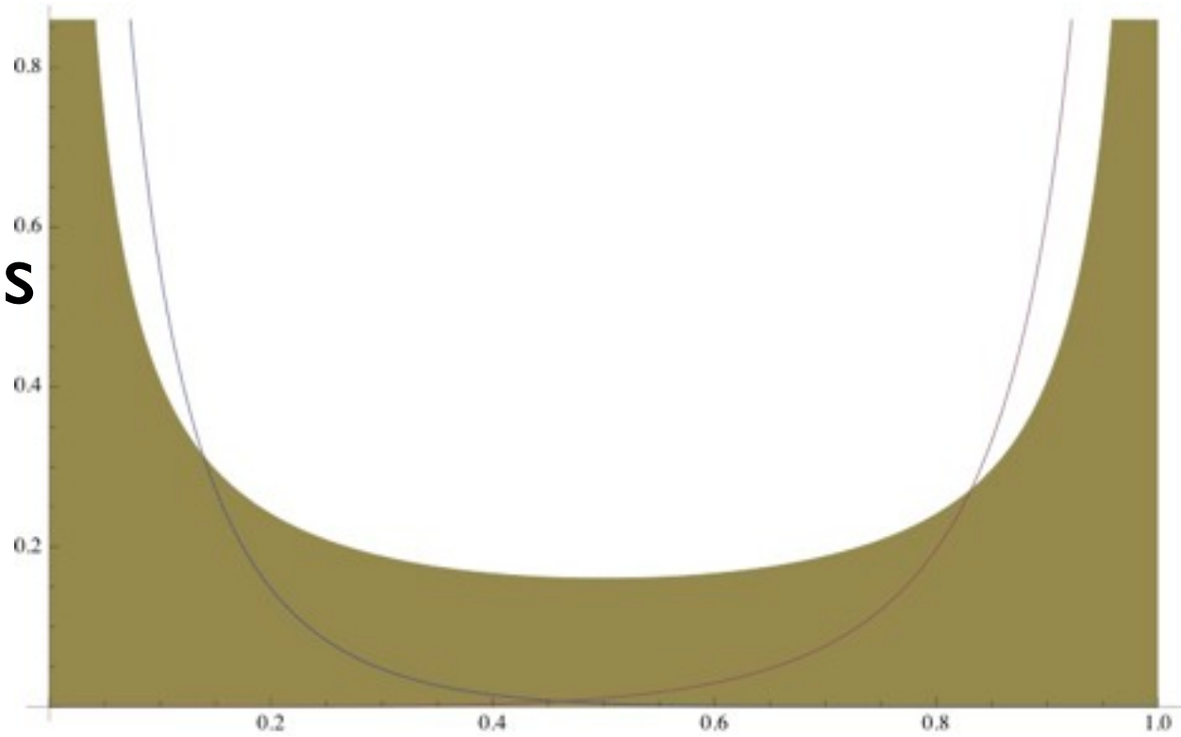

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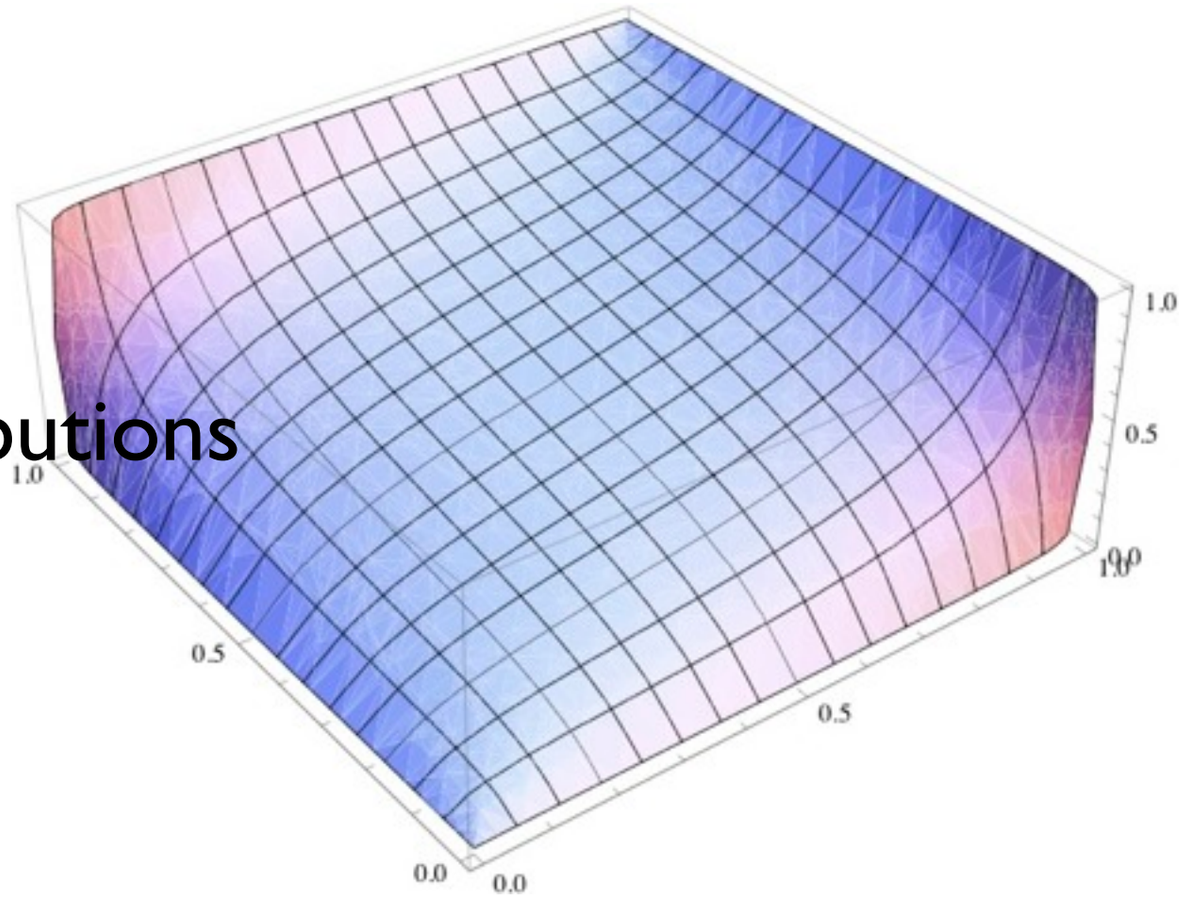
Bernoulli Distributions



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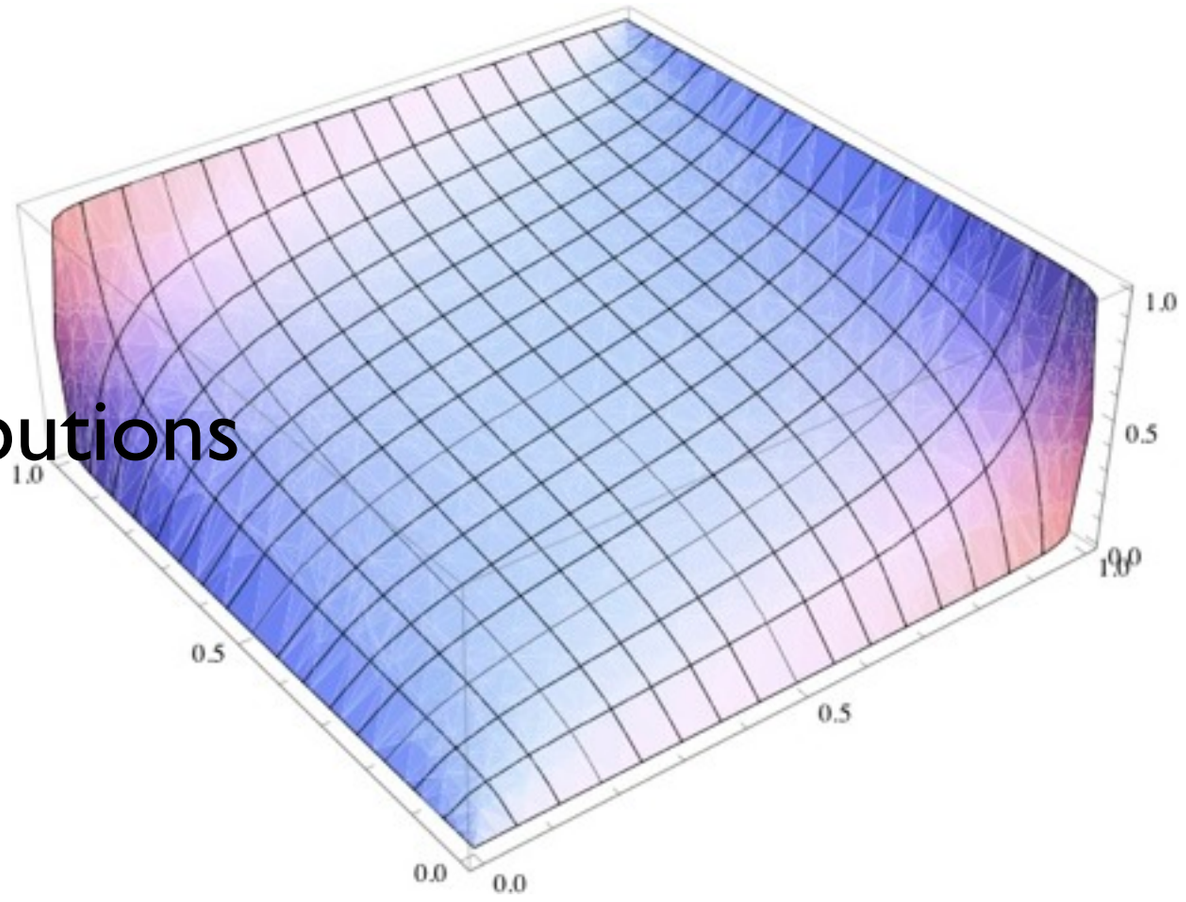
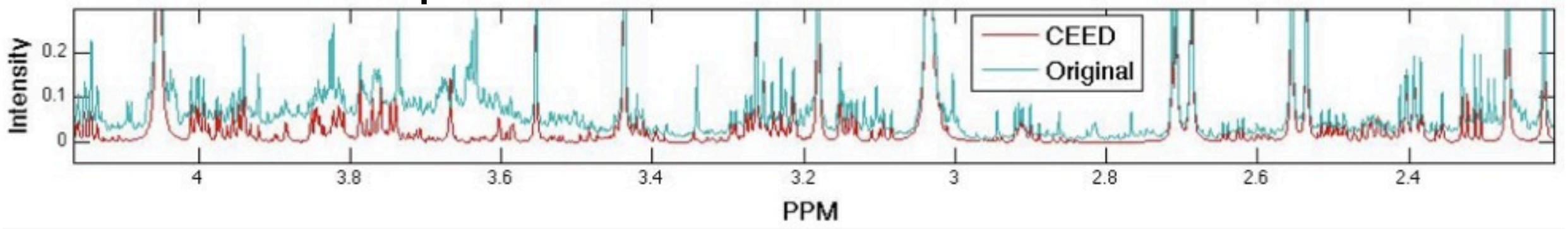
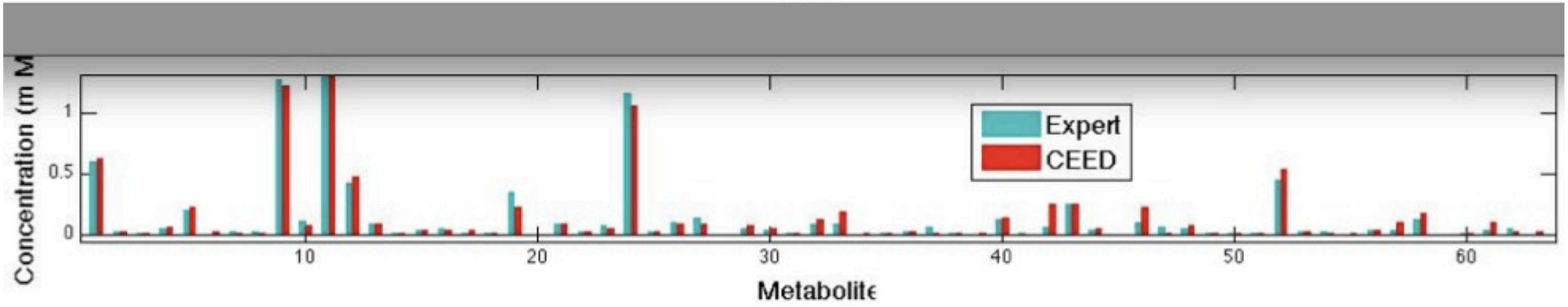


A Tweak for NMR problem

Experimental Results

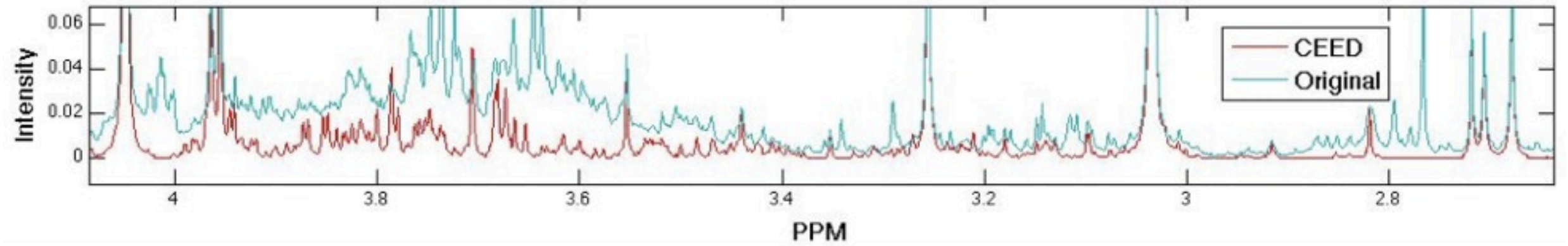
800 MHz Spectra

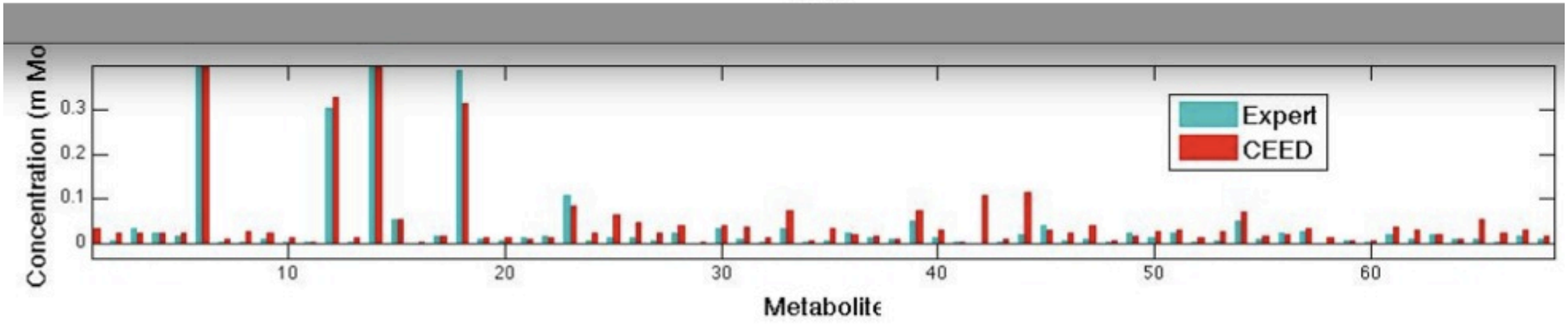




Experimental Results

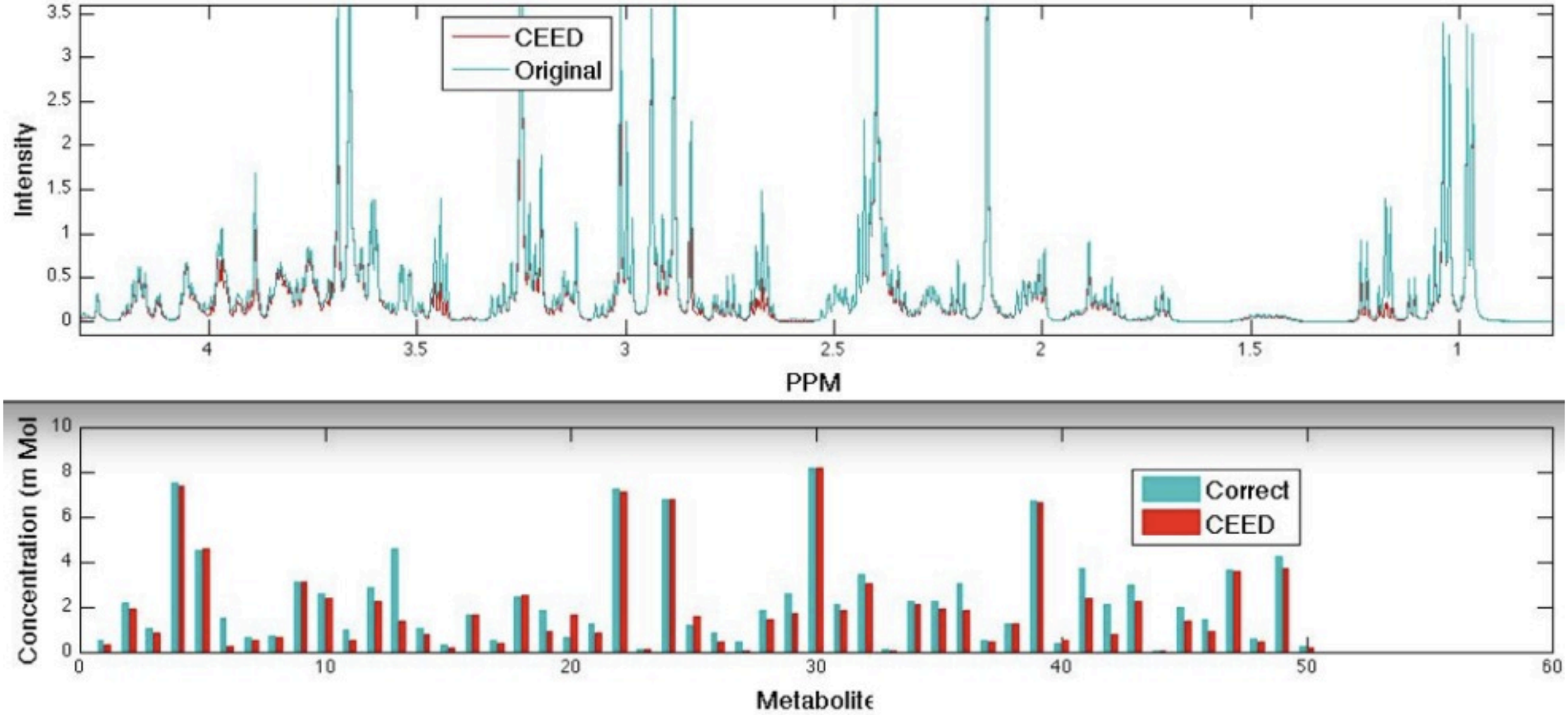
500 MHz Spectra





Experimental Results

Simulated Spectra



Experimental Results

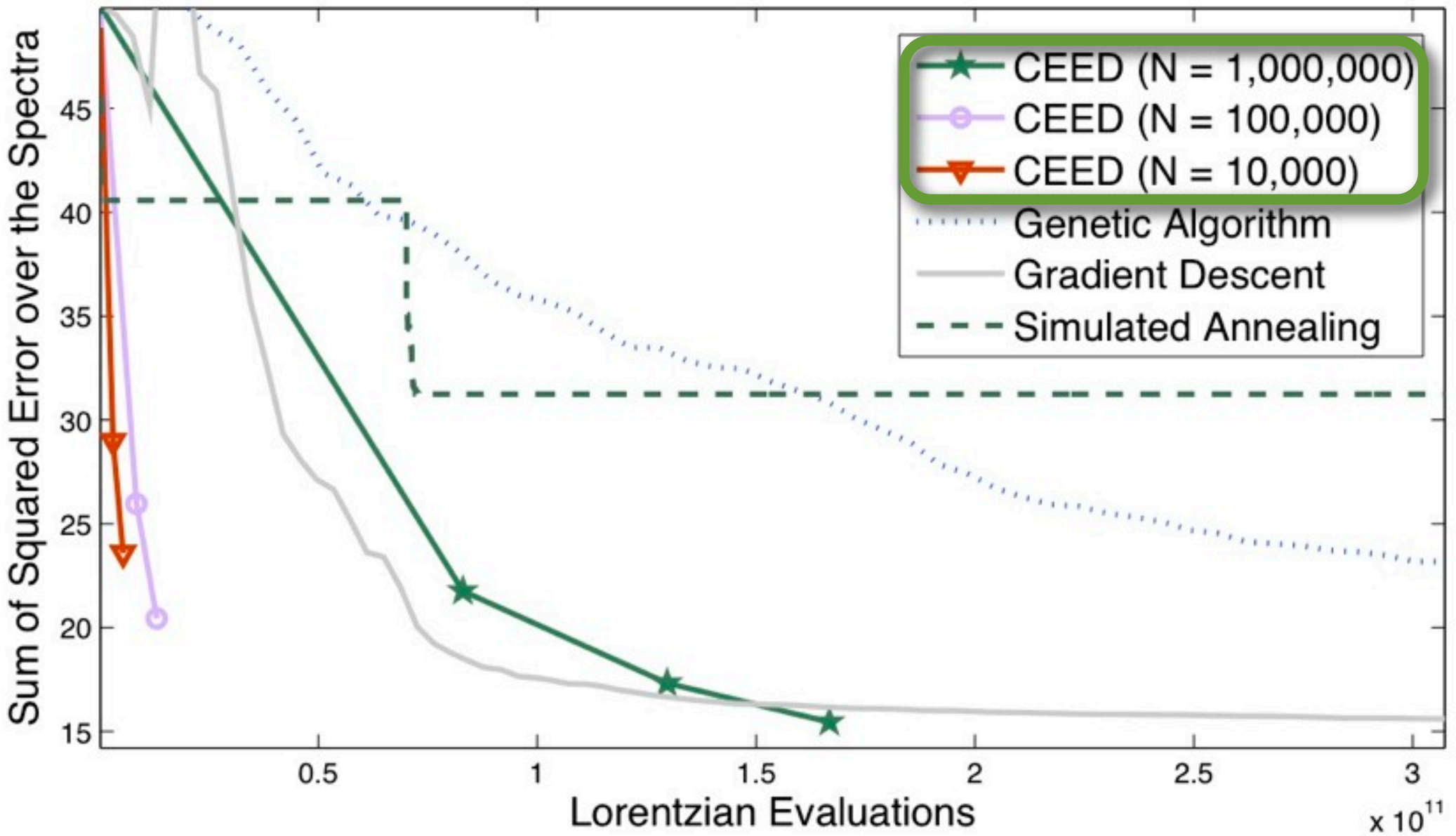
Comparison with ChenomX Inc. automated fitting software

Quantification Task

Metabolite Detection Task
Threshold: .02 mMol

Alg	Avg. Relative Error	Precision	Recall	F-measure
Us	**.39** ± .05	**.83** ± .08	.93 ± .06	**.87** ± .06
Them	.76 ± .05	.68 ± .13	.97 ± .03	.79 ± .10

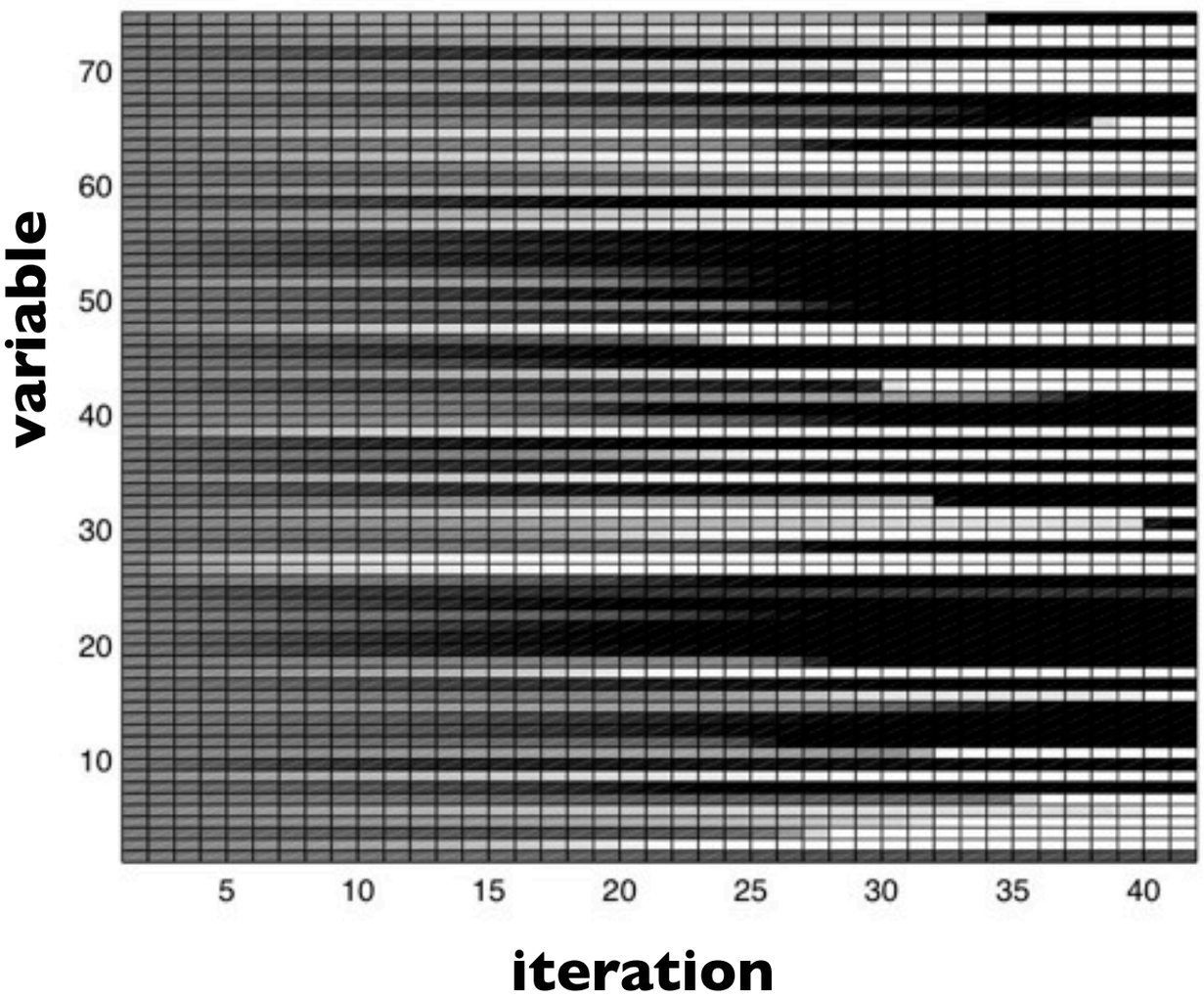
Experimental Results



MaxSAT

Analytically update of dist's is possible.

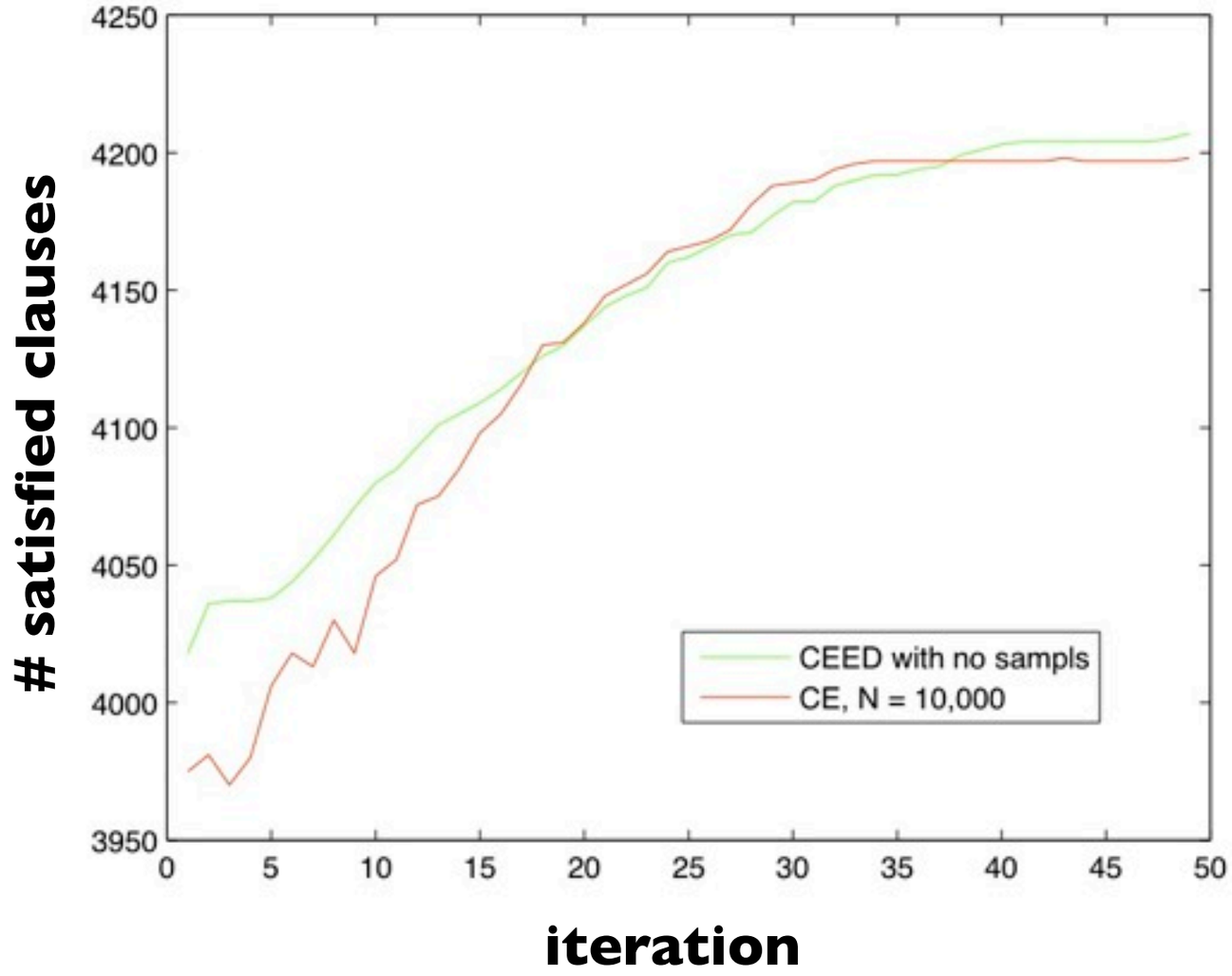
Convergence of distributions to the correct assignment



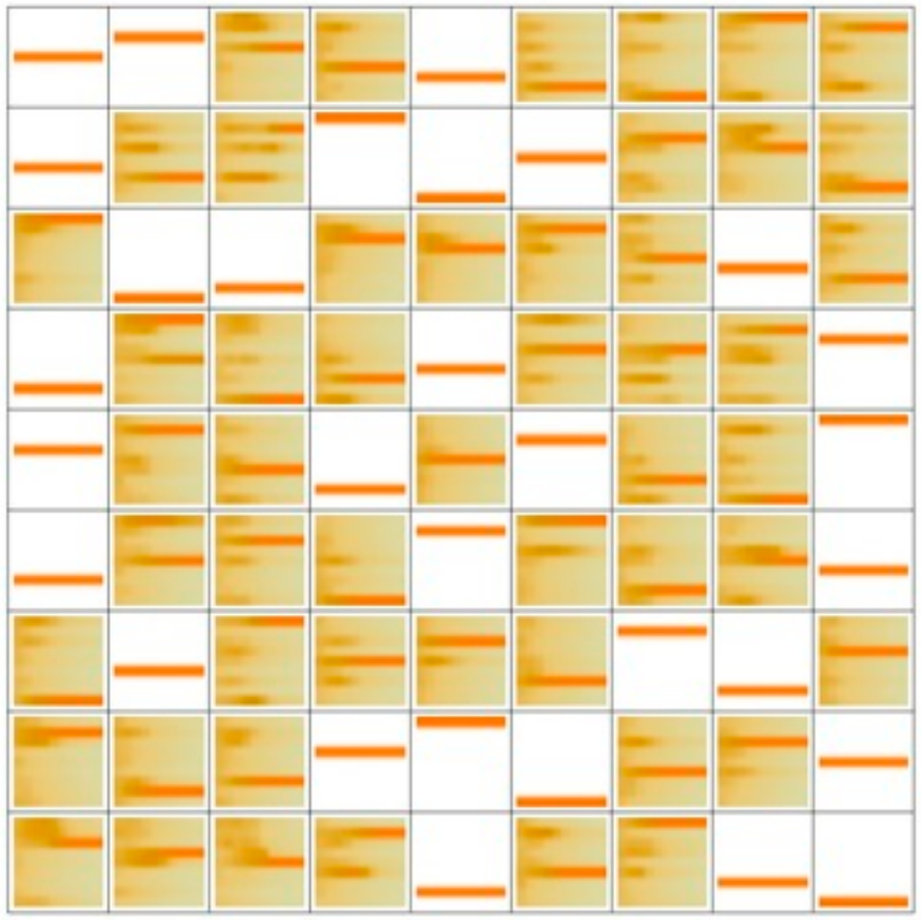
MaxSAT

Convergence of CE and CEED

#clauses 4250
#vars 1000



Sudoku



5	3	4	6	7	8	9	1	2
6	7	2	1	9	5	3	4	8
1	9	8	3	4	2	5	6	7
8	5	9	7	6	1	4	2	3
4	2	6	8	5	3	7	9	1
7	1	3	9	2	4	8	5	6
9	6	1	5	3	7	2	8	4
2	8	7	4	1	9	6	3	5
3	4	5	2	8	6	1	7	9

solution in red

Using categorical distribution

Our Alg. is **5** times faster than **CE**

Conclusion

Our method successfully exploits partial decomposability in targeted profiling of NMR spectra
as well as some combinatorial problems (i.e. SAT & Sudoku)

Maximum likelihood estimates could be linearly combined based on their certainty using their Fisher Information.

Thank you!

Computing Science Dept. University of Alberta

Alberta Ingenuity Centre for Machine Learning