

nmrfam's Home

BMRB_metabolites

GISSMO_v2

Maybridge_RO3_library

GISSMO (Guided Ideographic Spin System Model Optimization) - v. 2.0

File Tools Help

Get integral Draw full experimental spectrum Reset integral

Show 2D figure

Water region (PPM):
4.6 5
☒ delete water region

DSS region (PPM):
-0.1 0.1
☒ delete DSS region

ROI region: -1 to 12 get ROI from exp. spectrum

database entries:

bmse000006(ATP)-Initialvalues

bmse000025(D-(+)-Threo-isocitricacid)-Initialvalues

bmse000026(D-(+)-Xylose)-Initialvalues

bmse000027(D-Xylose)-Initialvalues

bmse000028(L-Alanine)-Initialvalues

bmse000029(L-Arginine)-Initialvalues

bmse000030(L-Asparagine)-Initialvalues

bmse000031(L-Asparticacid)-Initialvalues

bmse000032(L-Citrulline)-Initialvalues

bmse000033(L-Cystathionine)-Initialvalues

bmse000034(L-Cysteine)-Initialvalues

bmse000035(L-Cystine)-Initialvalues

bmse000036(L-(-)-Fucose)-Initialvalues

bmse000037(L-Glutamicacid)-Initialvalues

bmse000038(L-Glutamine)-Initialvalues

bmse000039(L-Histidine)-Initialvalues

bmse000040(L-Homoserine)-Initialvalues

bmse000041(L-Isoleucine)-Initialvalues

bmse000042(L-Leucine)-Initialvalues

bmse000043(L-Lysine)-Initialvalues

bmse000044(L-Methionine)-Initialvalues

Load

Num points: 2^14
☒ Same as exp. data

Line width: 0.3

Field strength: 500
☒ Same as exp. data

Lorentzian coeff: 0.8

Gaussian coeff: 0.2

☒ Save automatically

simulation info

Process Optimization Group and optimize

Copy selected cells Paste cells Swap two cells

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-0.1 0.1
☒ delete DSS region

ROI region: -1 to 12 get ROI from exp. spectrum

☒ Save automatically simulation info

database entries: bmse000030(L-Asparagine)-Initialvalues Load

compound_name

	1	2
1		
2		
3		
4		

Process Optimization Group and optimize
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☒ delete DSS region

ROI region: -1 to 12 get ROI from exp. spectrum

database entries: bmse000030(L-Asparagine)-Initialvalues Load

Same compounds in the library: bmse000912 Copy matrix

L-Asparagine(merged)

	10	11	12
10	2.7811	-12.4000	6.3700
11	-12.4000	2.8911	6.3700
12	6.3700	6.3700	3.9620

Process Optimization Group and optimize

Copy selected cells Paste cells Swap two cells

☒ Save automatically simulation info

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Show 2D figure

Water region (PPM):
4.6 5
☒ delete water region

DSS region (PPM):
-0.1 0.1
☒ delete DSS region

Get integral

Draw full experimental spectrum

Reset integral

Num points: 32768
☒ Same as exp. data

Line width: 1

Field strength: 499.84
☒ Same as exp. data

Lorentzian coeff: 0.8

Gaussian coeff: 0.2

database entries: bmse000030(L-Asparagine)-Initialvalues Load

Same compounds in the library: bmse000912 Copy matrix

L-Asparagine(merged)

	10	11	12
10	2.7811	-12.4000	6.3700
11	-12.4000	2.8911	6.3700
12	6.3700	6.3700	3.9620

ROI region: -1 to 12 get ROI from exp. spectrum

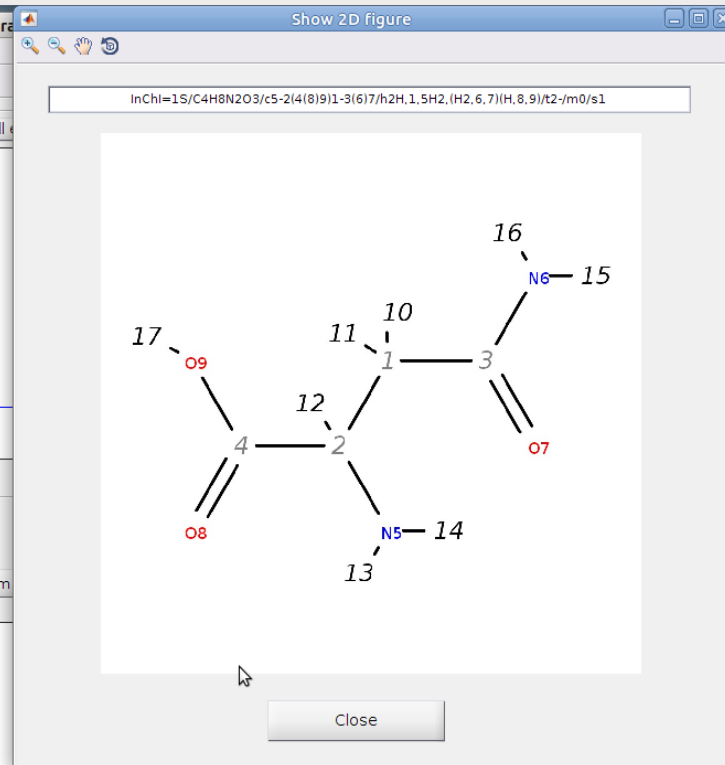
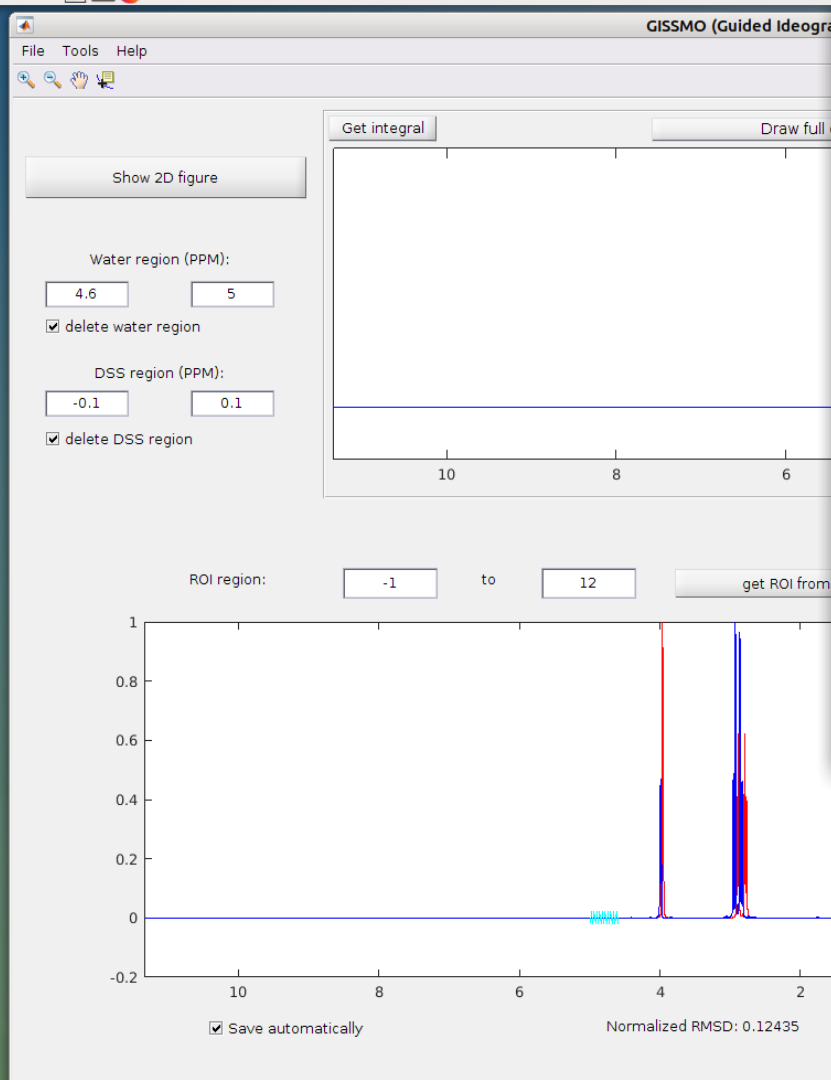
Normalized RMSD: 0.12435

☒ Save automatically

Process Optimization Group and optimize

Copy selected cells Paste cells Swap two cells

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GISSMO_v2
Maybridge_RO3_library



32768

ne as exp. data

1

499.84

me as exp. data

eff: 0.8

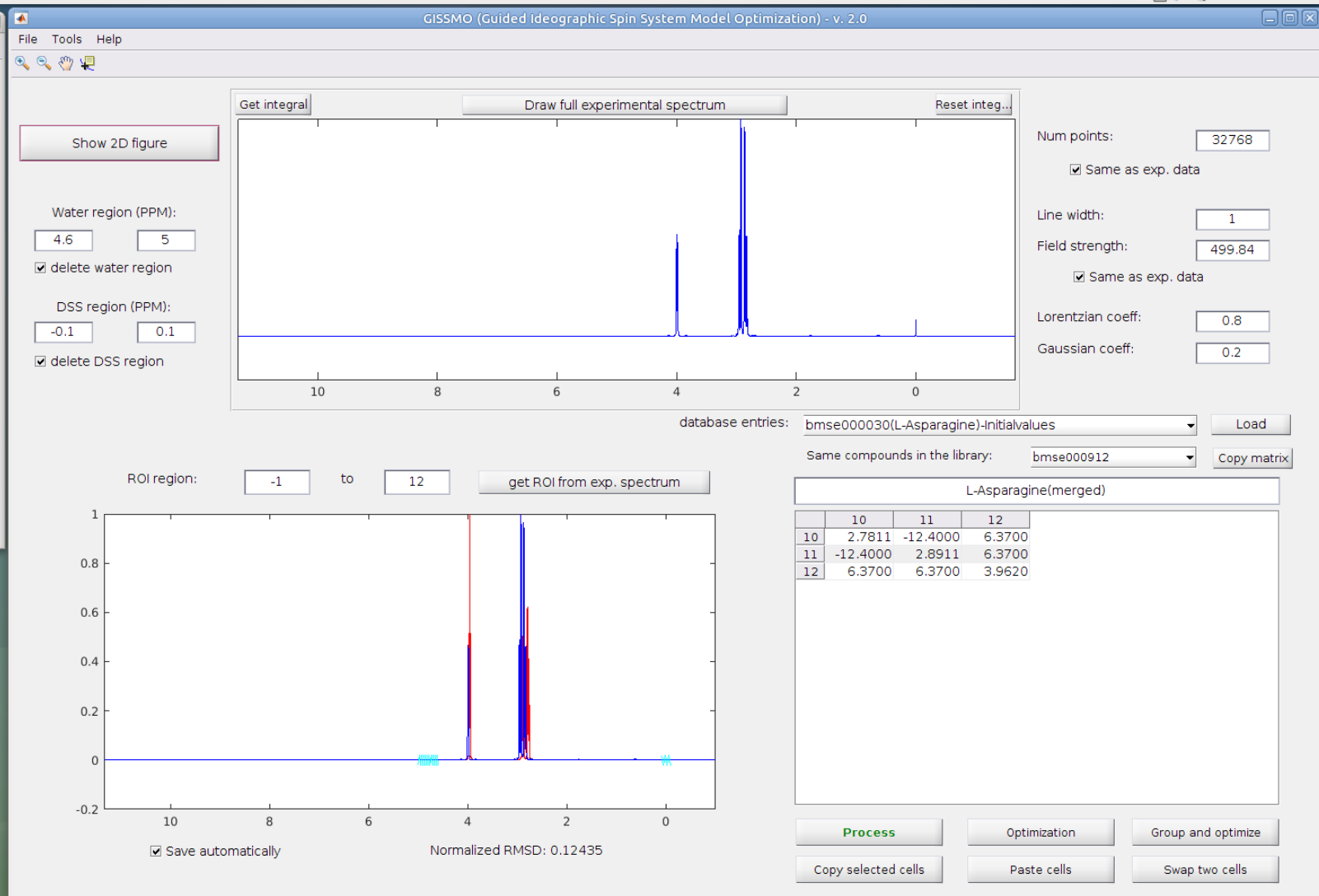
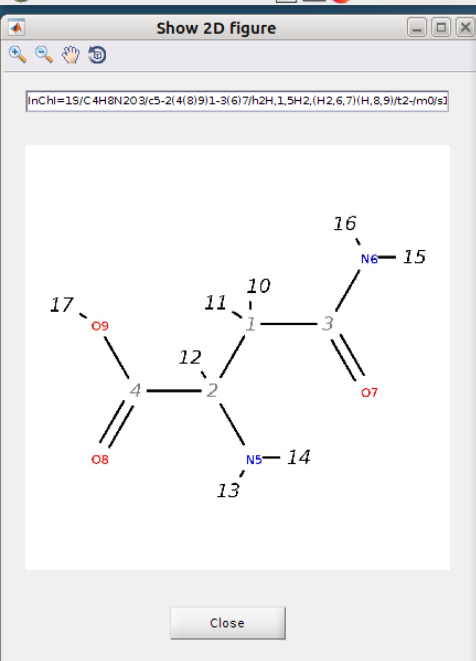
ff: 0.2

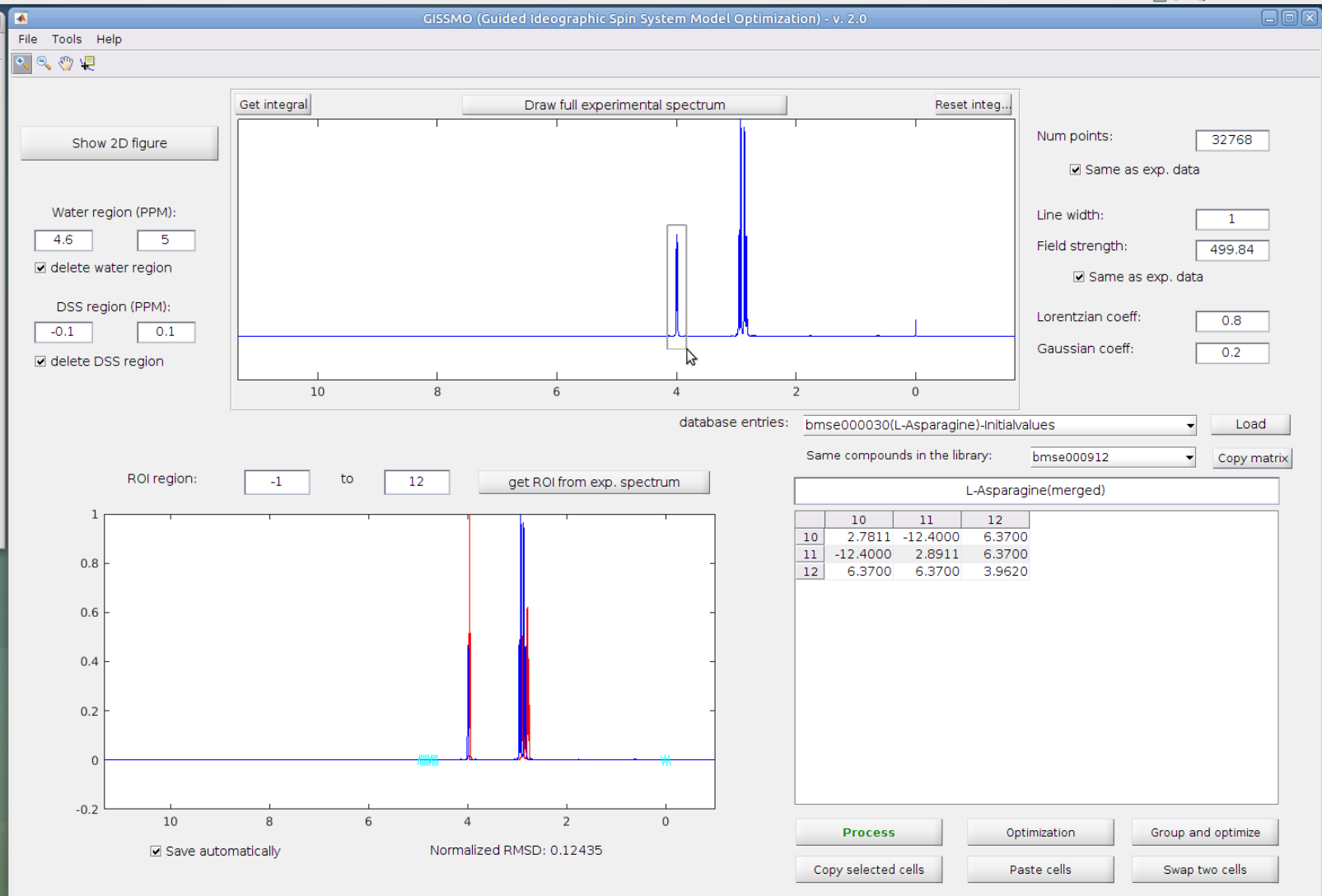
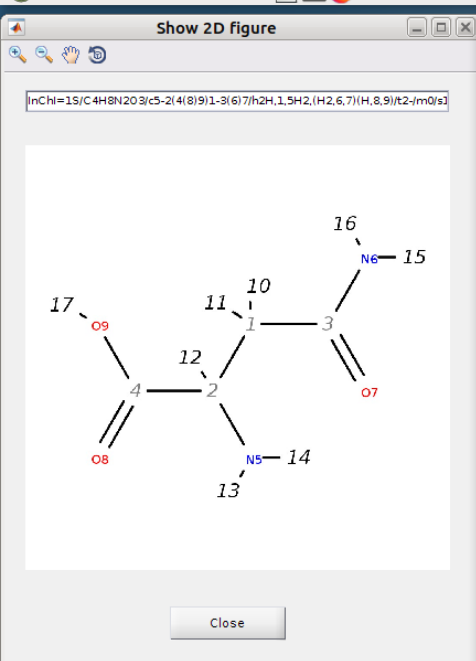
Load

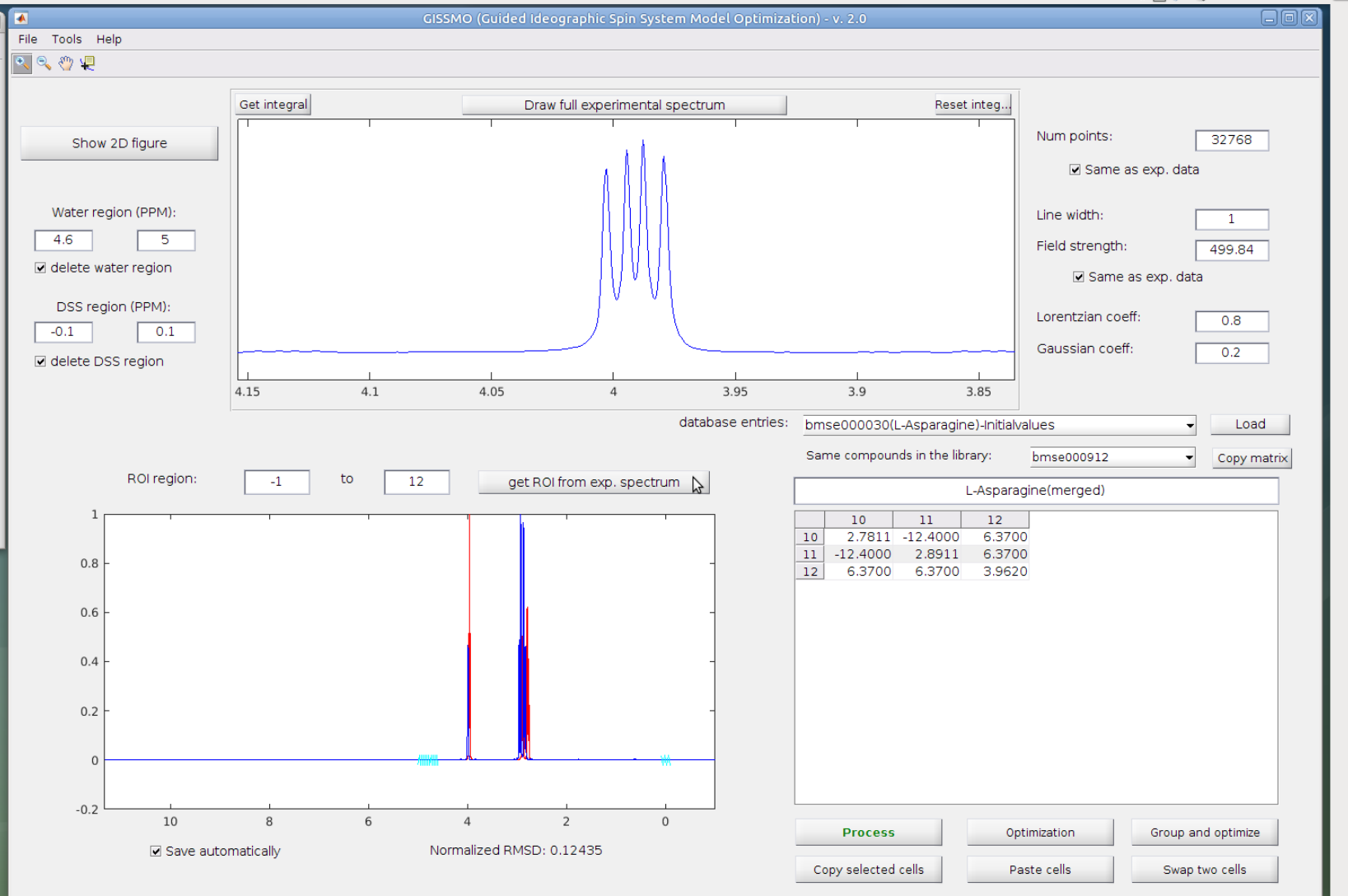
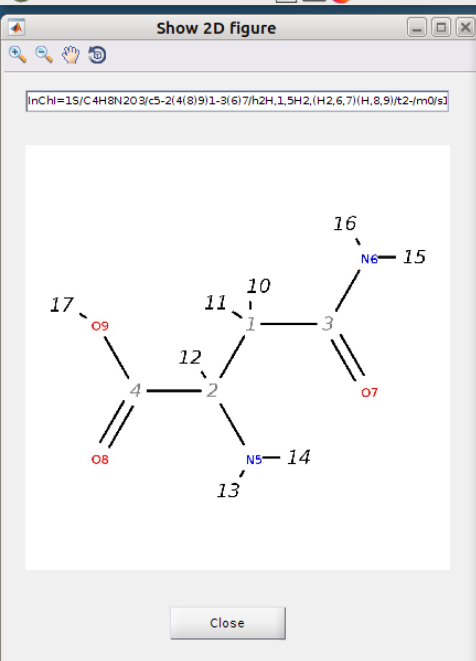
Copy matrix

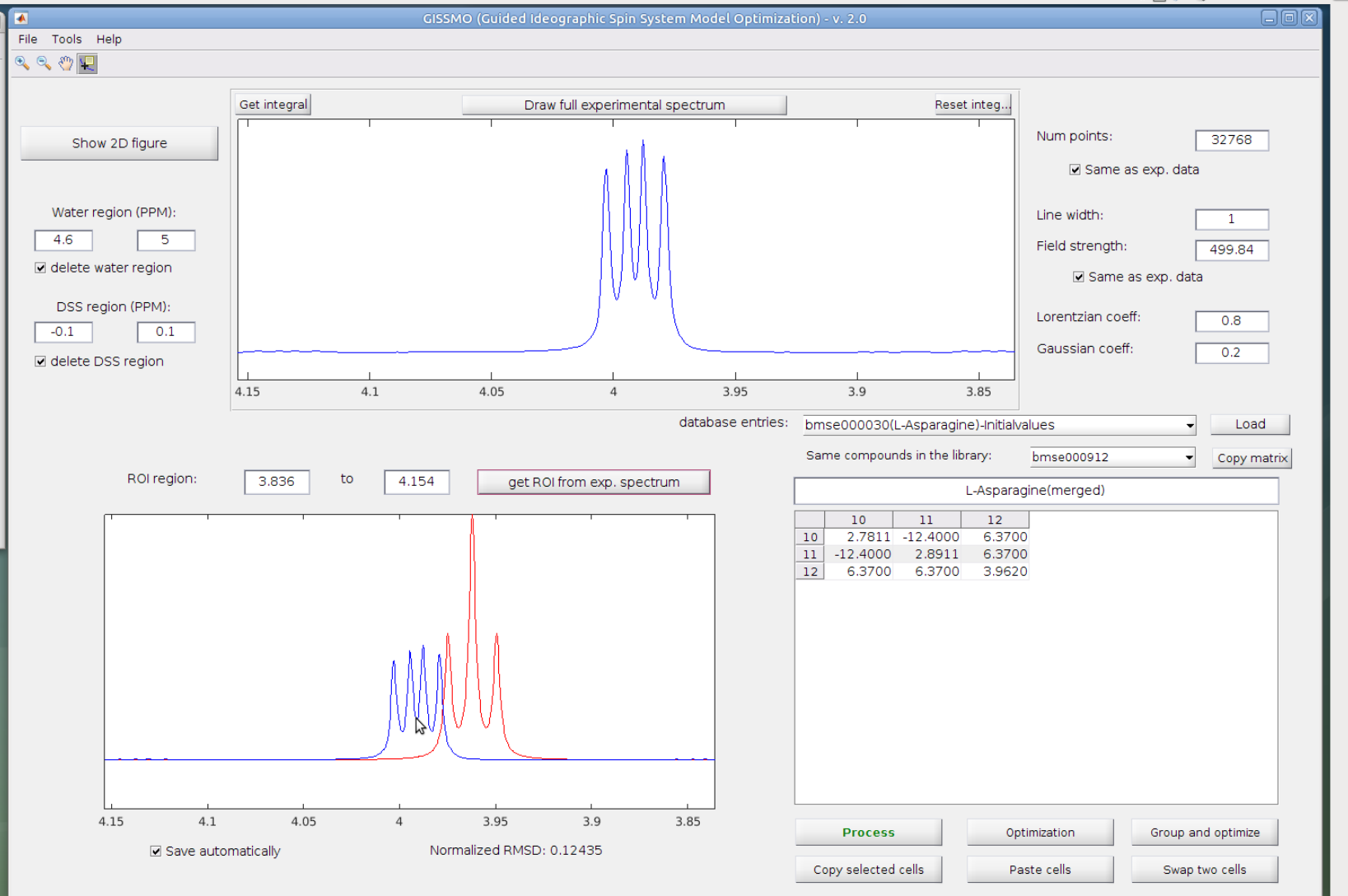
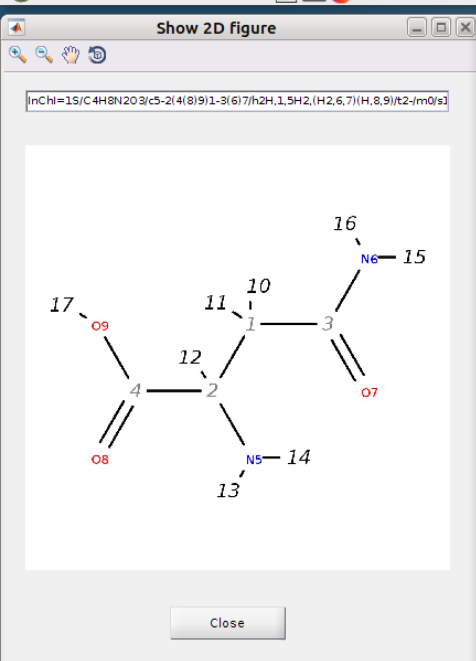
Process Optimization Group and optimize

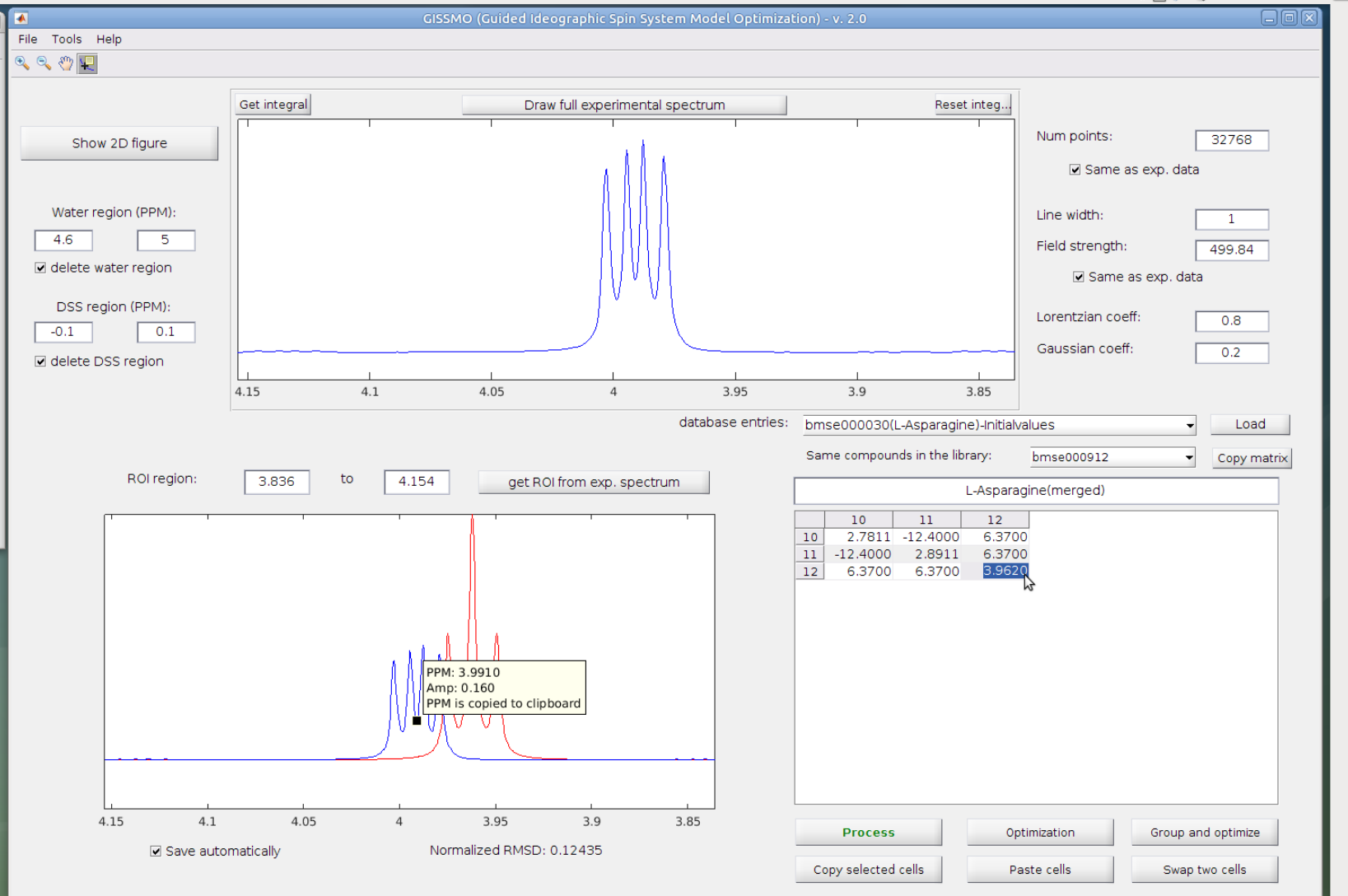
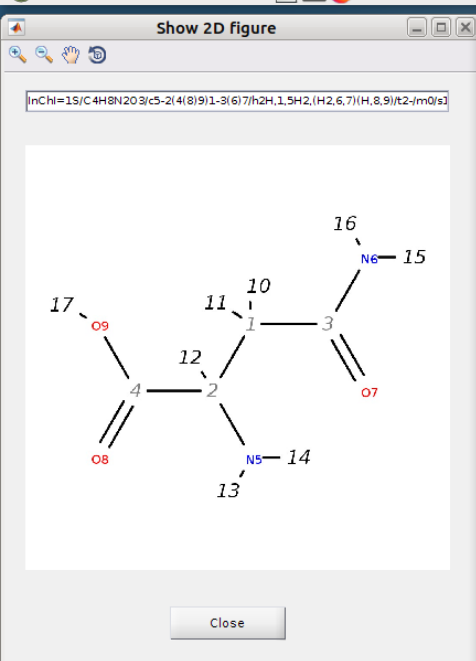
Copy selected cells Paste cells Swap two cells

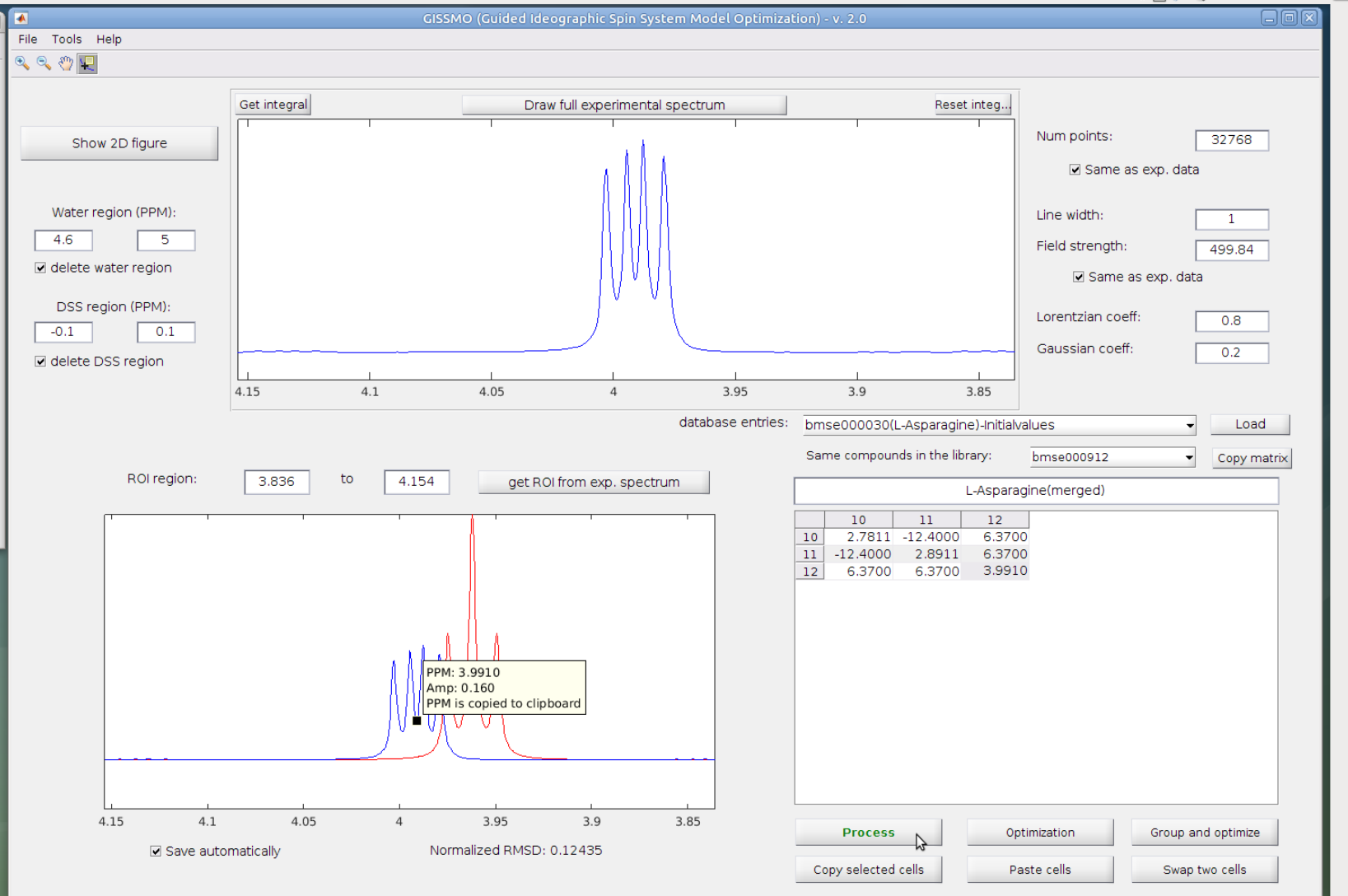
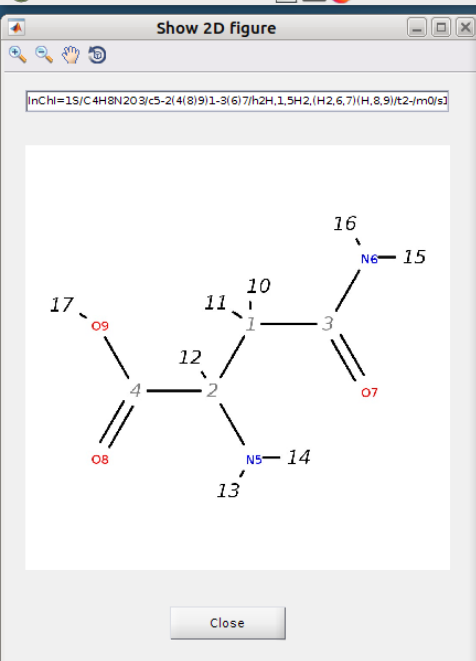


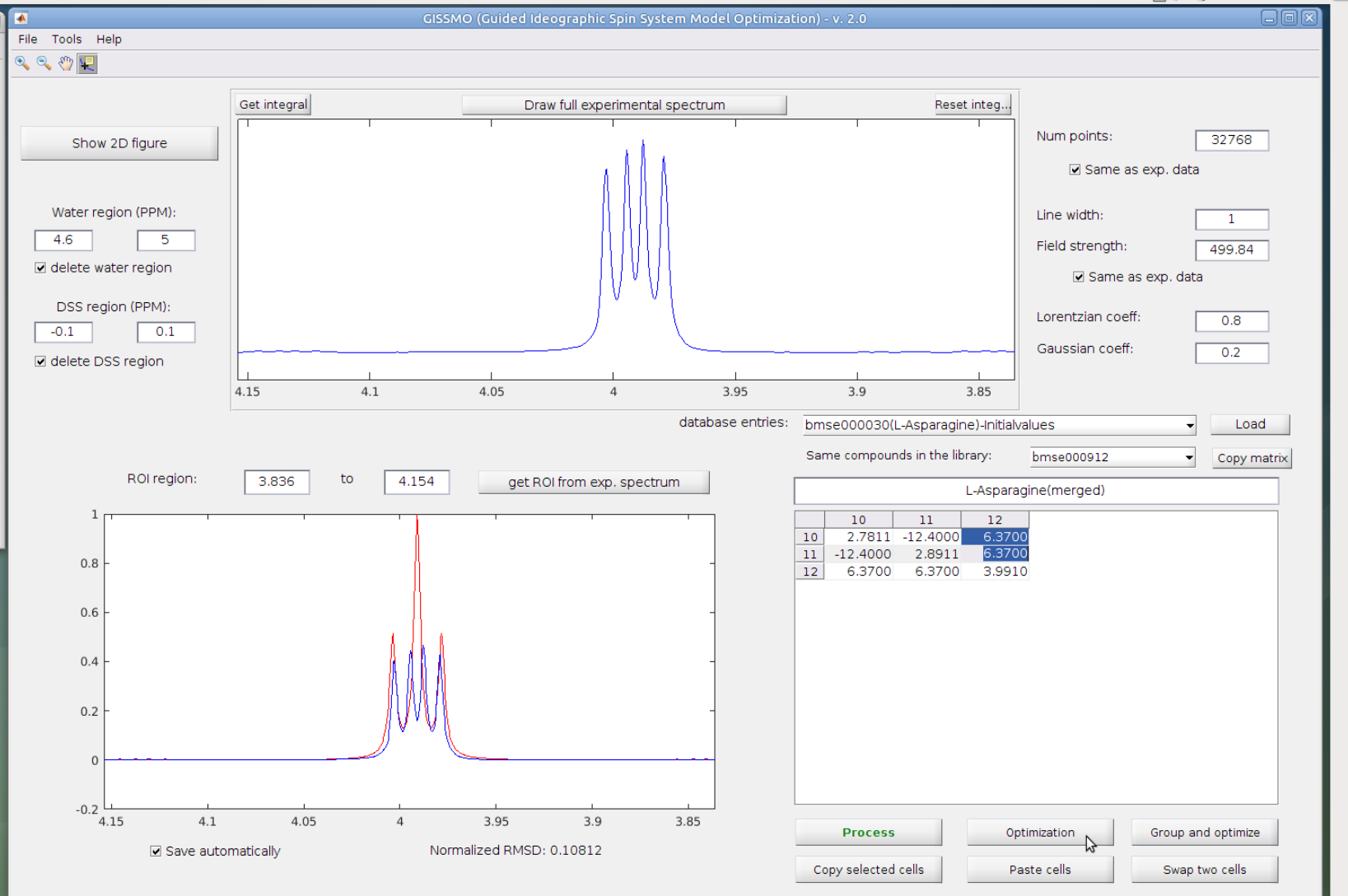
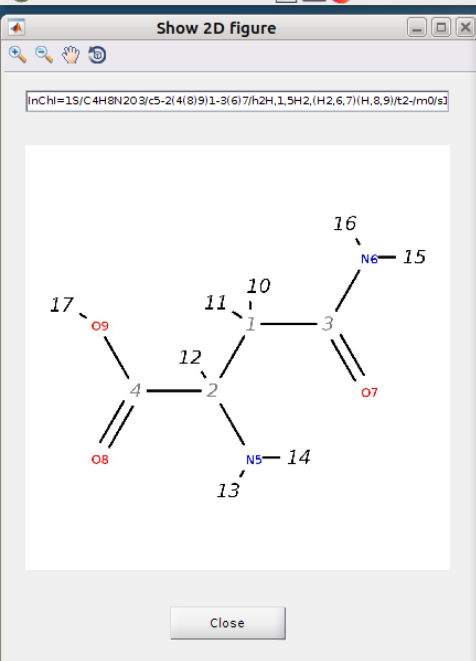


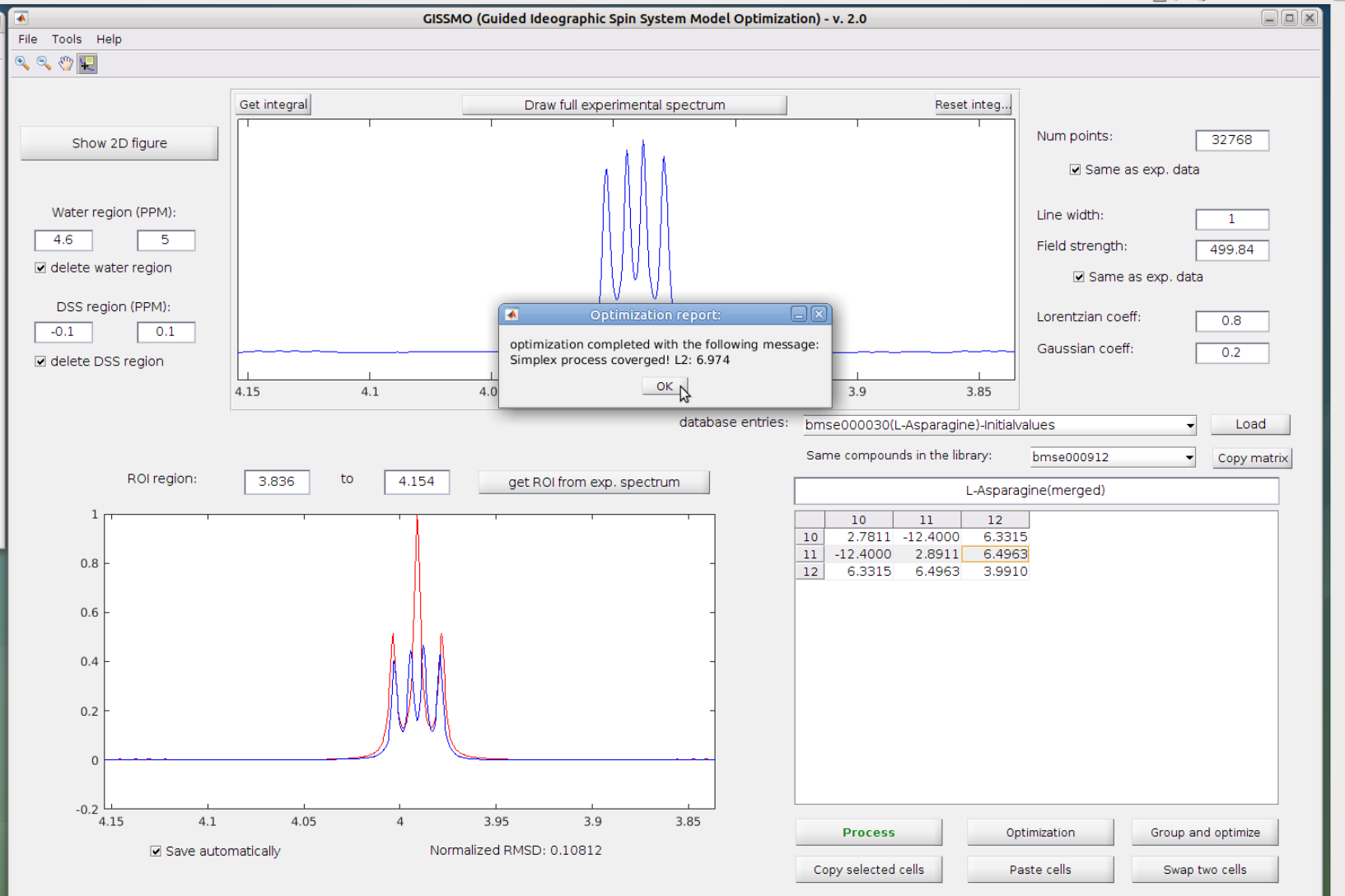
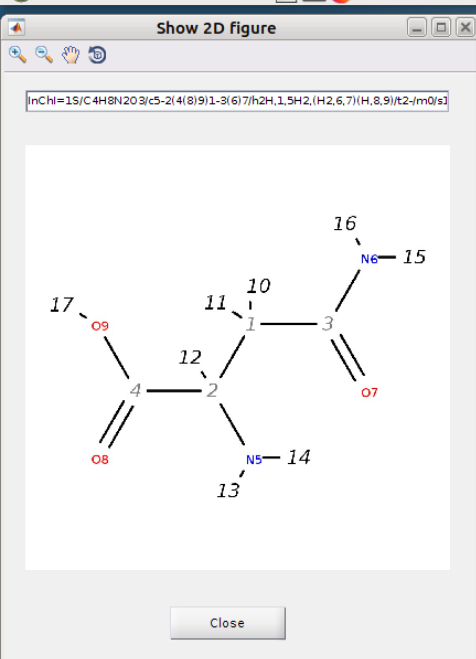


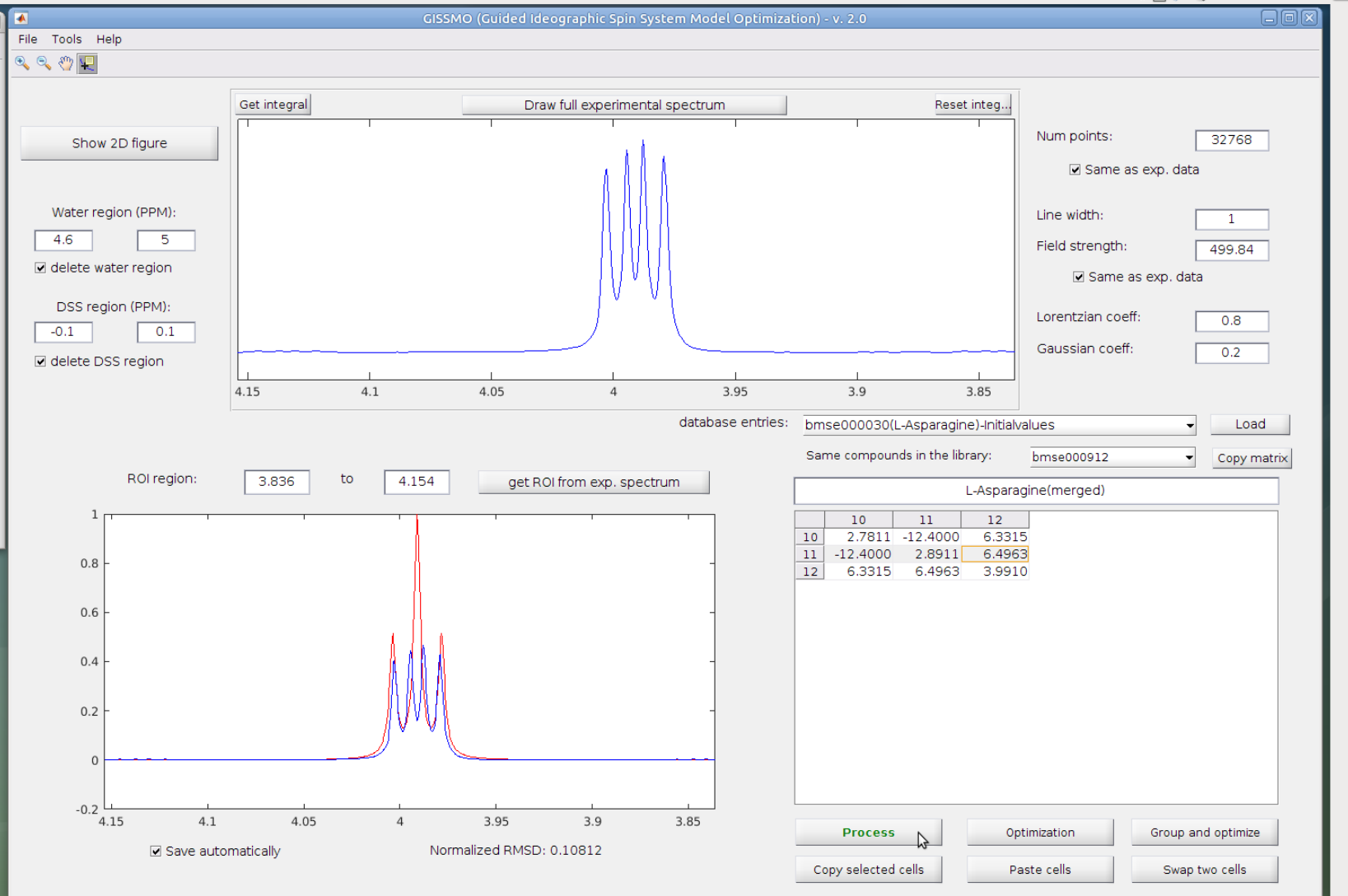
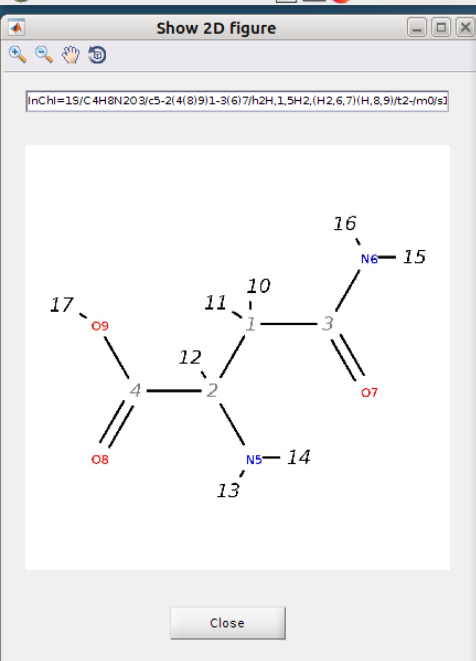


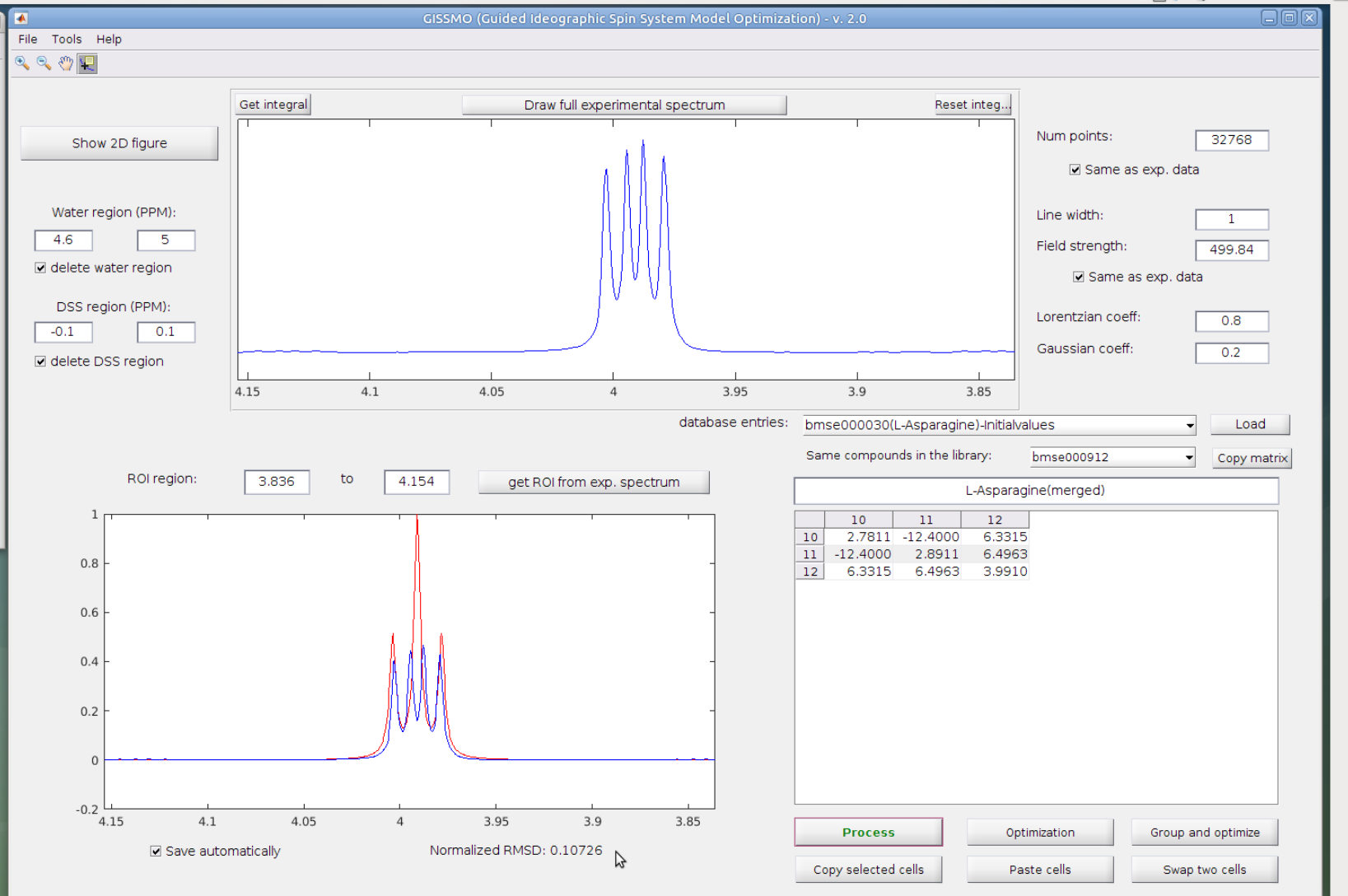
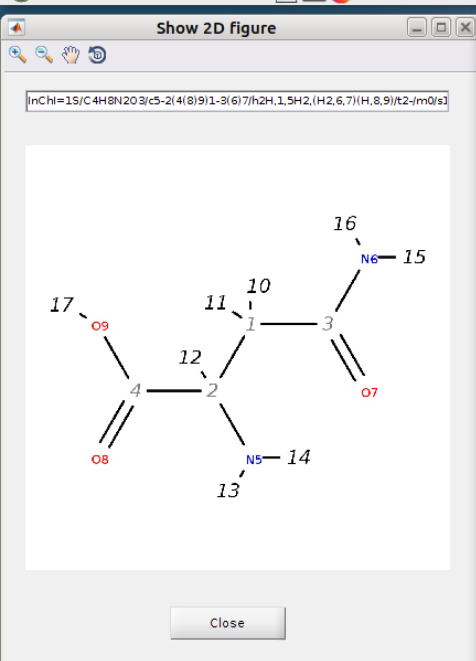


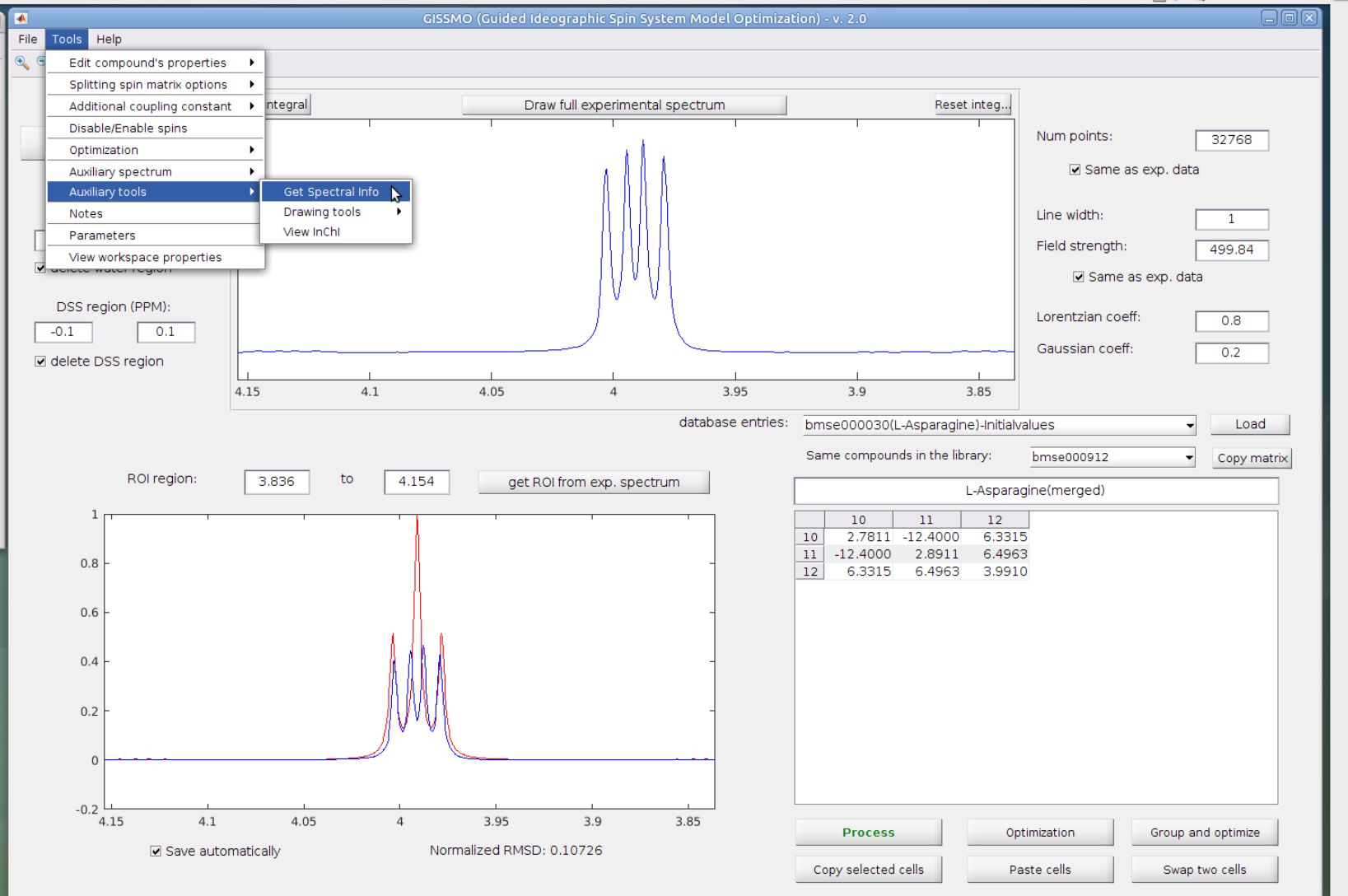
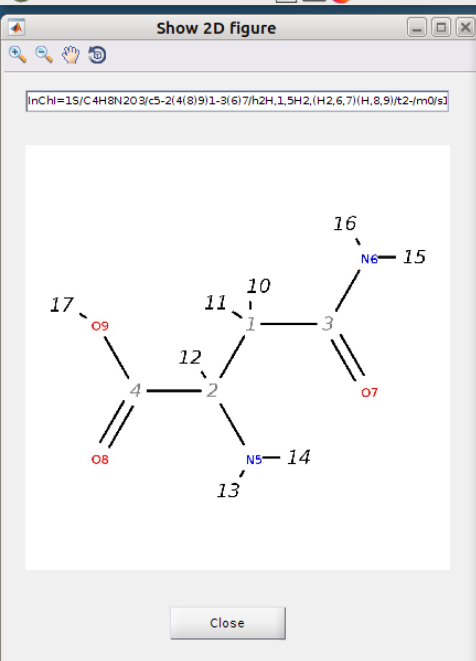


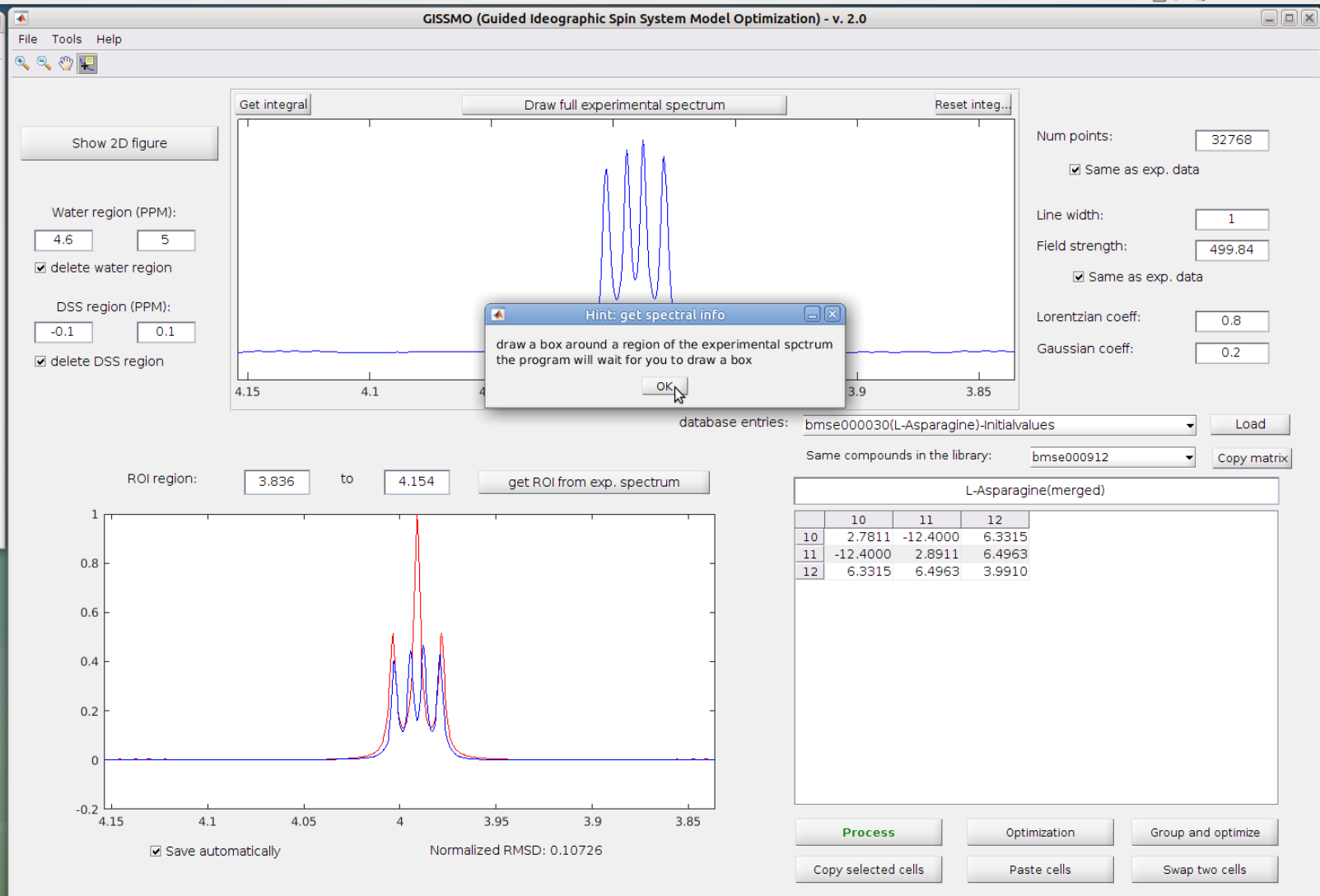
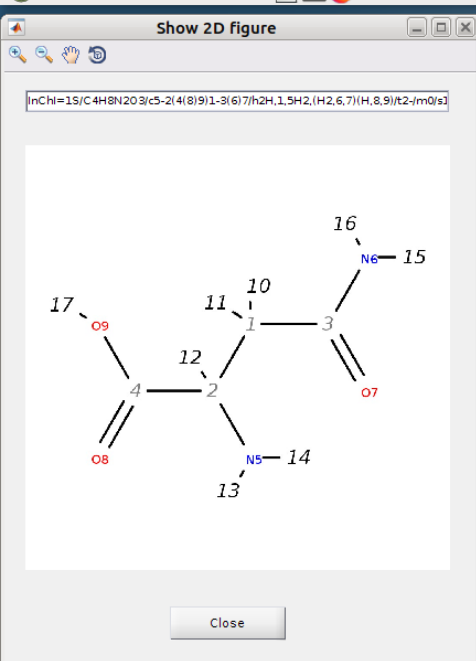


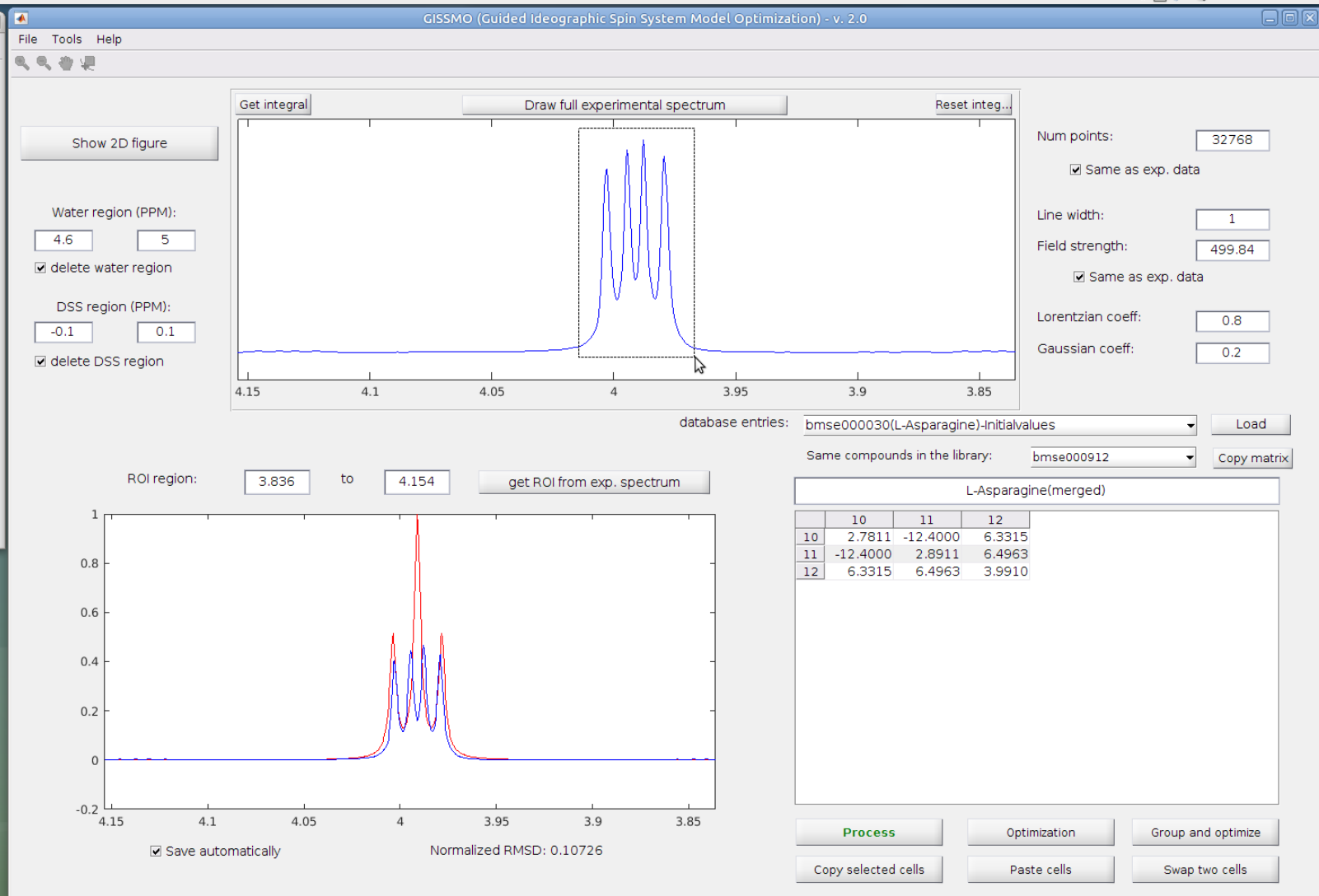
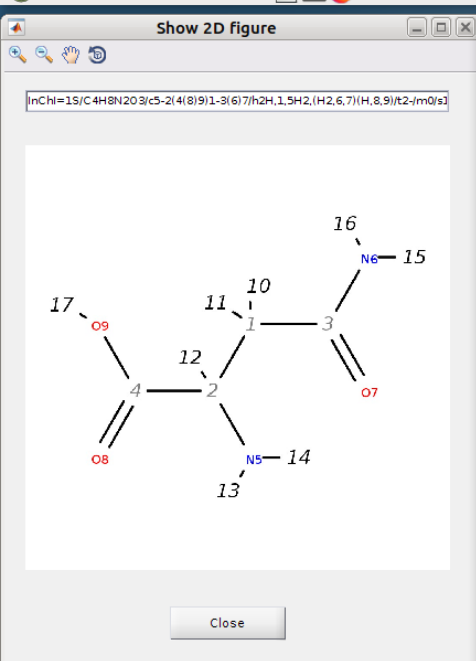


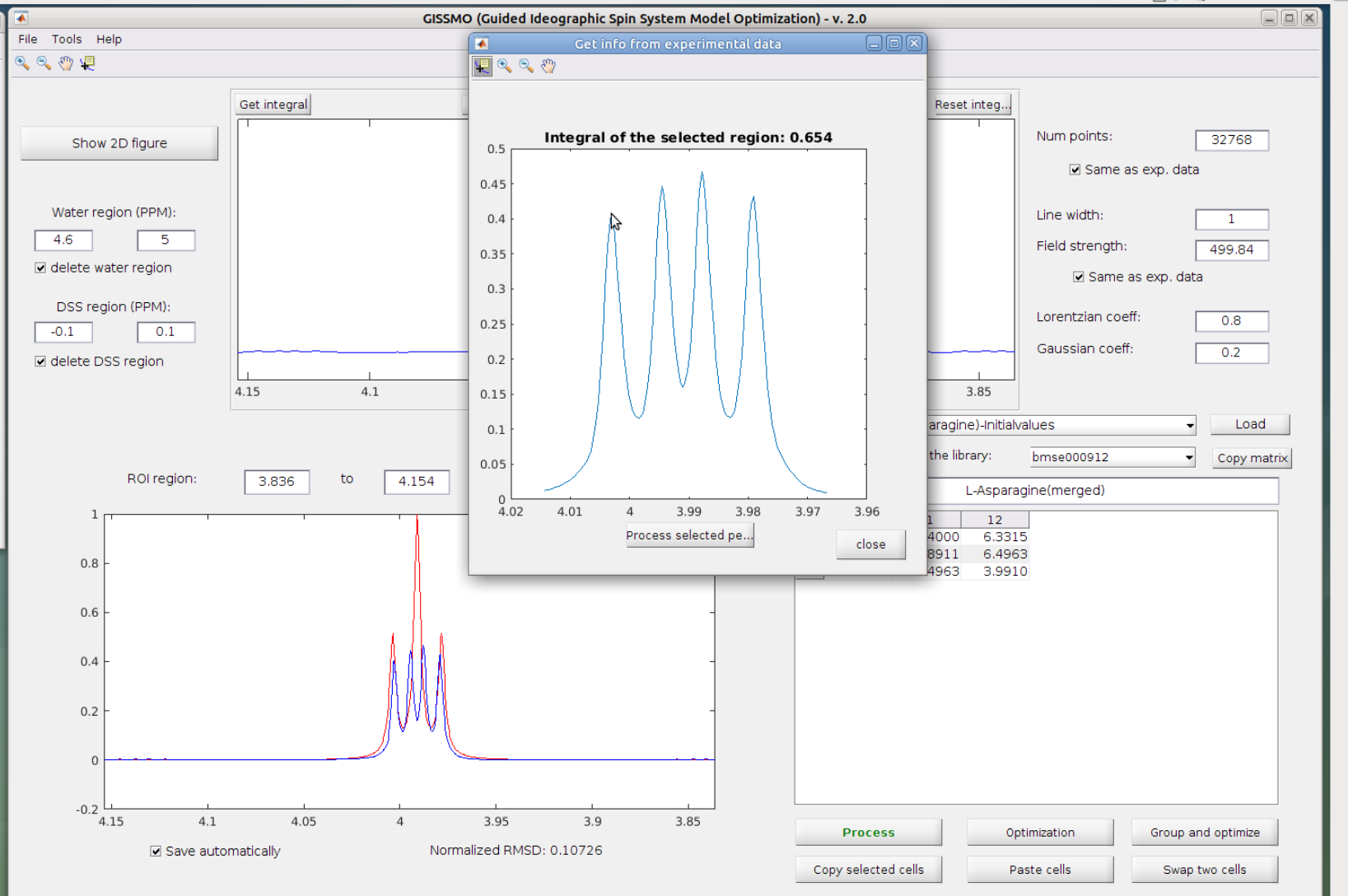
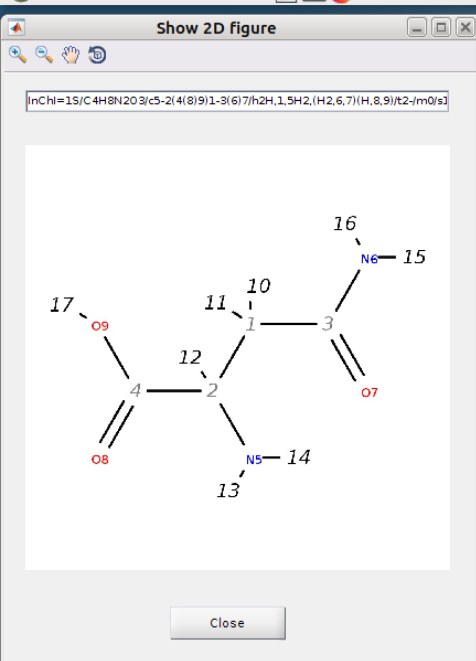


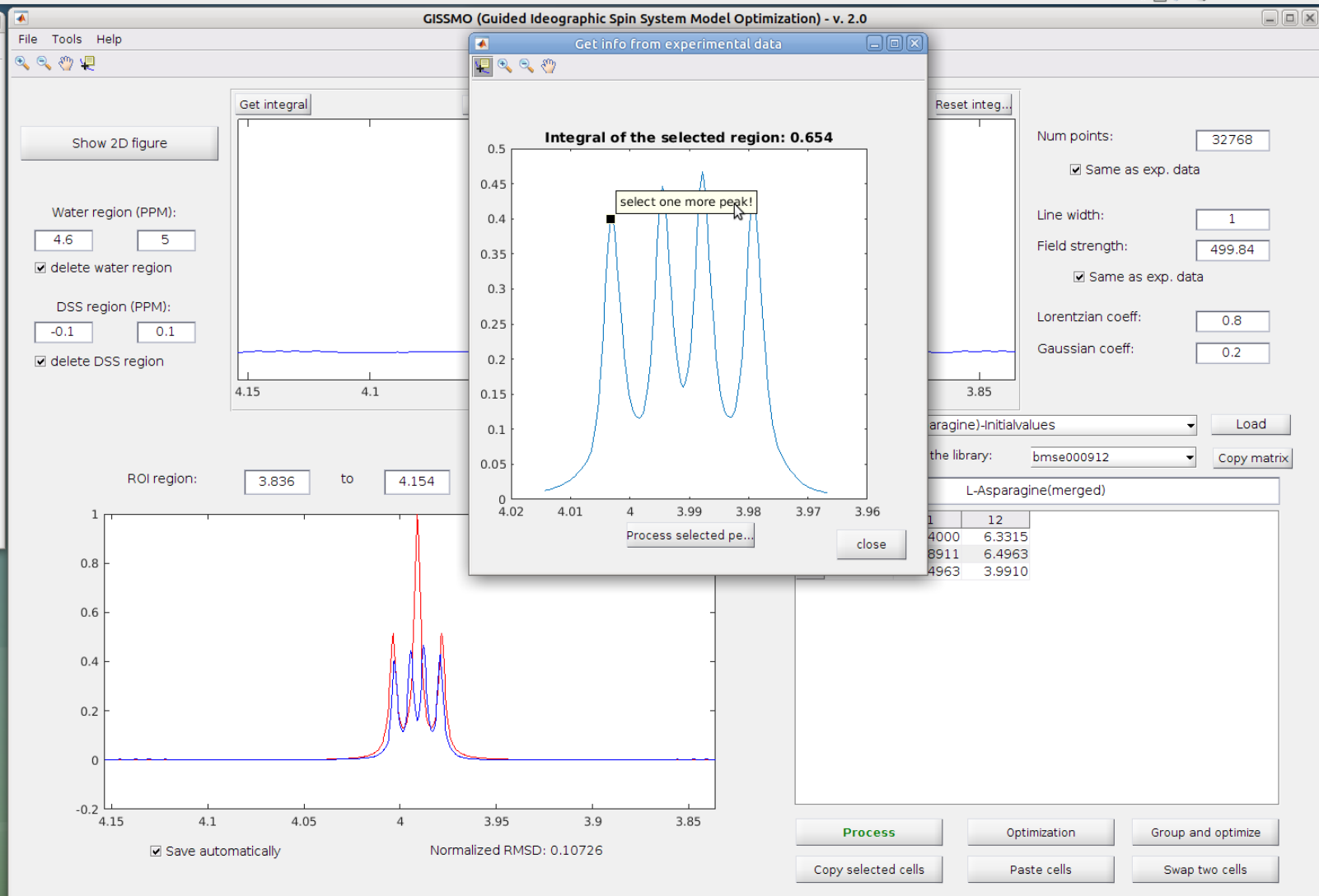
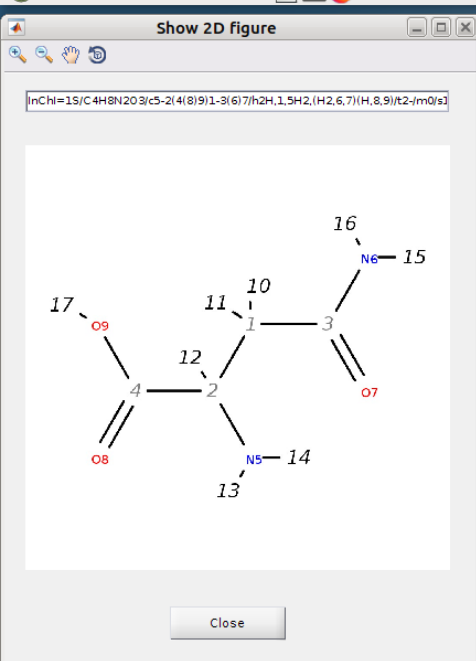


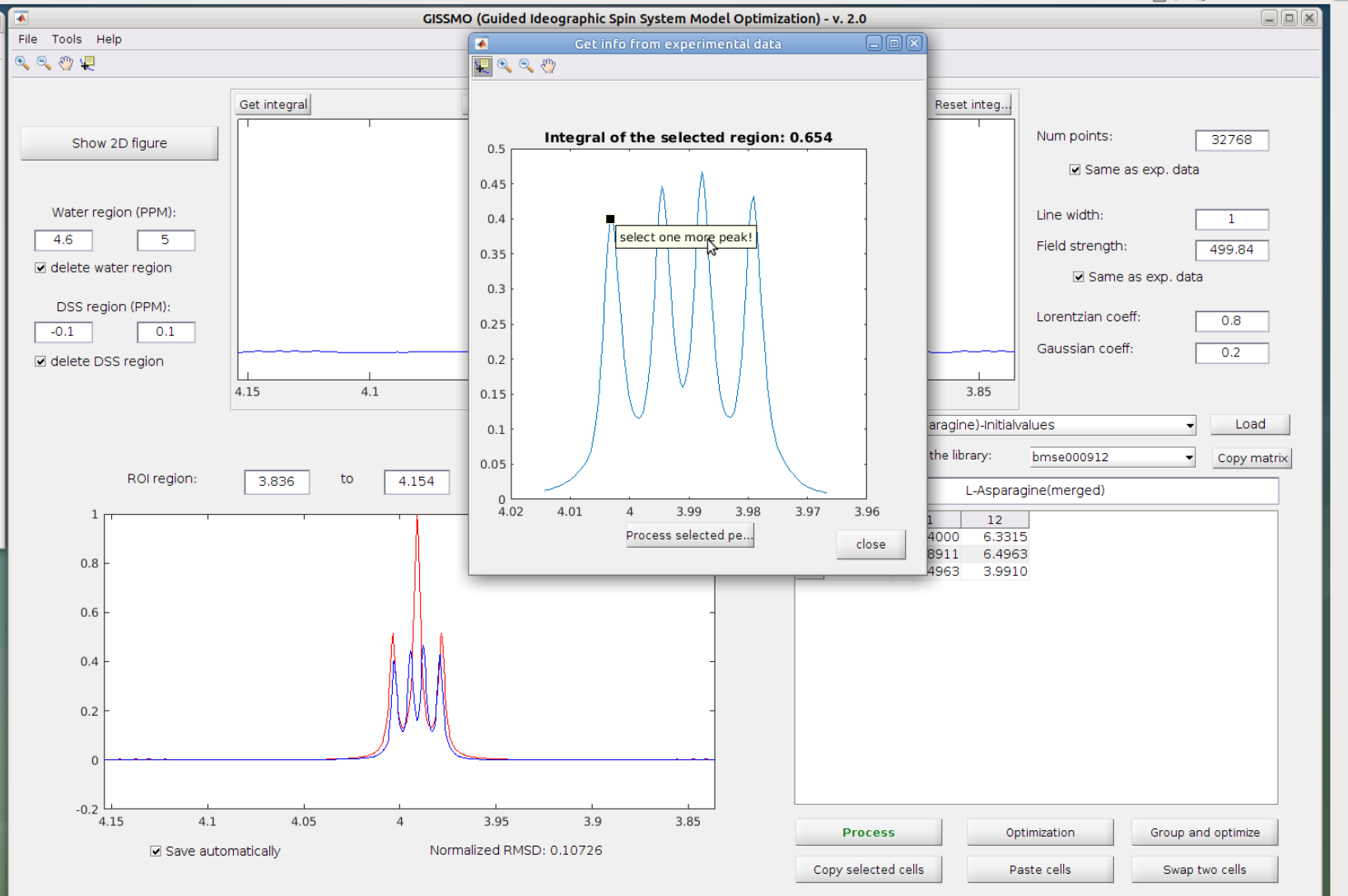
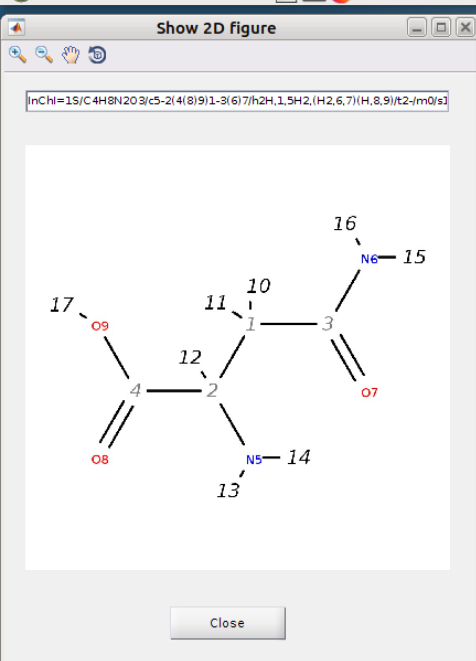


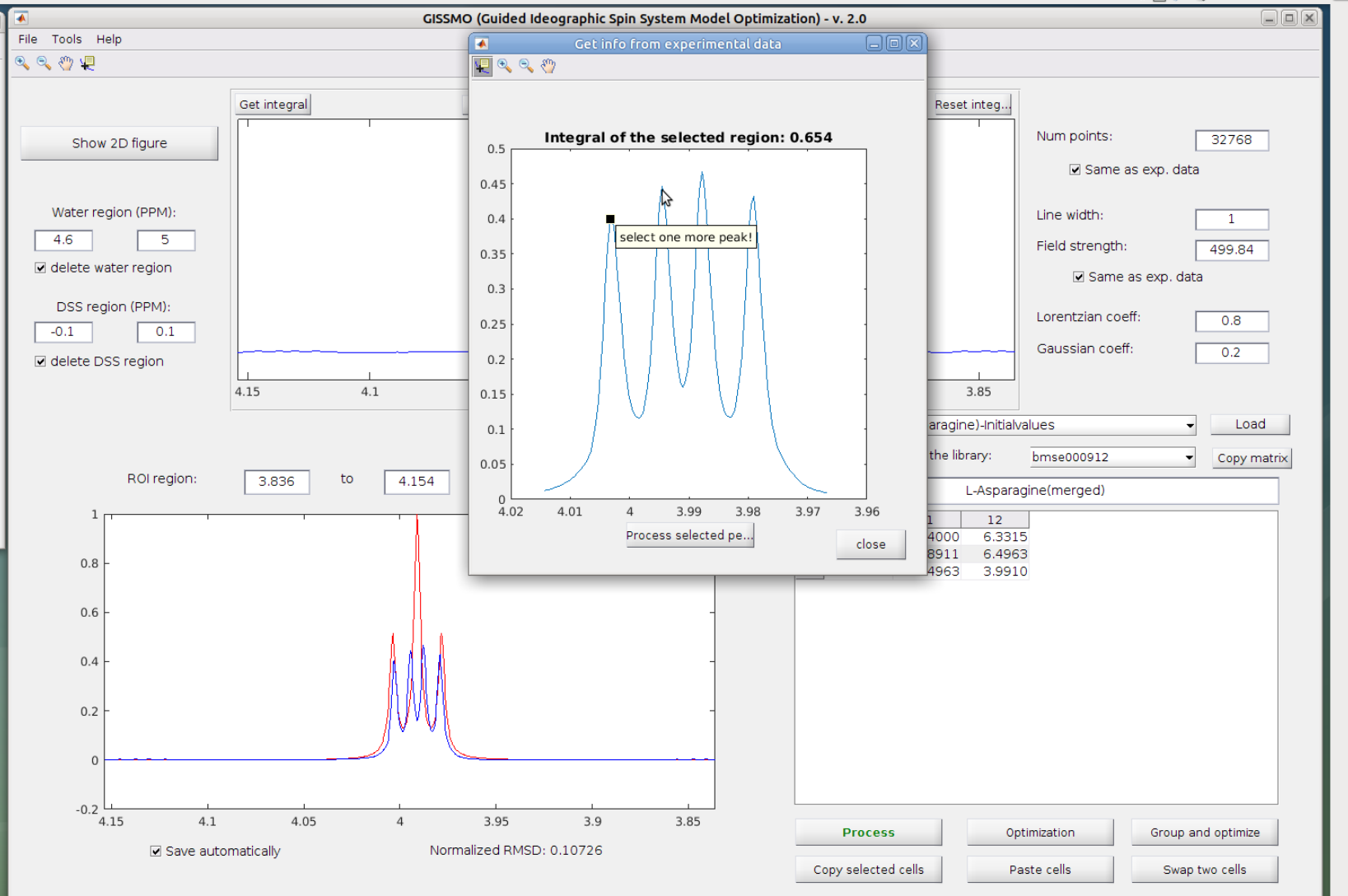
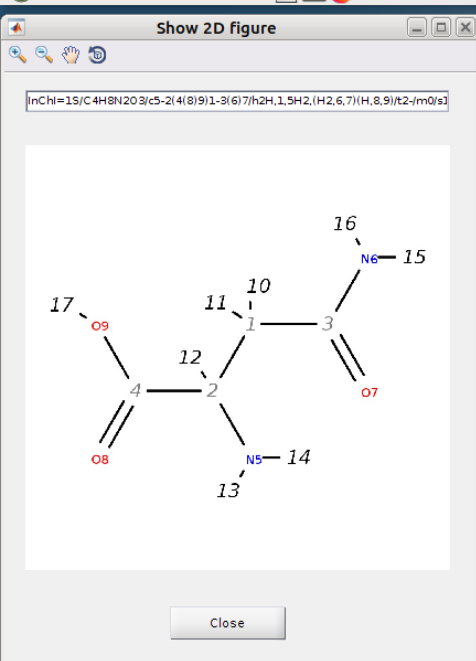


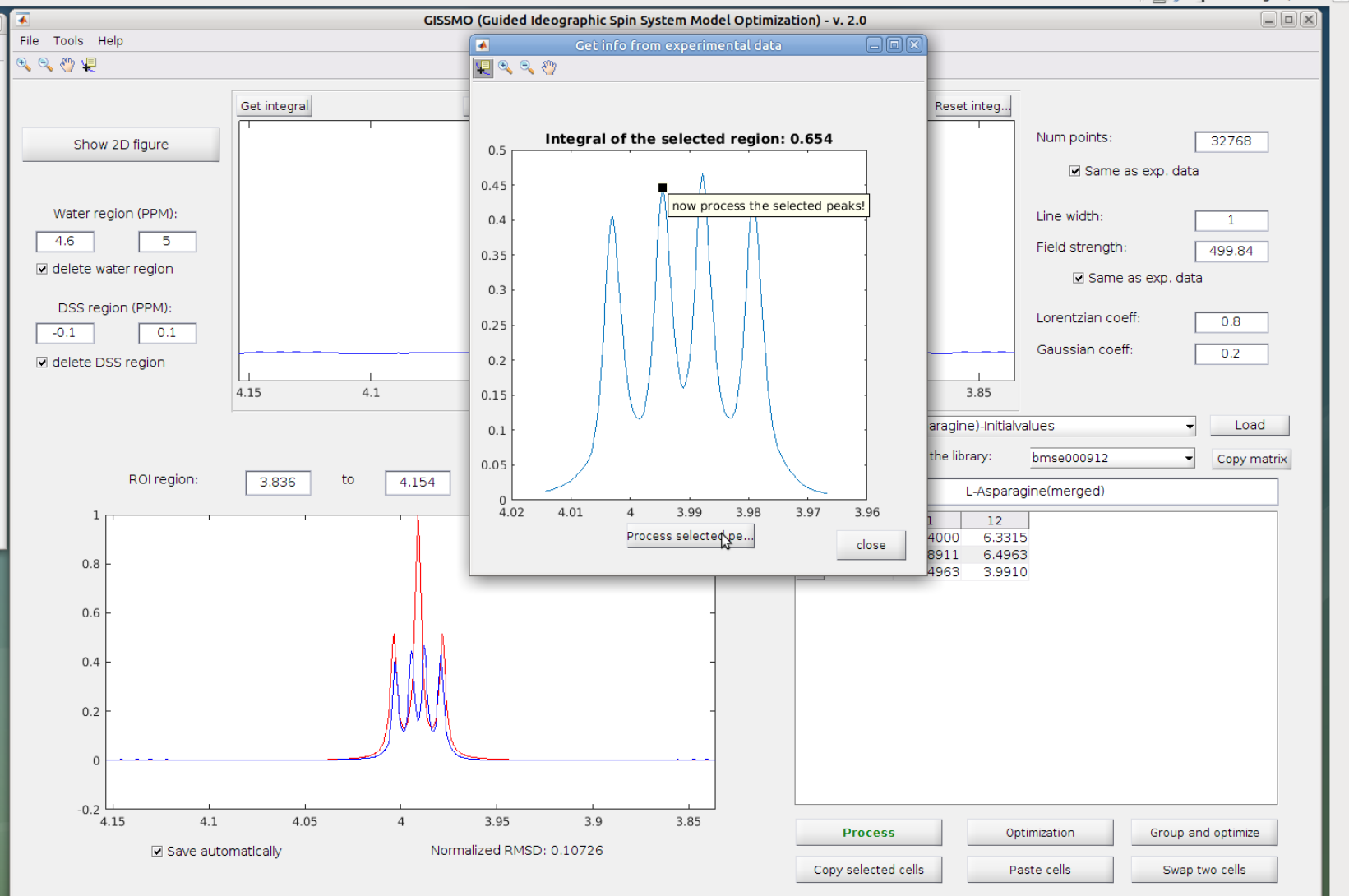


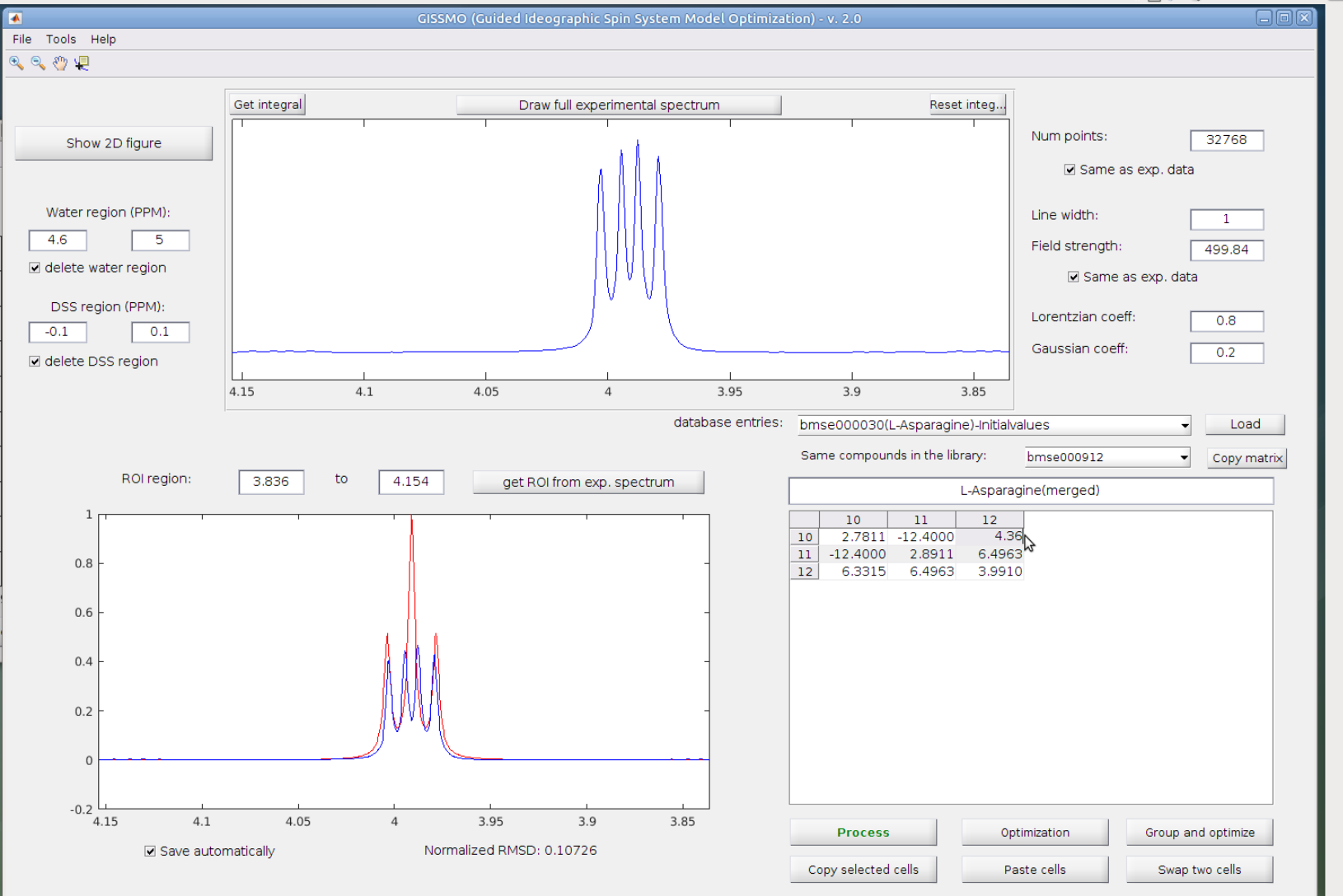
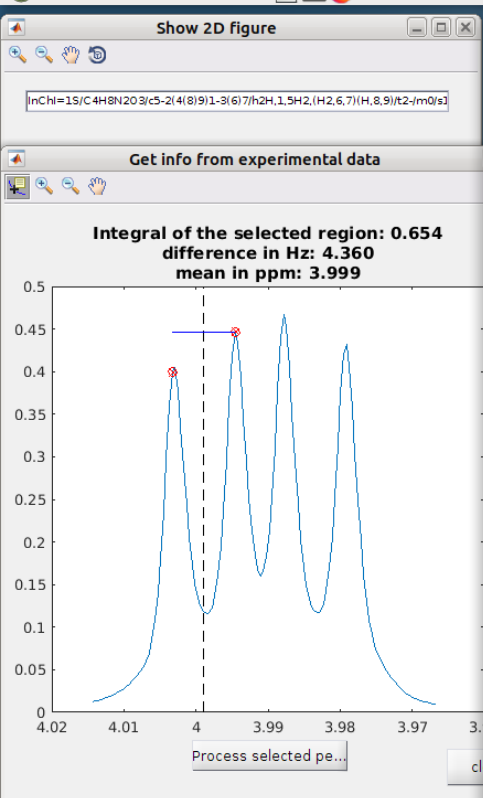








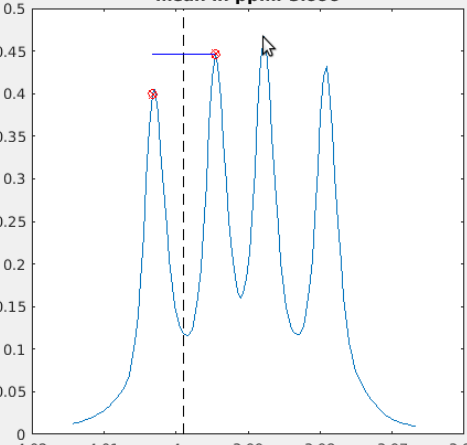




Show 2D figure

Get info from experimental data

Integral of the selected region: 0.654
difference in Hz: 4.360
mean in ppm: 3.999



Process selected pe... close

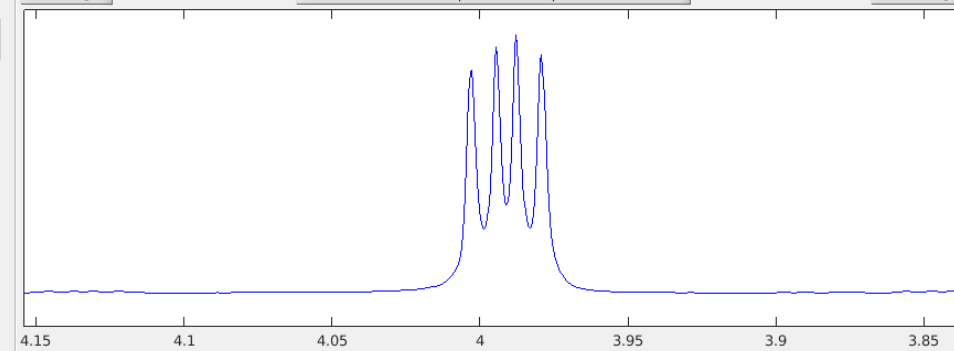
GISSMO (Guided Ideographic Spin System Model Optimization) - v. 2.0

File Tools Help

Get integral

Draw full experimental spectrum

Reset integ...



water region (PPM):
5 to 5

ete water region

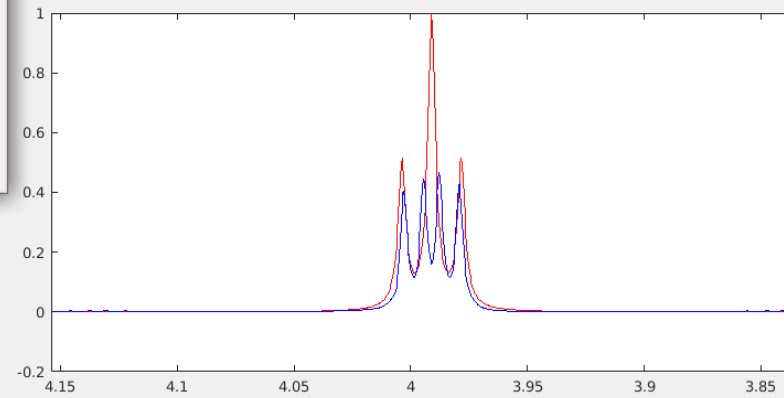
DSS region (PPM):
1 to 0.1

ete DSS region

database entries: bmse000030(L-Asparagine)-Initialvalues Load

Same compounds in the library: bmse000912 Copy matrix

ROI region: 3.836 to 4.154 get ROI from exp. spectrum



Save automatically Normalized RMSD: 0.10726

Process

Optimization

Group and optimize

Copy selected cells

Paste cells

Swap two cells

Num points: 32768

☒ Same as exp. data

Line width: 1

Field strength: 499.84

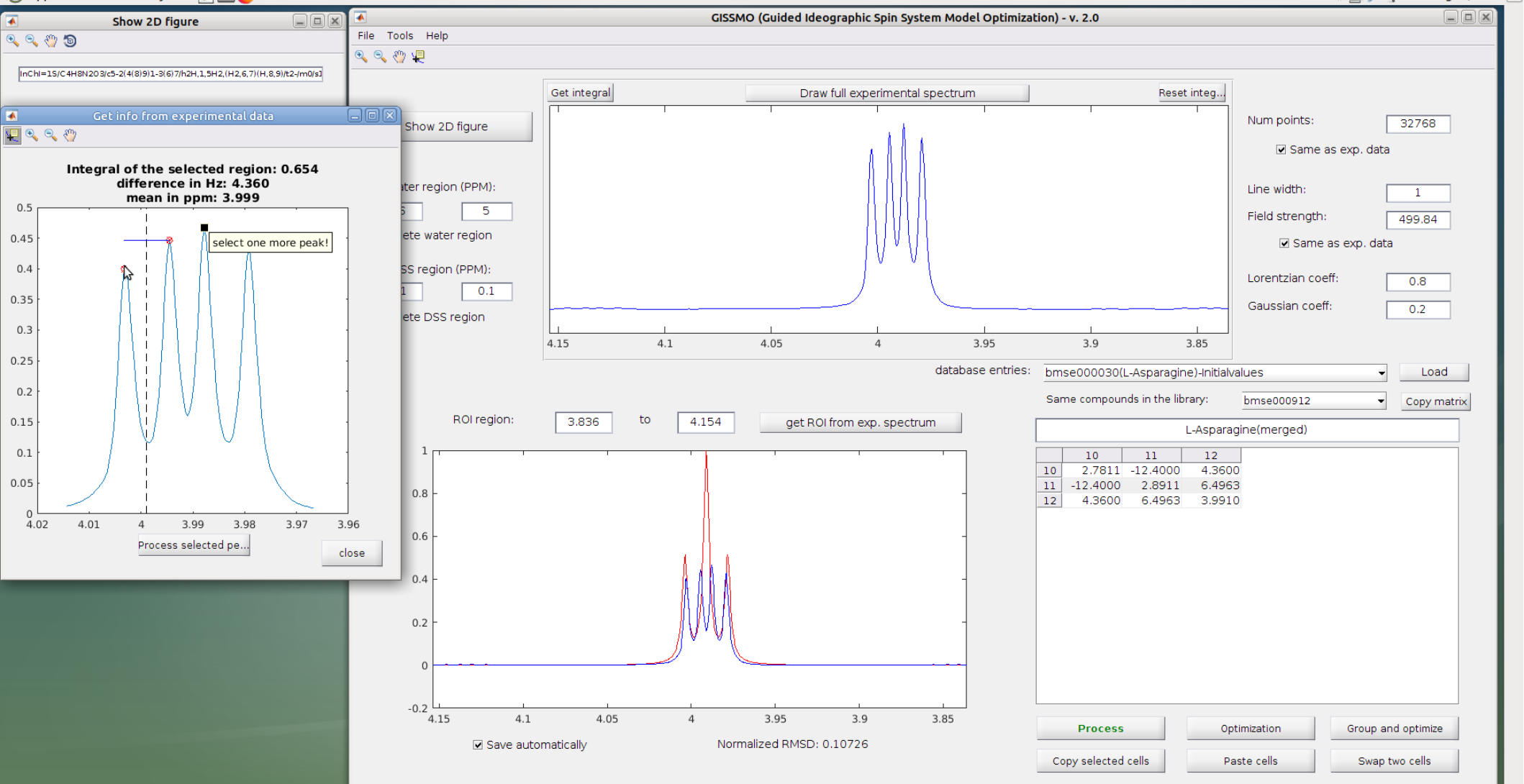
☒ Same as exp. data

Lorentzian coeff: 0.8

Gaussian coeff: 0.2

L-Asparagine(merged)

	10	11	12
10	2.7811	-12.4000	4.36
11	-12.4000	2.8911	6.4963
12	6.3315	6.4963	3.9910



Show 2D figure

Get info from experimental data

Integral of the selected region: 0.654
difference in Hz: 4.360
mean in ppm: 3.999

now process the selected peaks!

Process selected peaks...

close

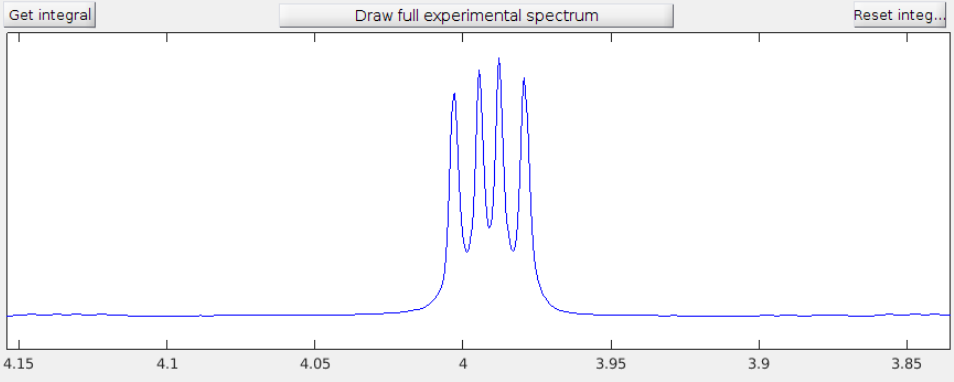
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File Tools Help

Get integral

Draw full experimental spectrum

Reset integ...

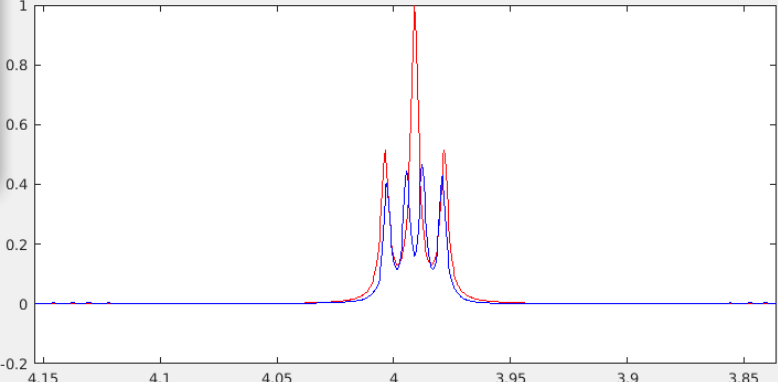


Num points: 32768
☒ Same as exp. data
 Line width: 1
 Field strength: 499.84
☒ Same as exp. data
 Lorentzian coeff: 0.8
 Gaussian coeff: 0.2

water region (PPM): 5 to 5
 delete water region
 DSS region (PPM): 1 to 0.1
 delete DSS region

database entries: bmse000030(L-Asparagine)-Initialvalues Load
 Same compounds in the library: bmse000912 Copy matrix

ROI region: 3.836 to 4.154 get ROI from exp. spectrum



Save automatically Normalized RMSD: 0.10726

Process

Optimization

Group and optimize

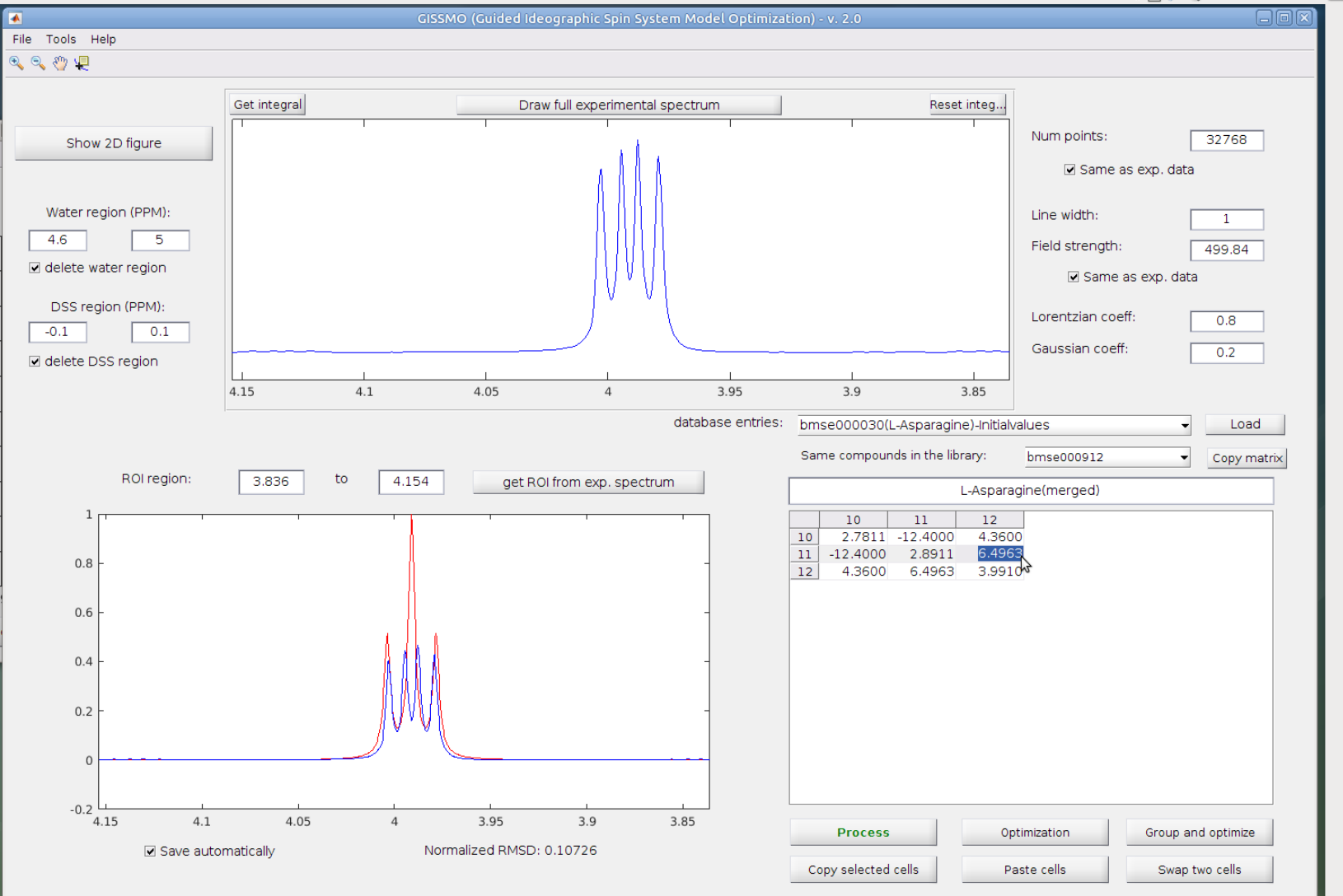
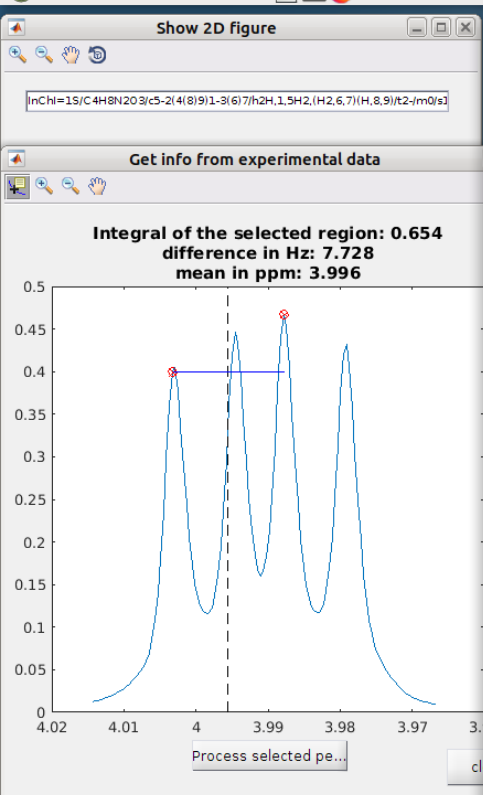
Copy selected cells

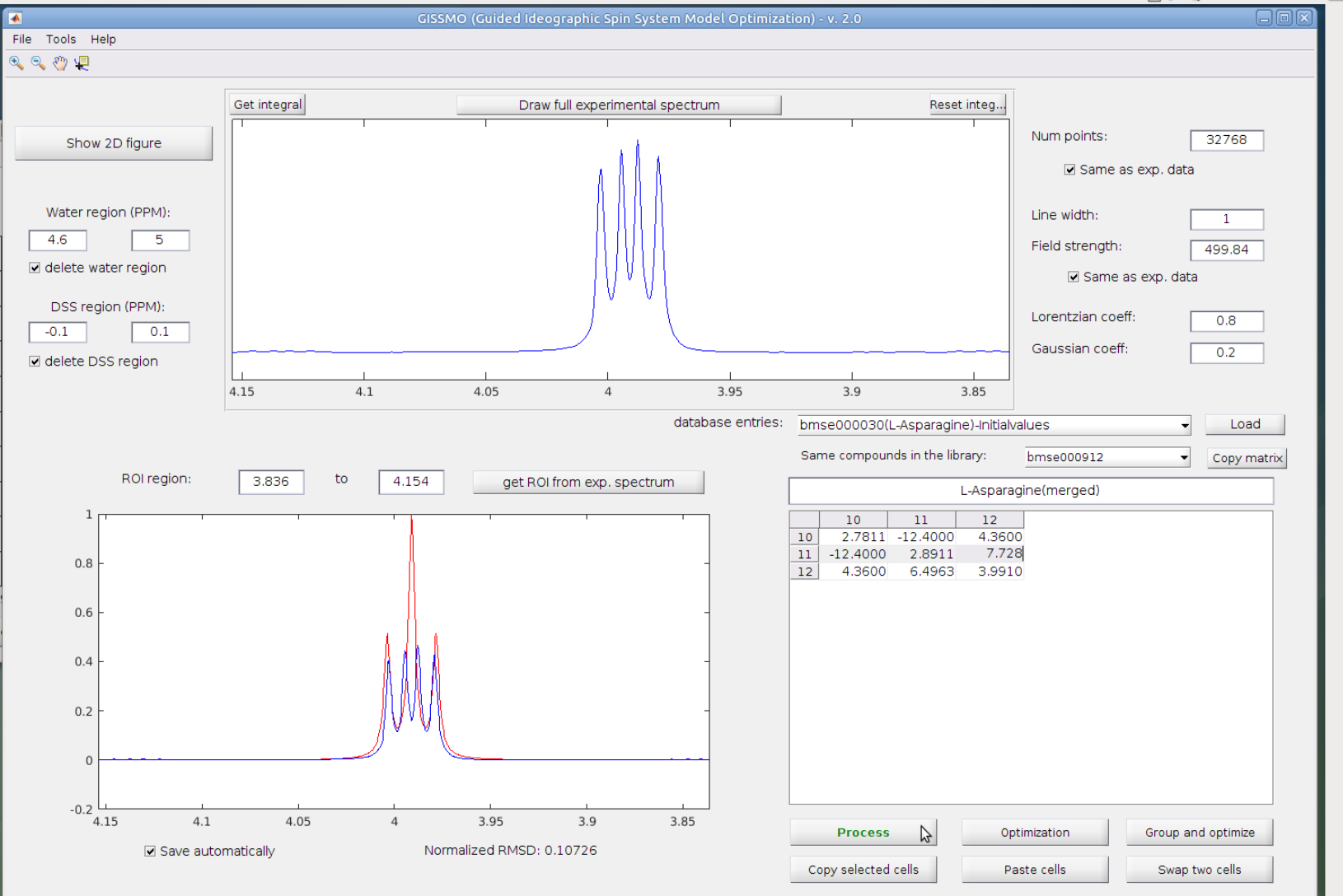
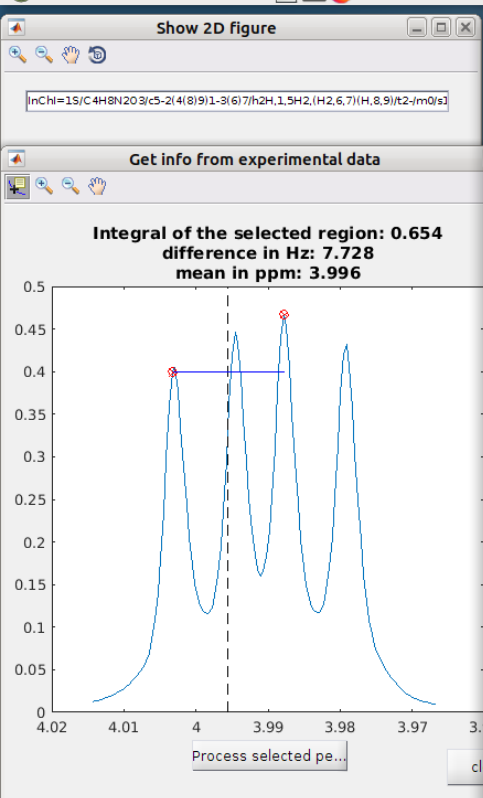
Paste cells

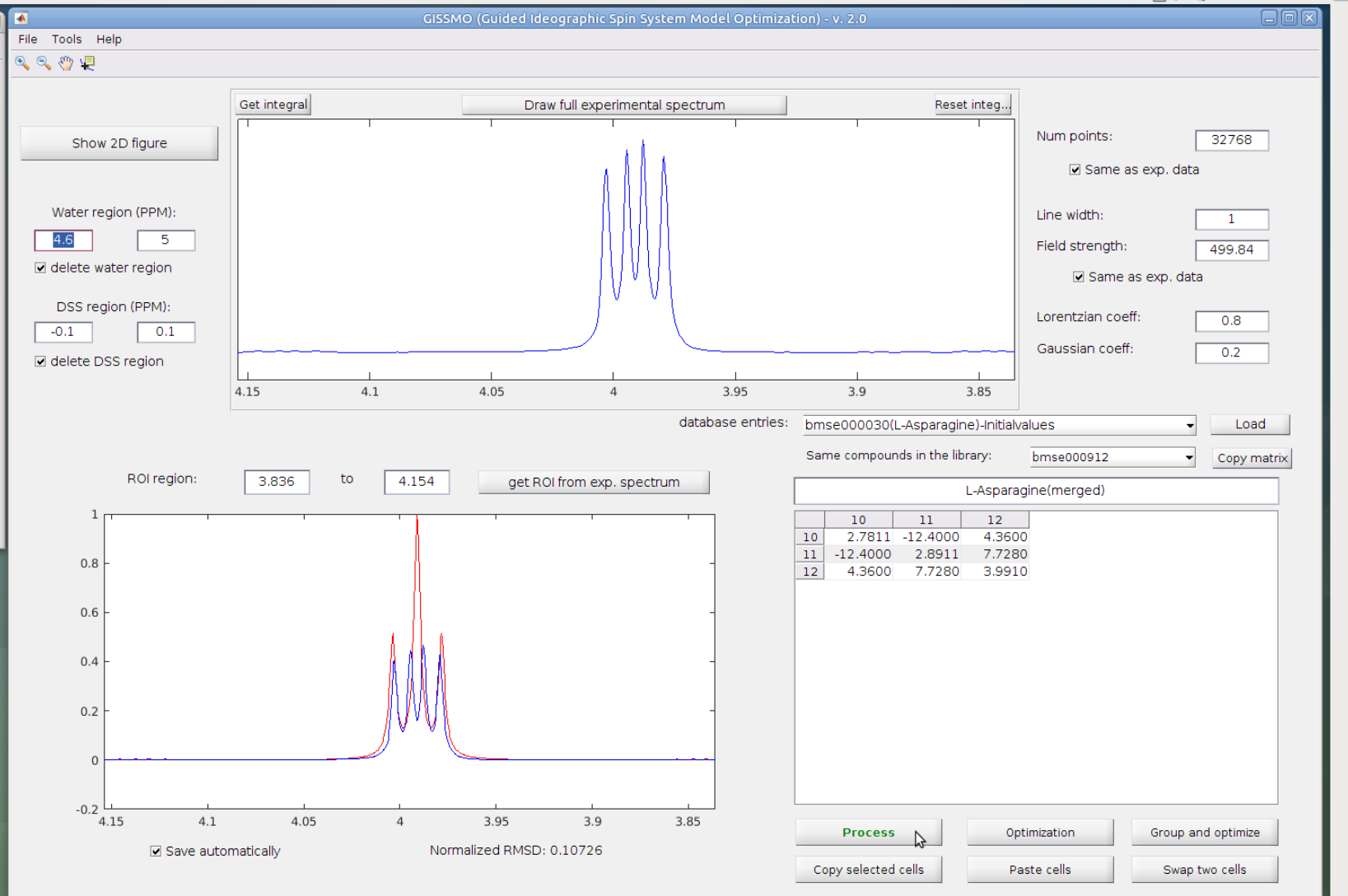
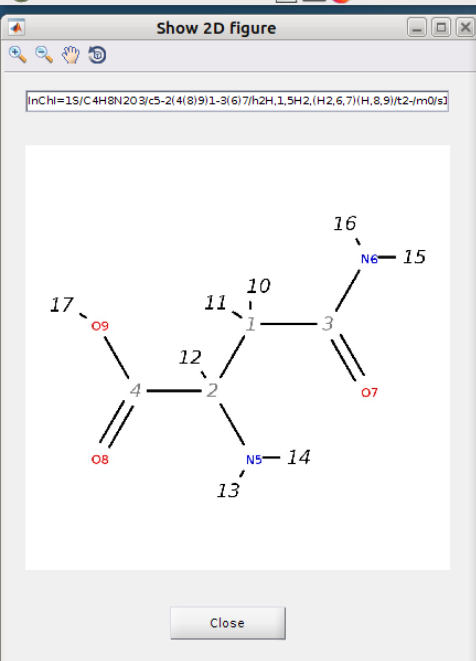
Swap two cells

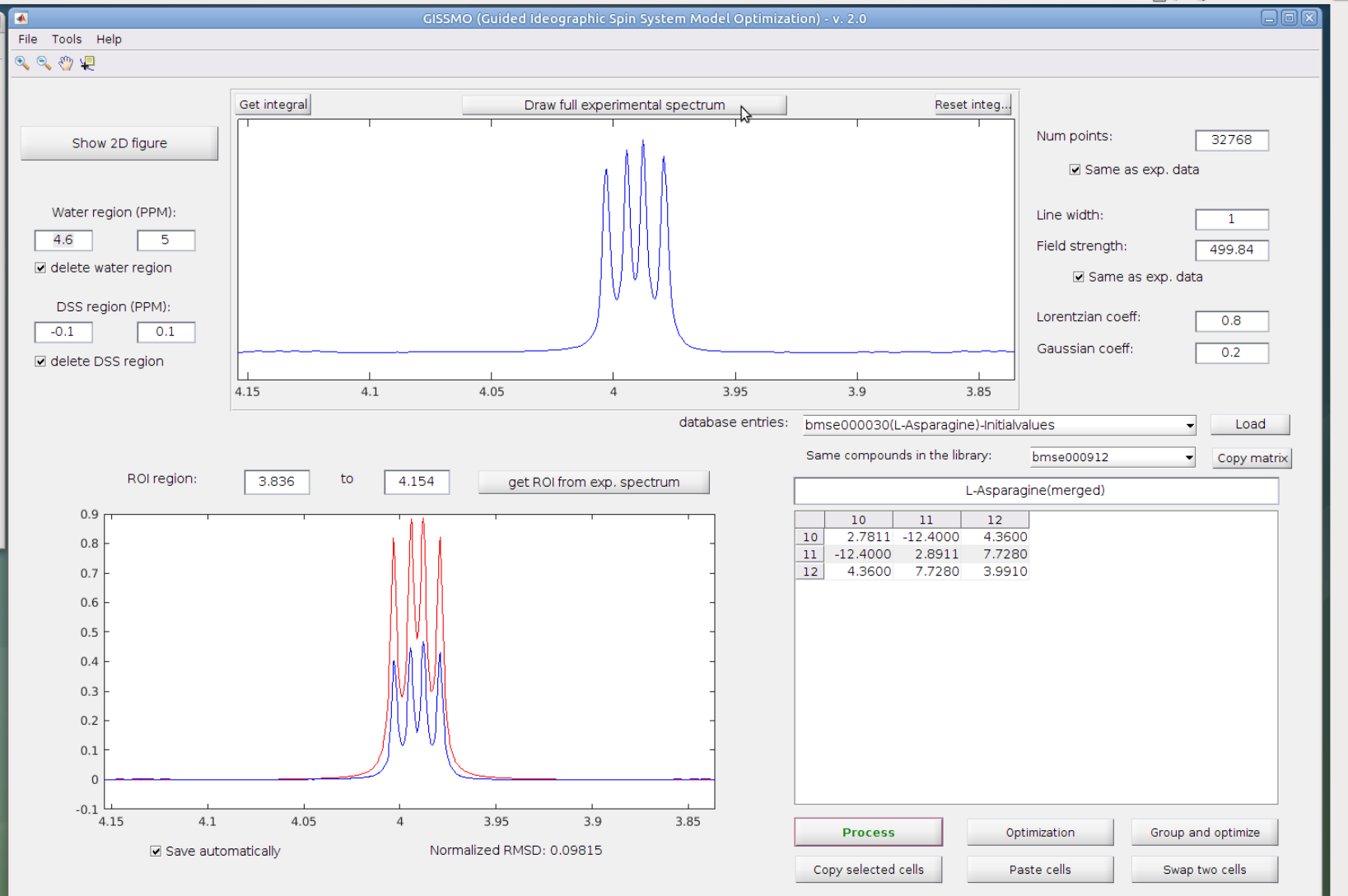
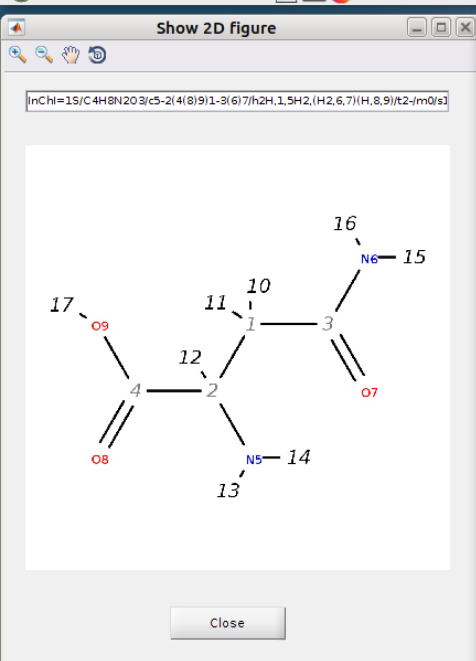
L-Asparagine(merged)

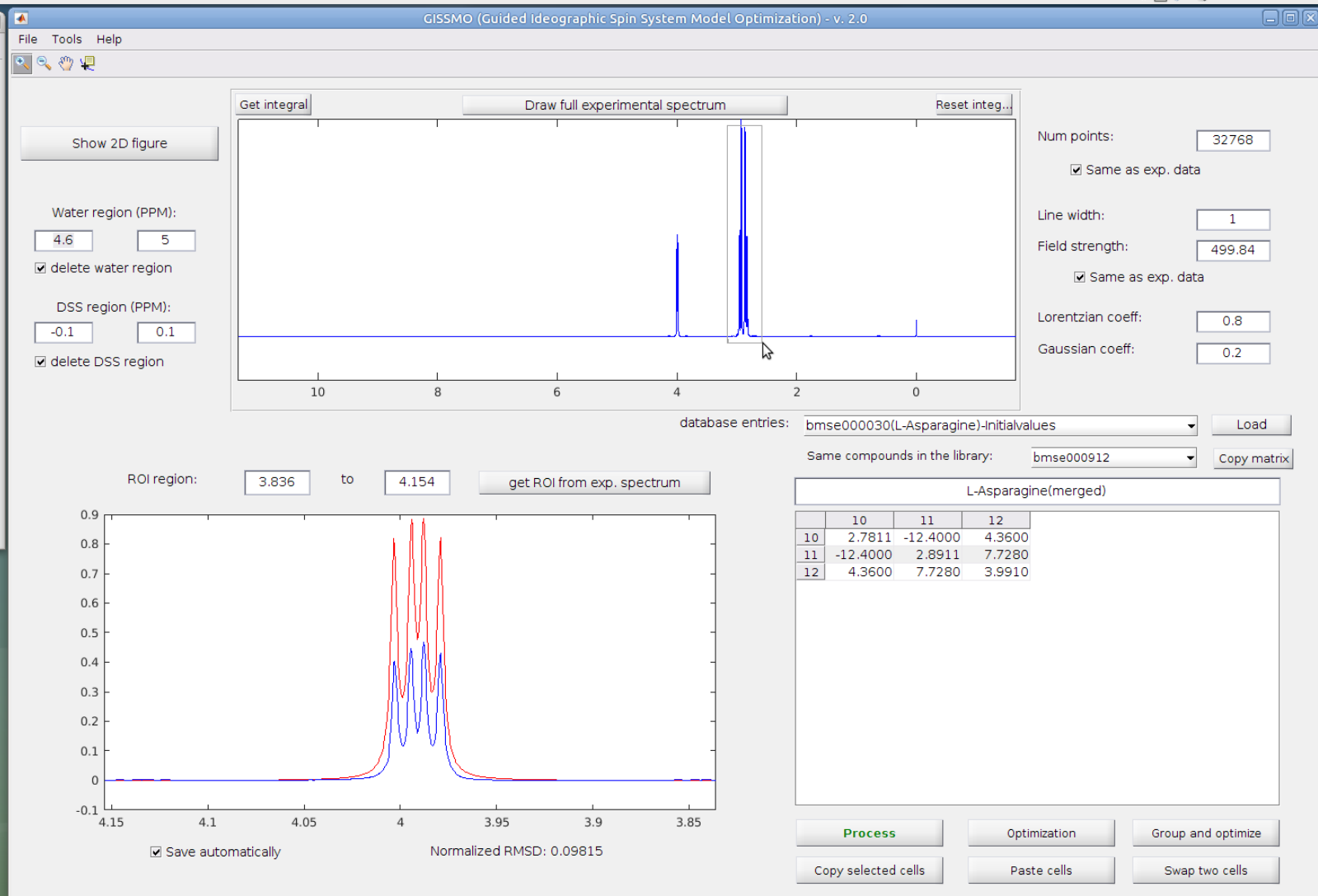
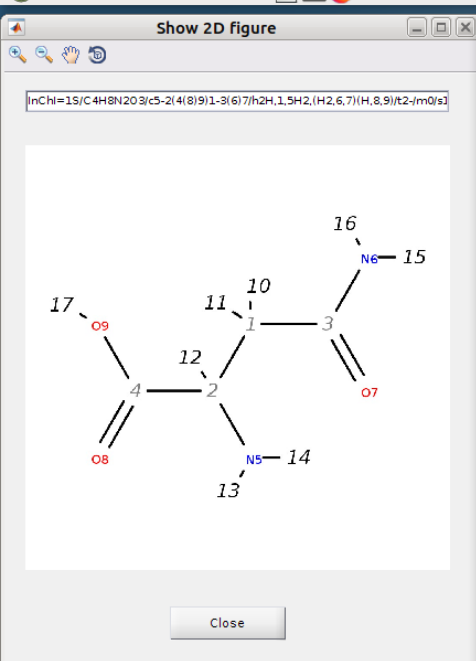
	10	11	12
10	2.7811	-12.4000	4.3600
11	-12.4000	2.8911	6.4963
12	4.3600	6.4963	3.9910

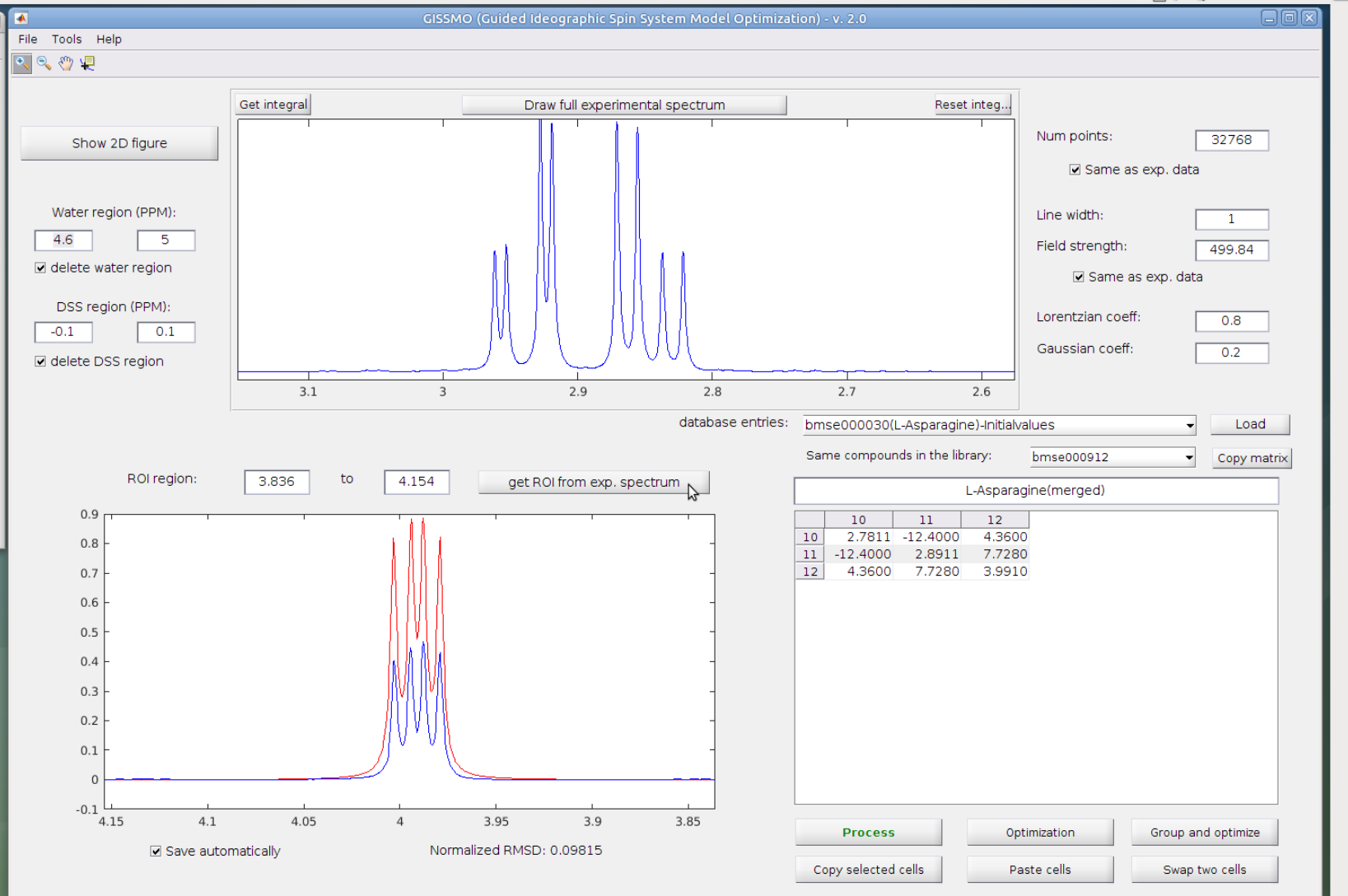
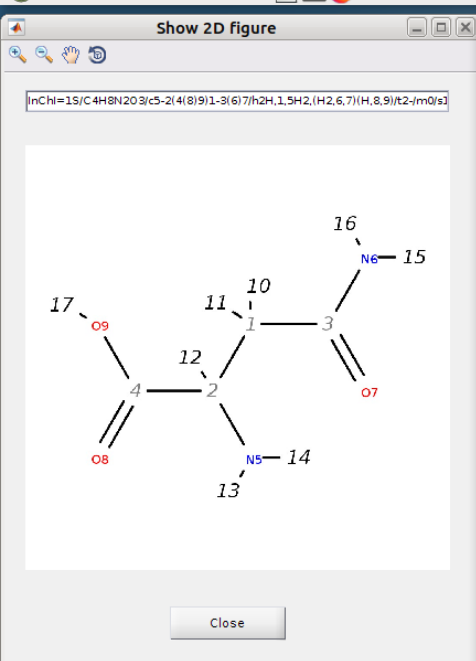


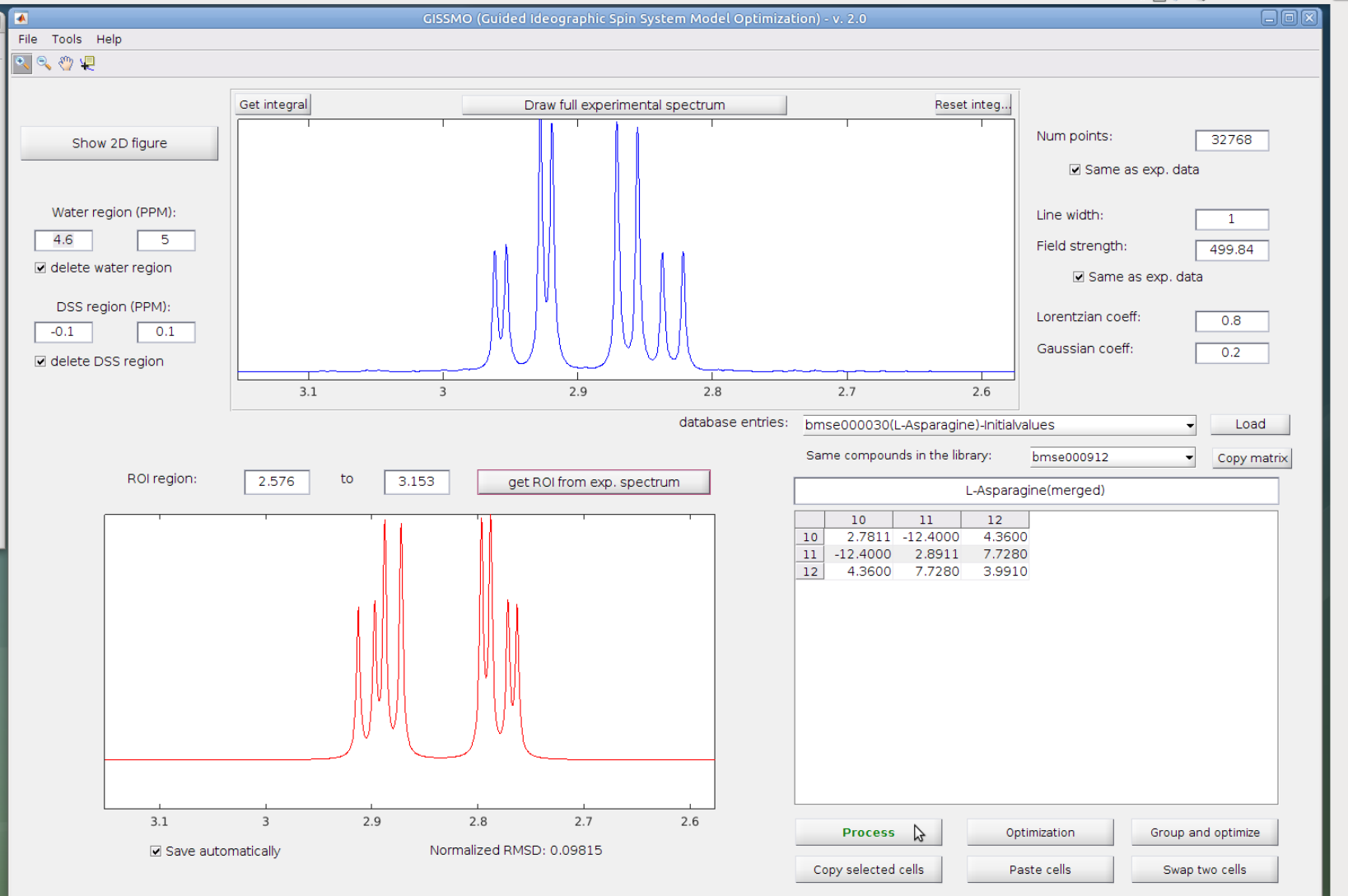
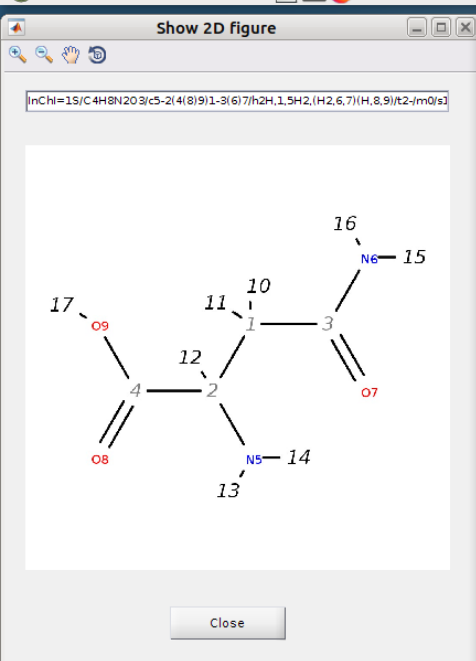


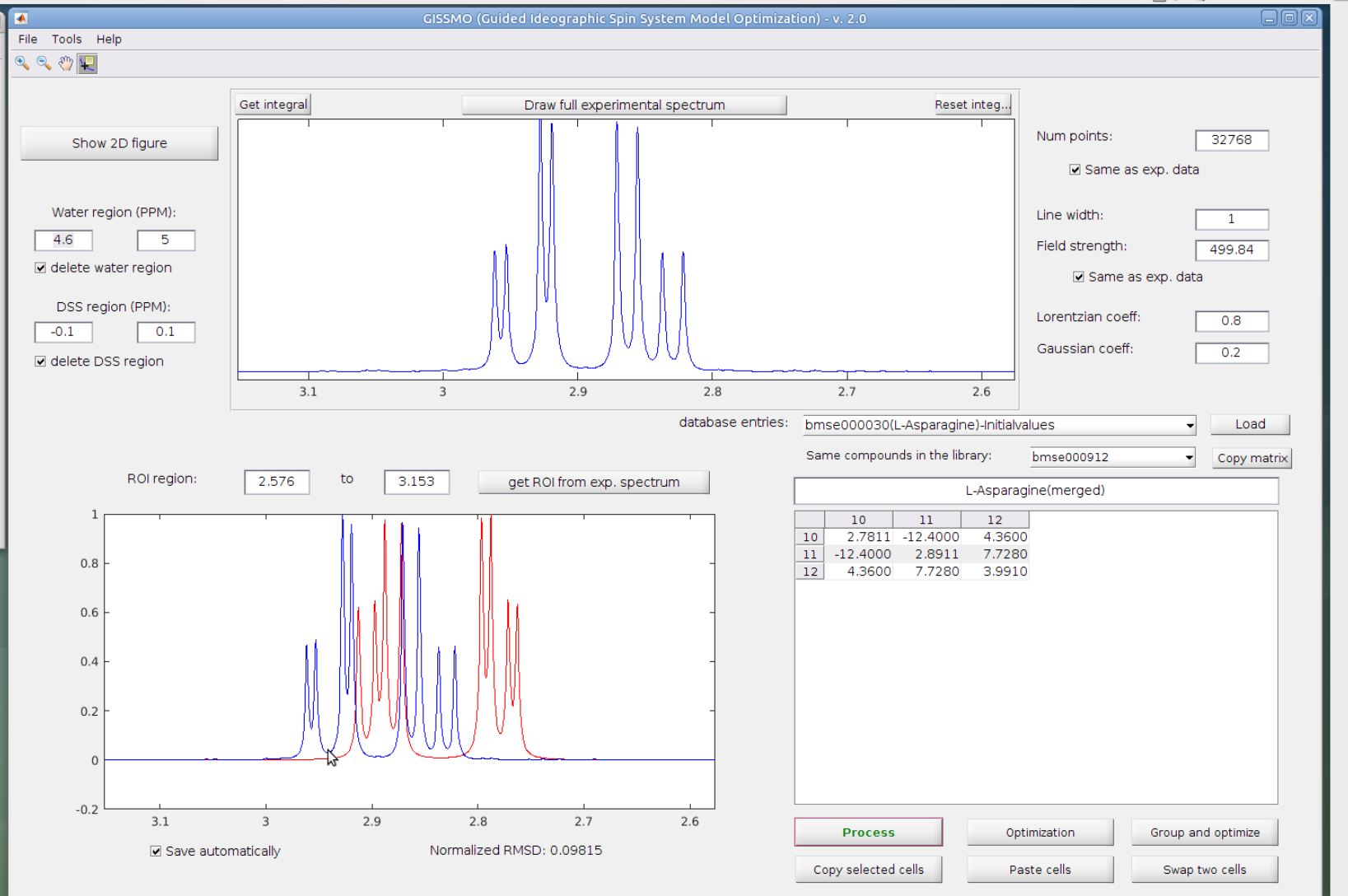
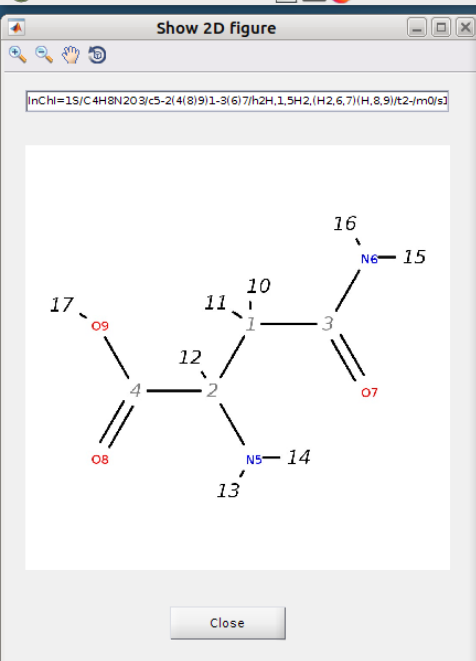


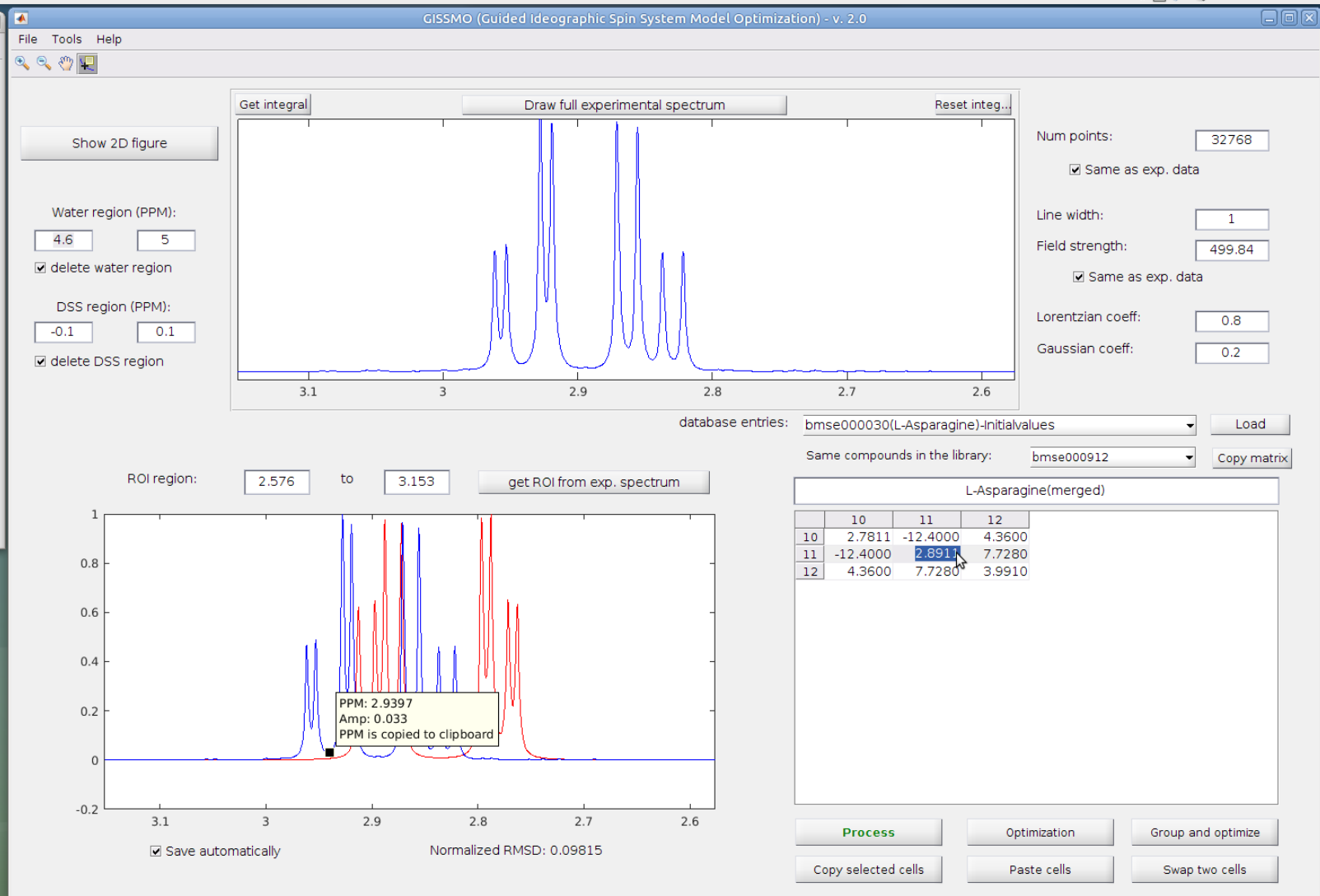
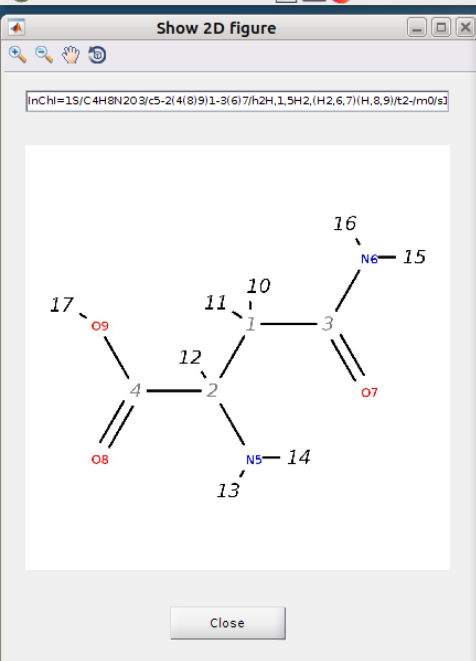


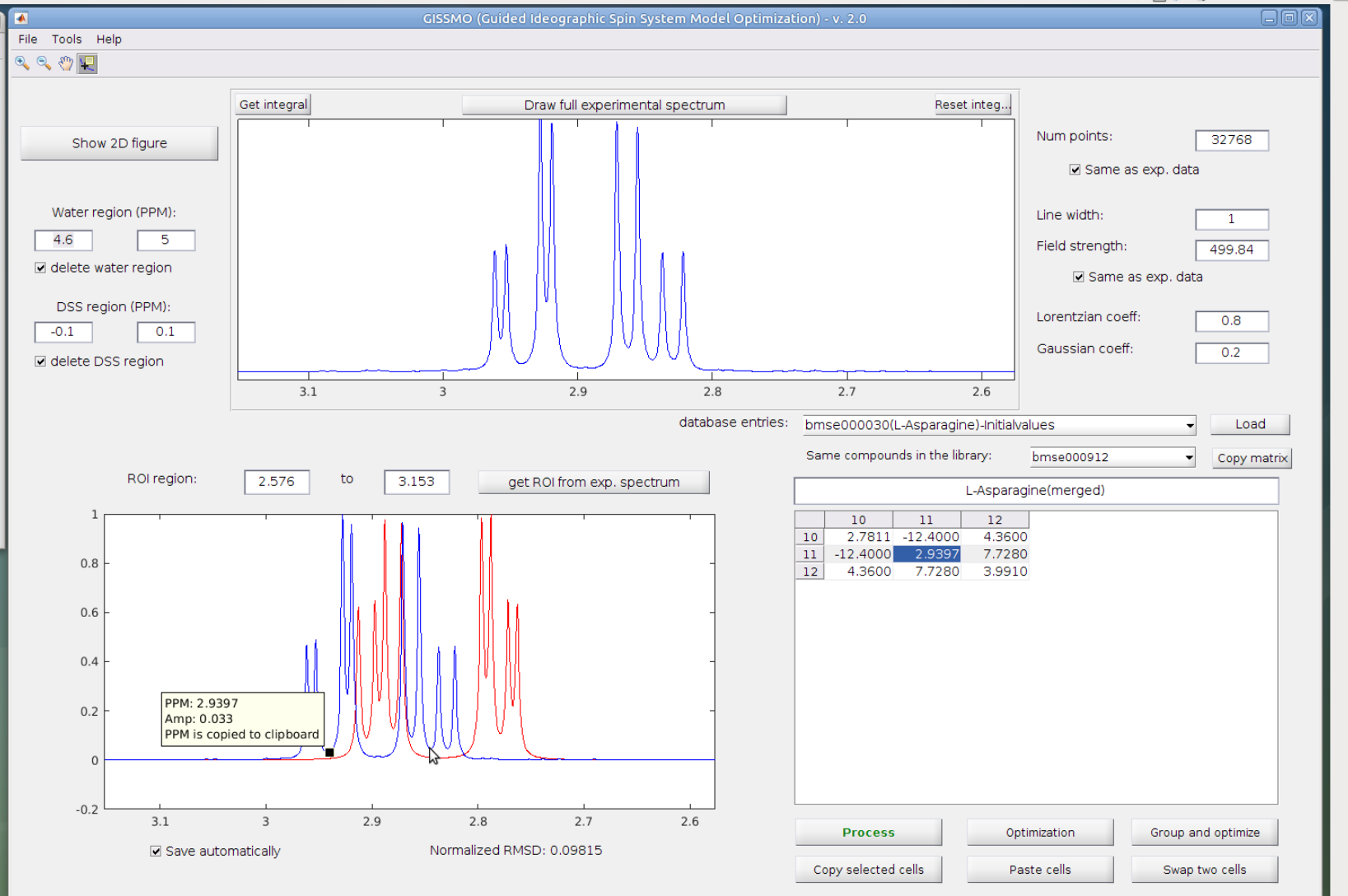
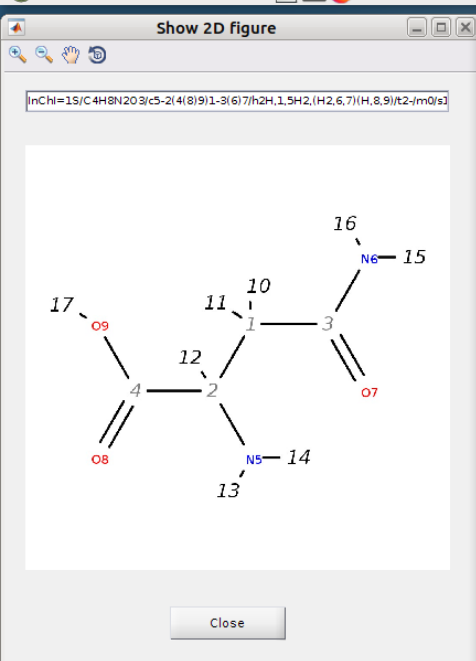


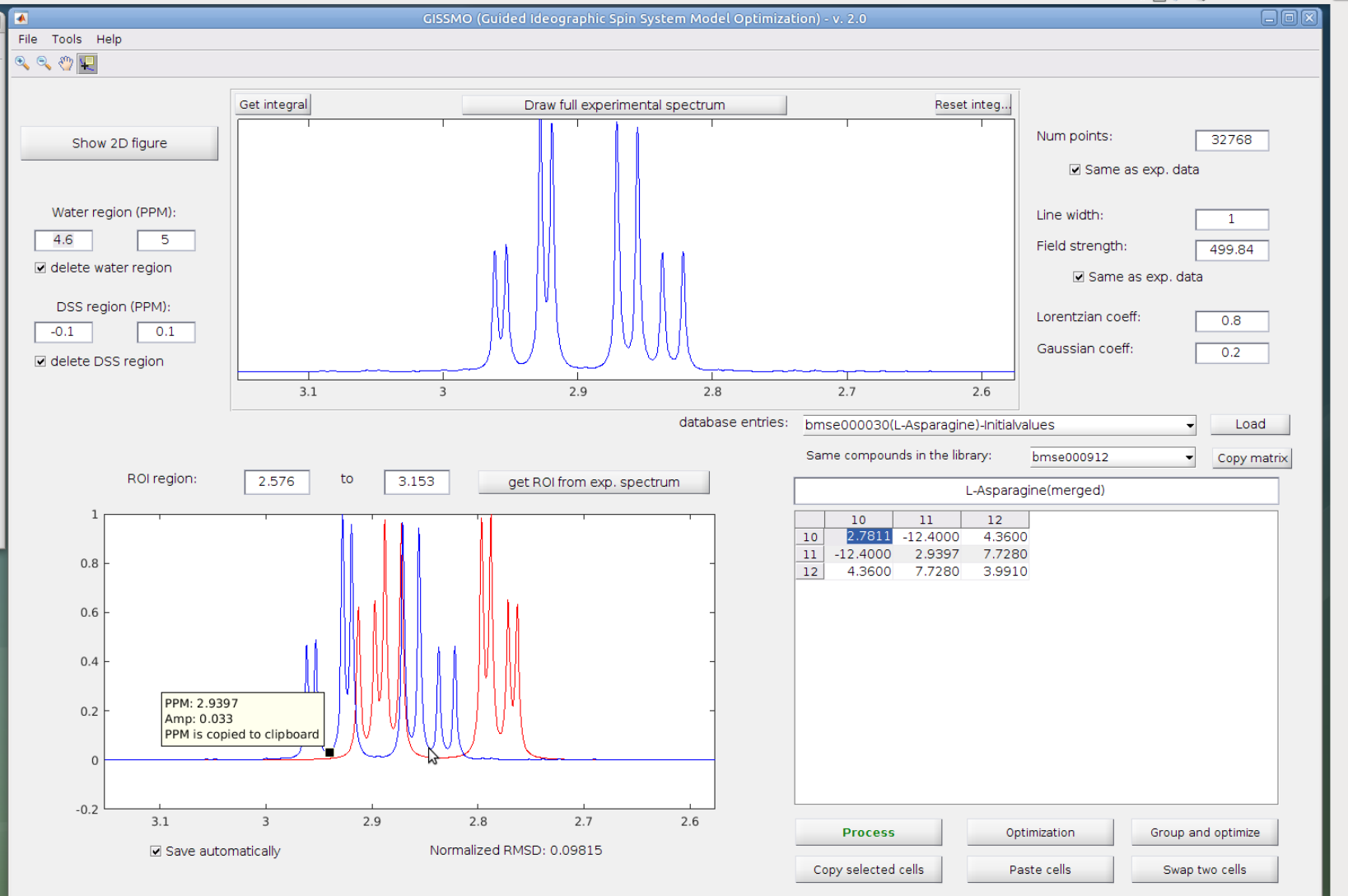
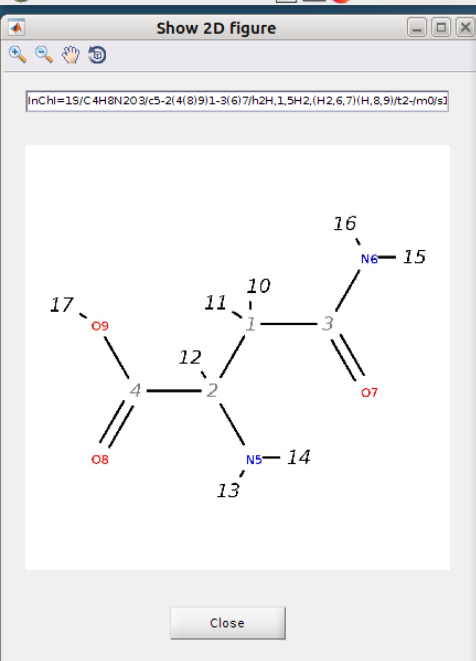


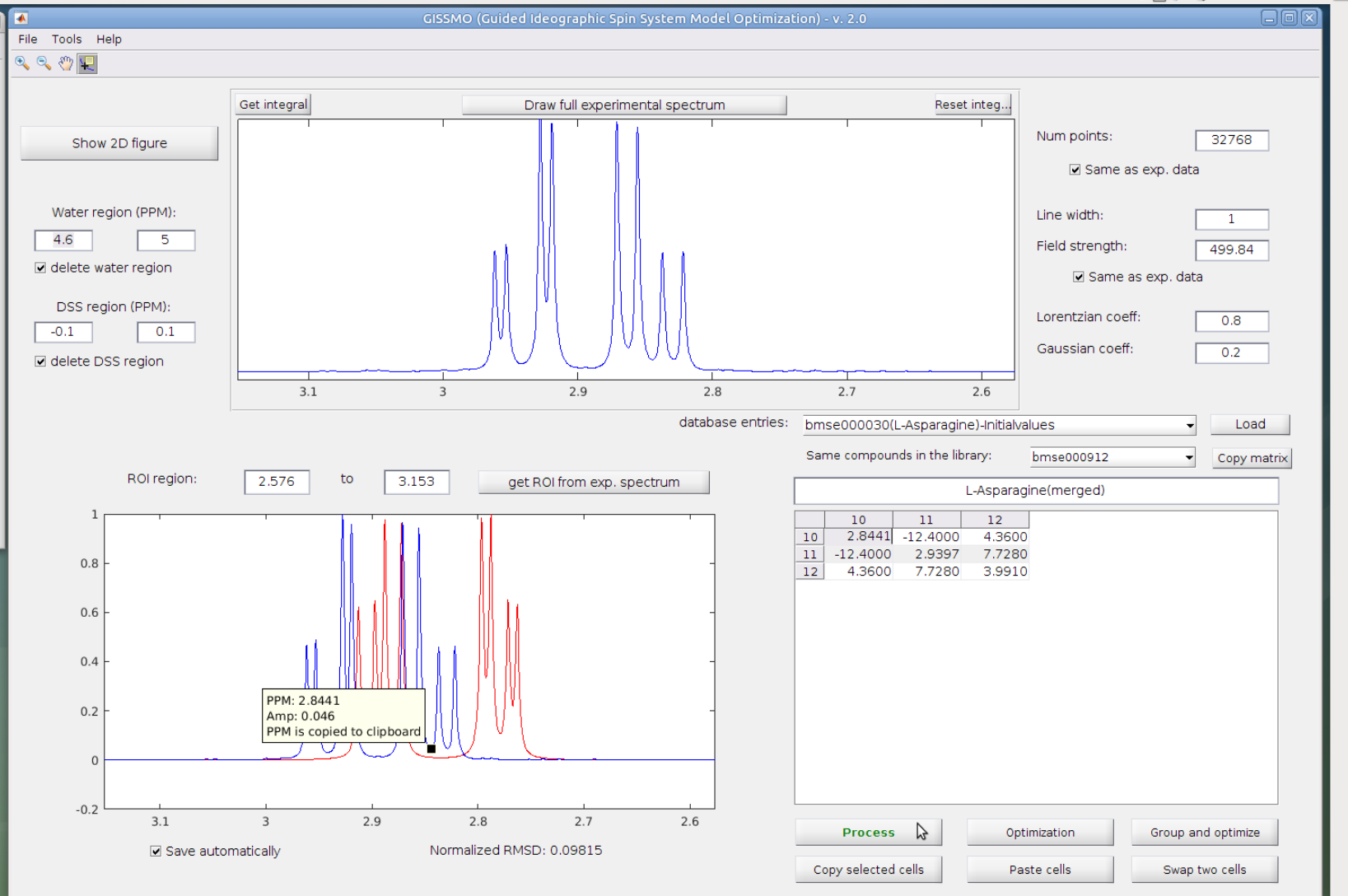
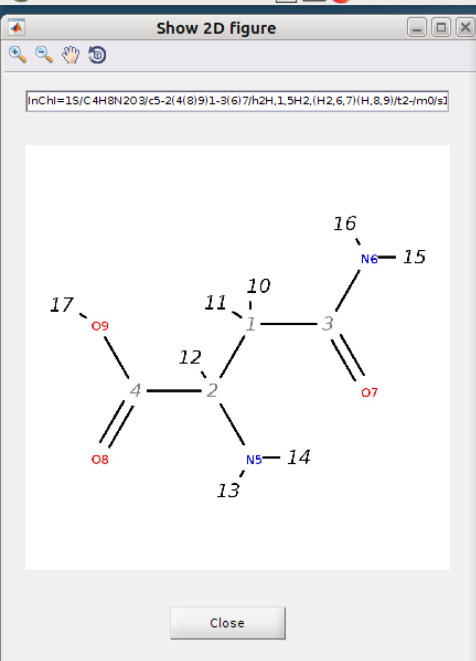


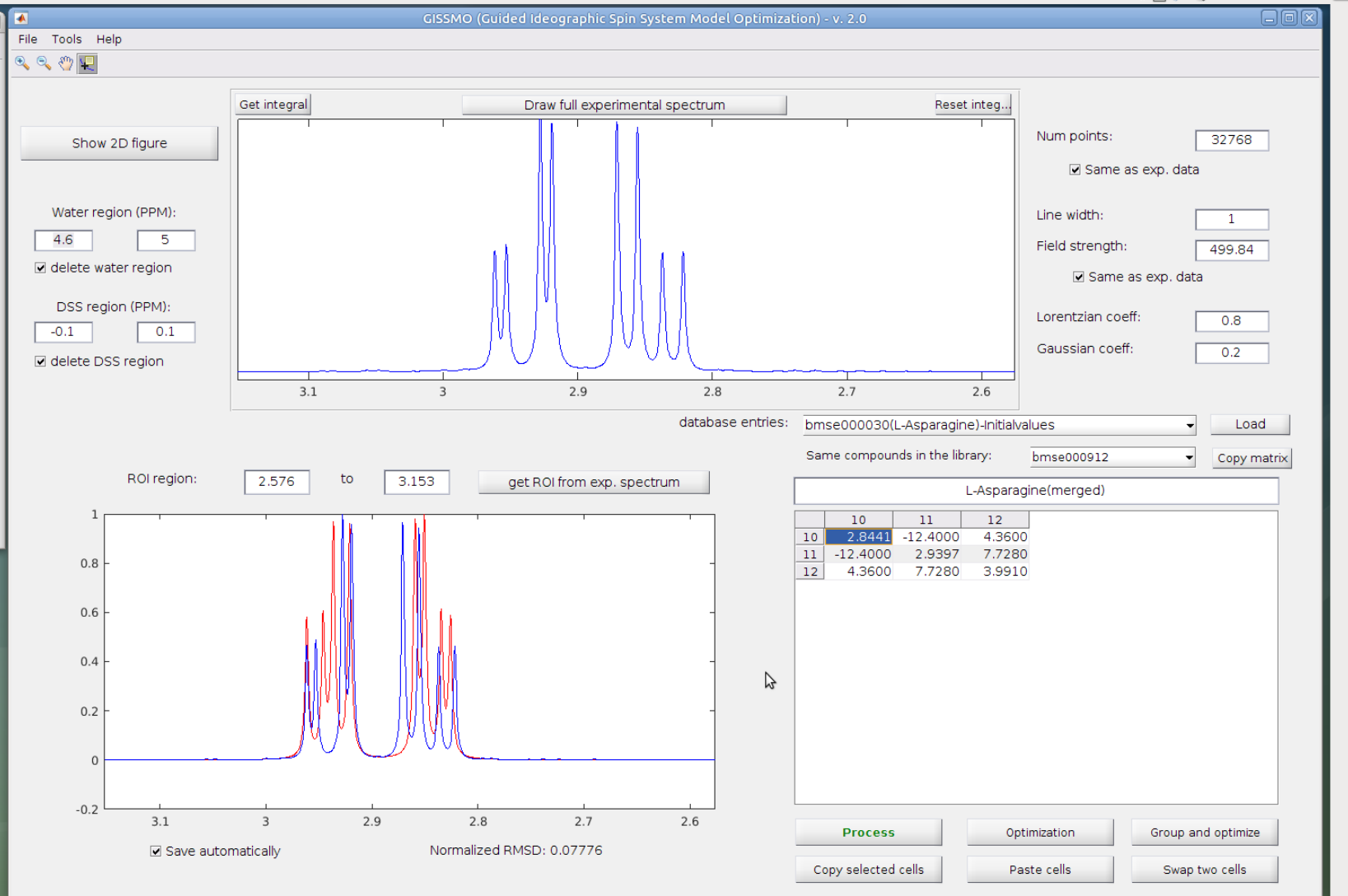
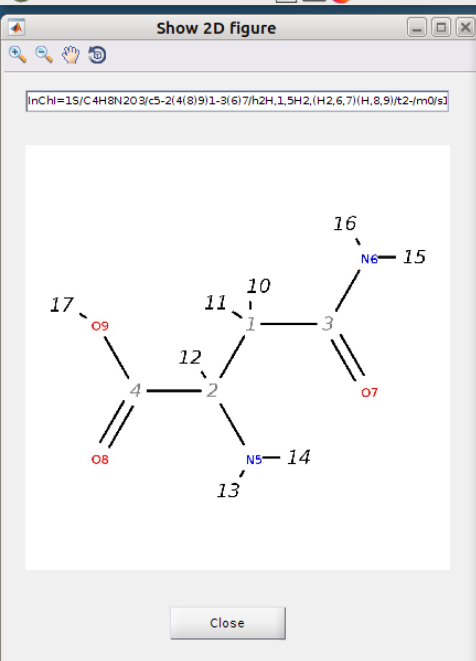


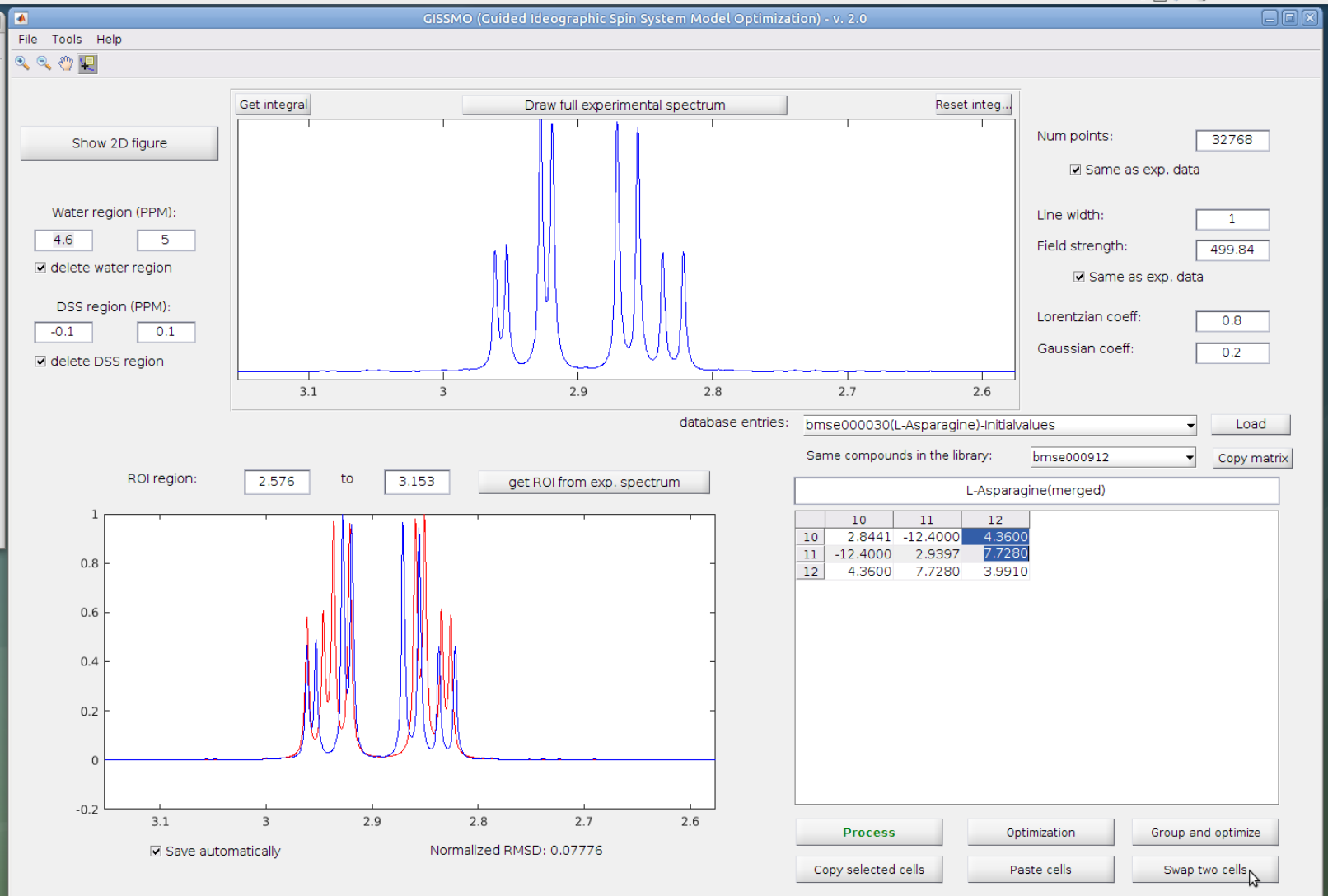
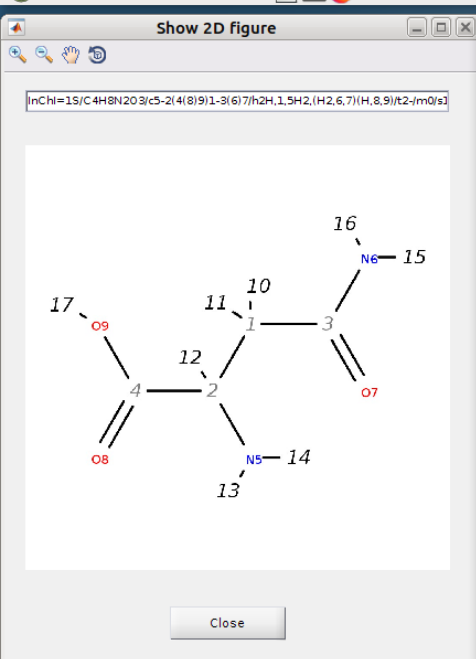


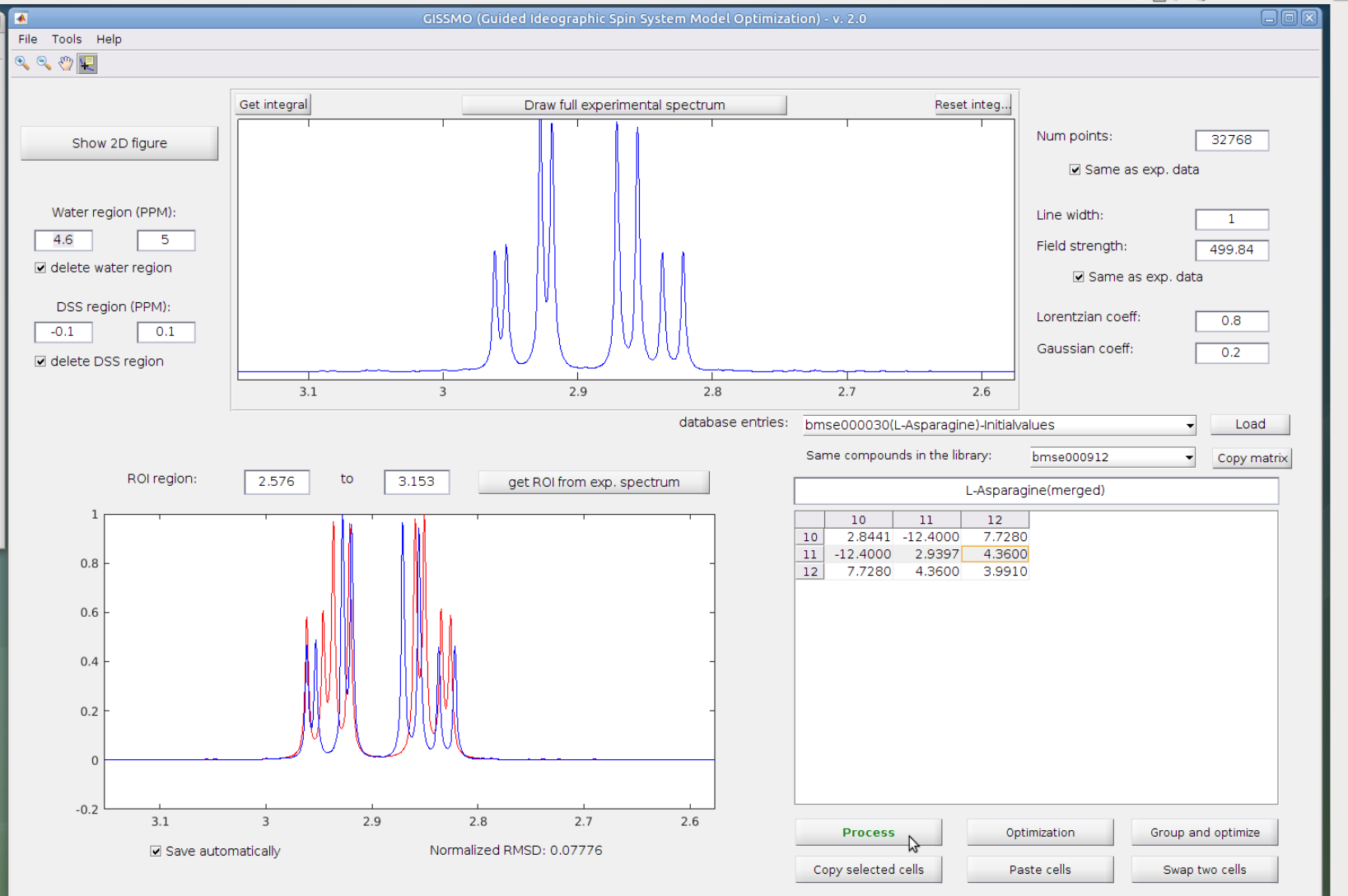
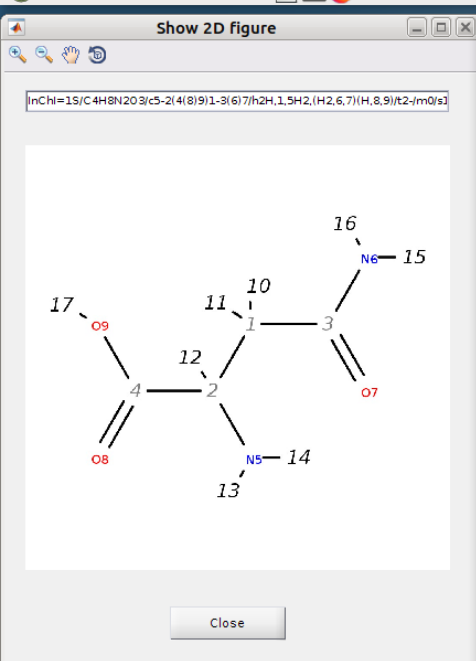


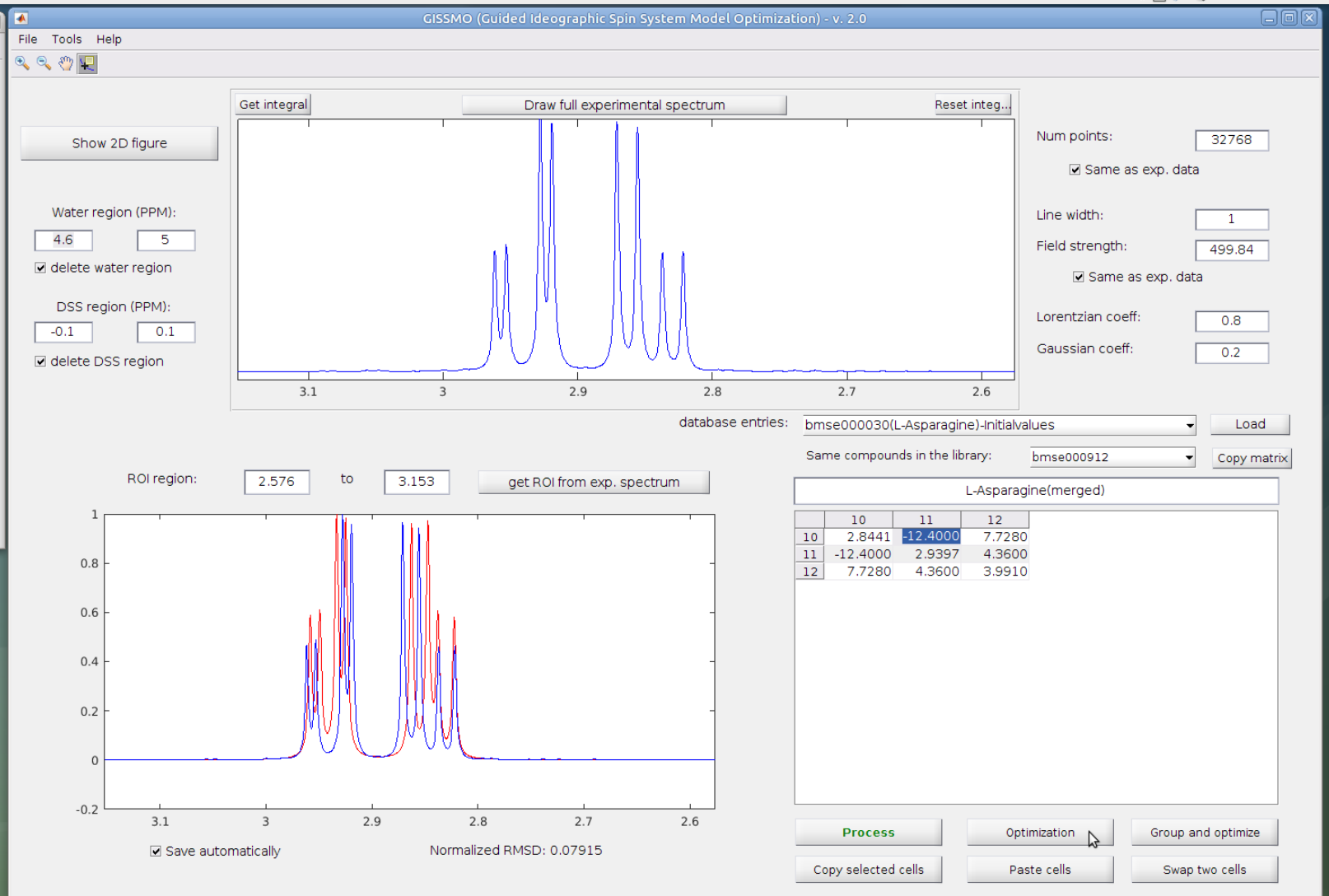
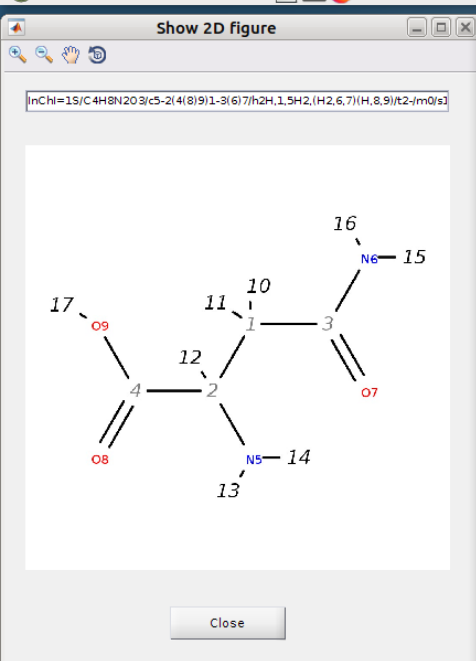


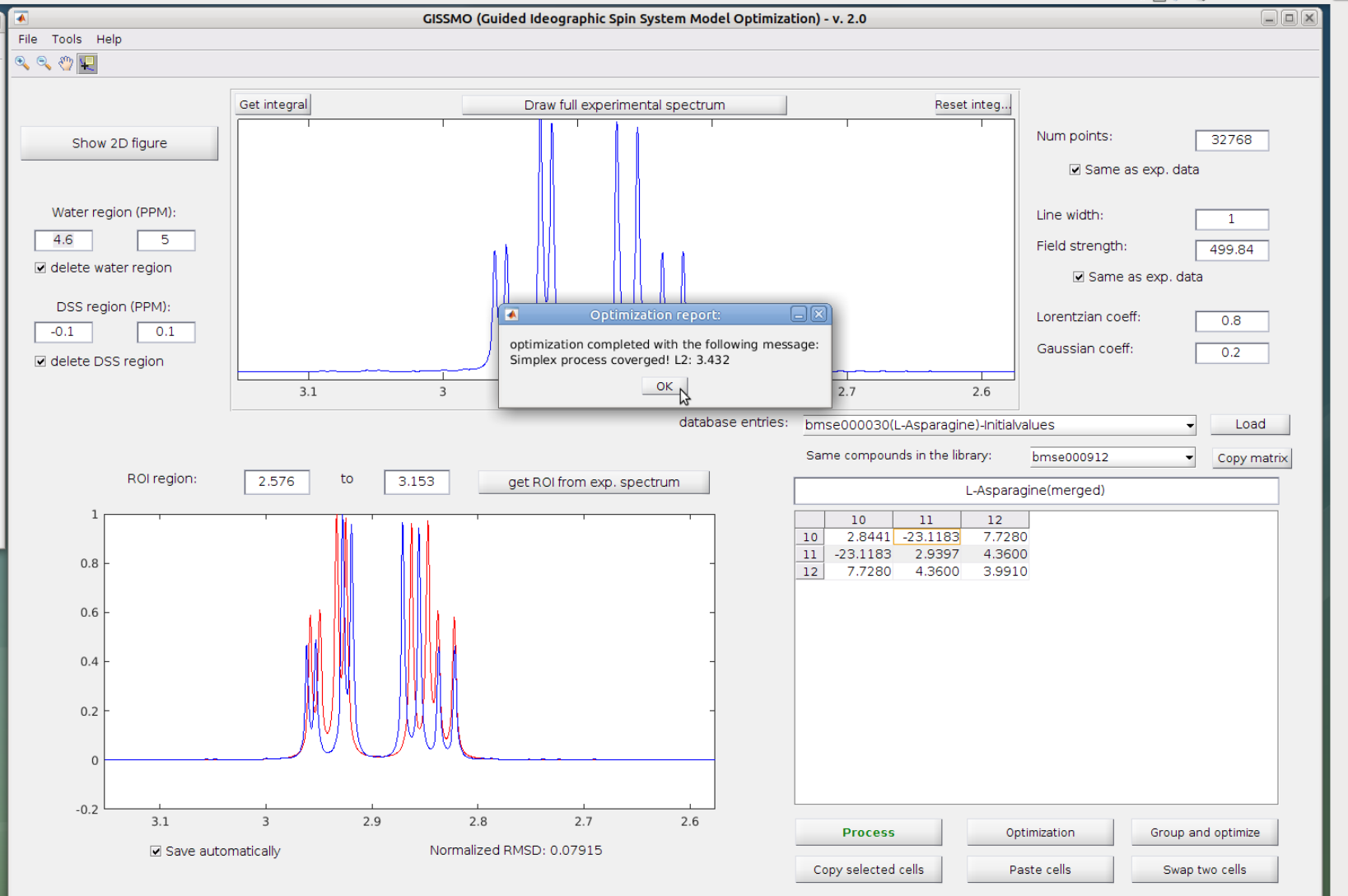
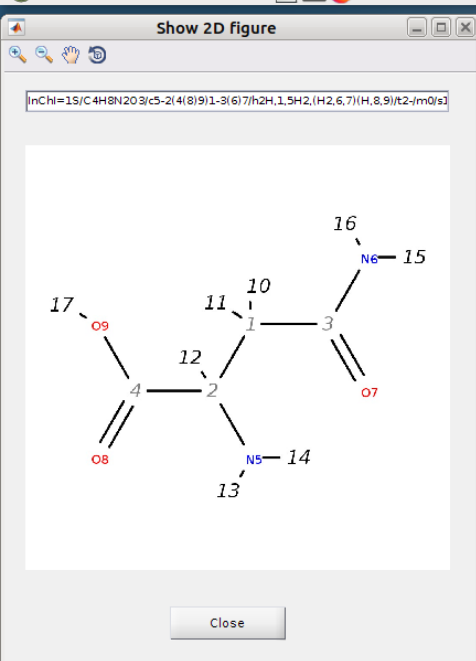


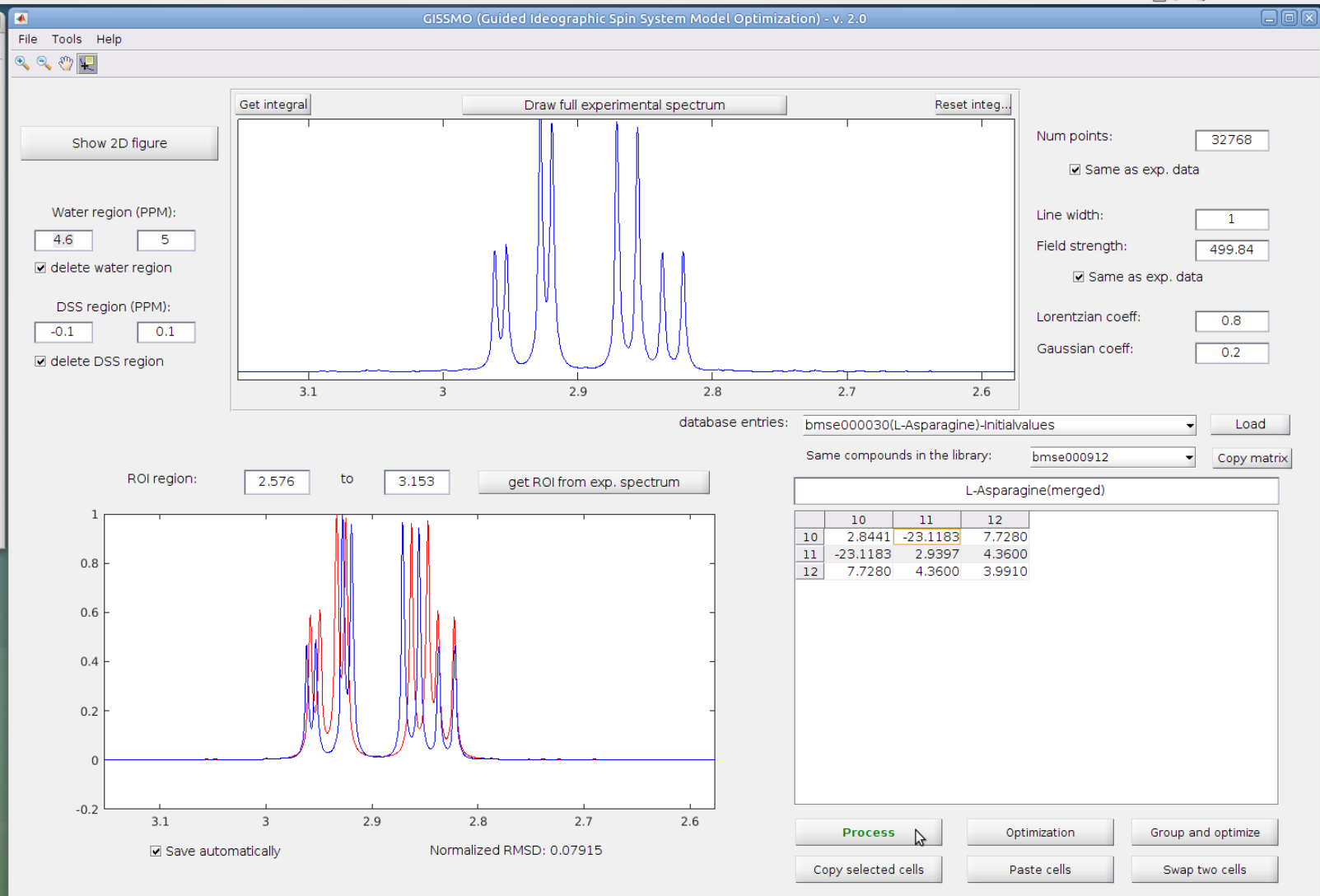
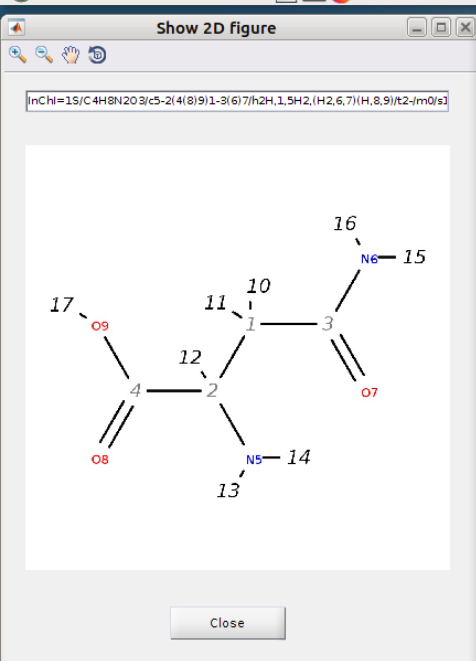


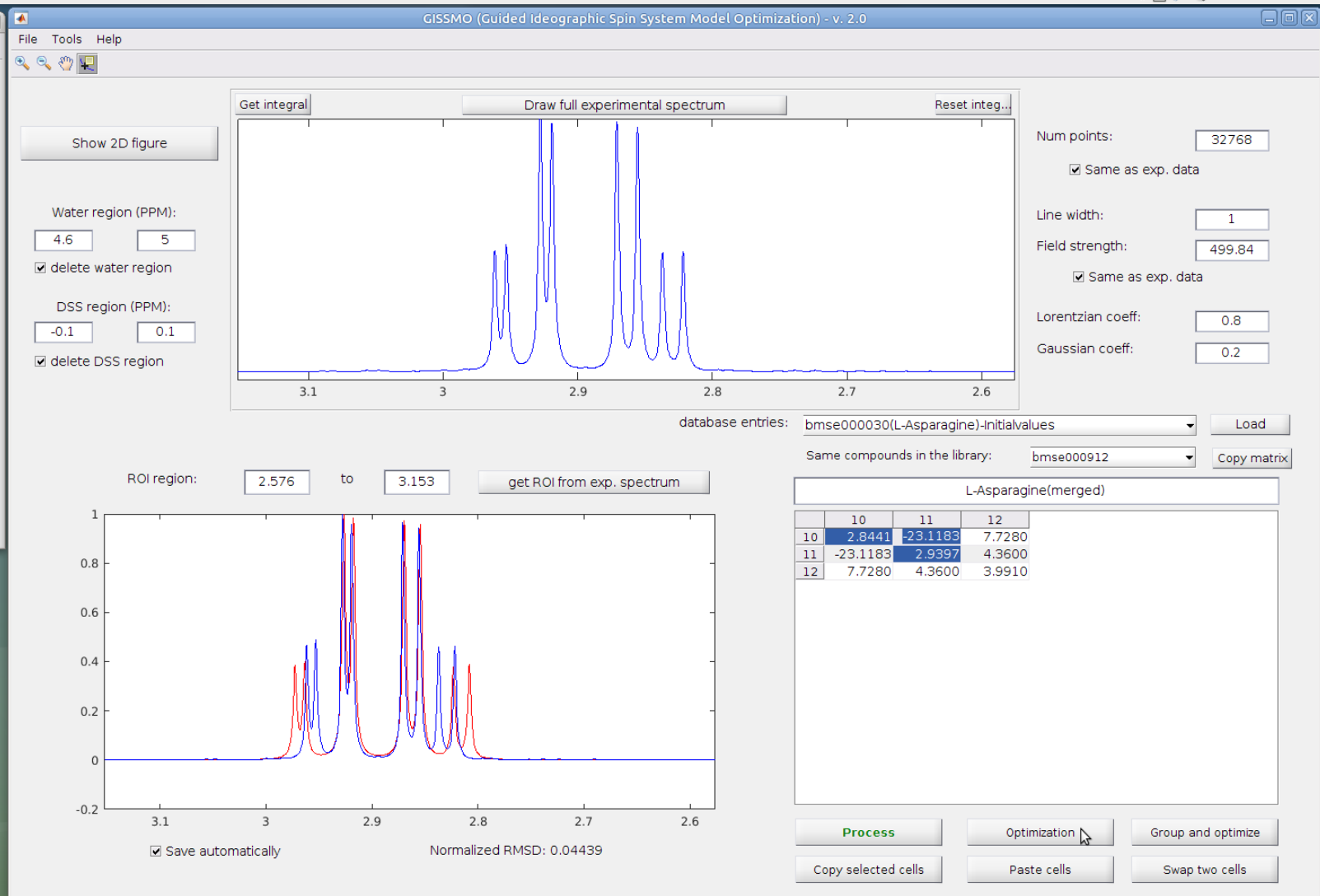
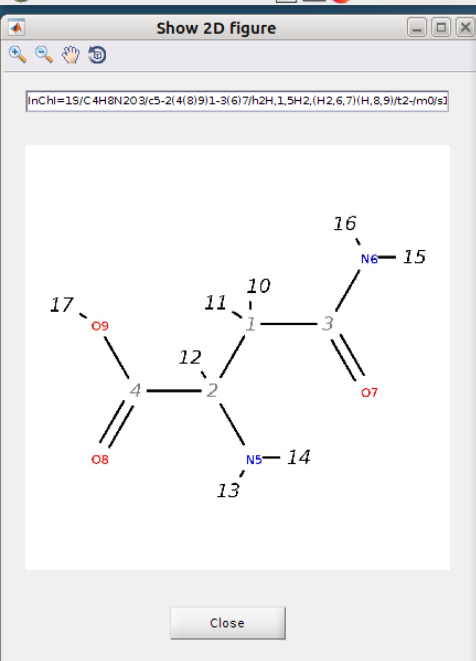


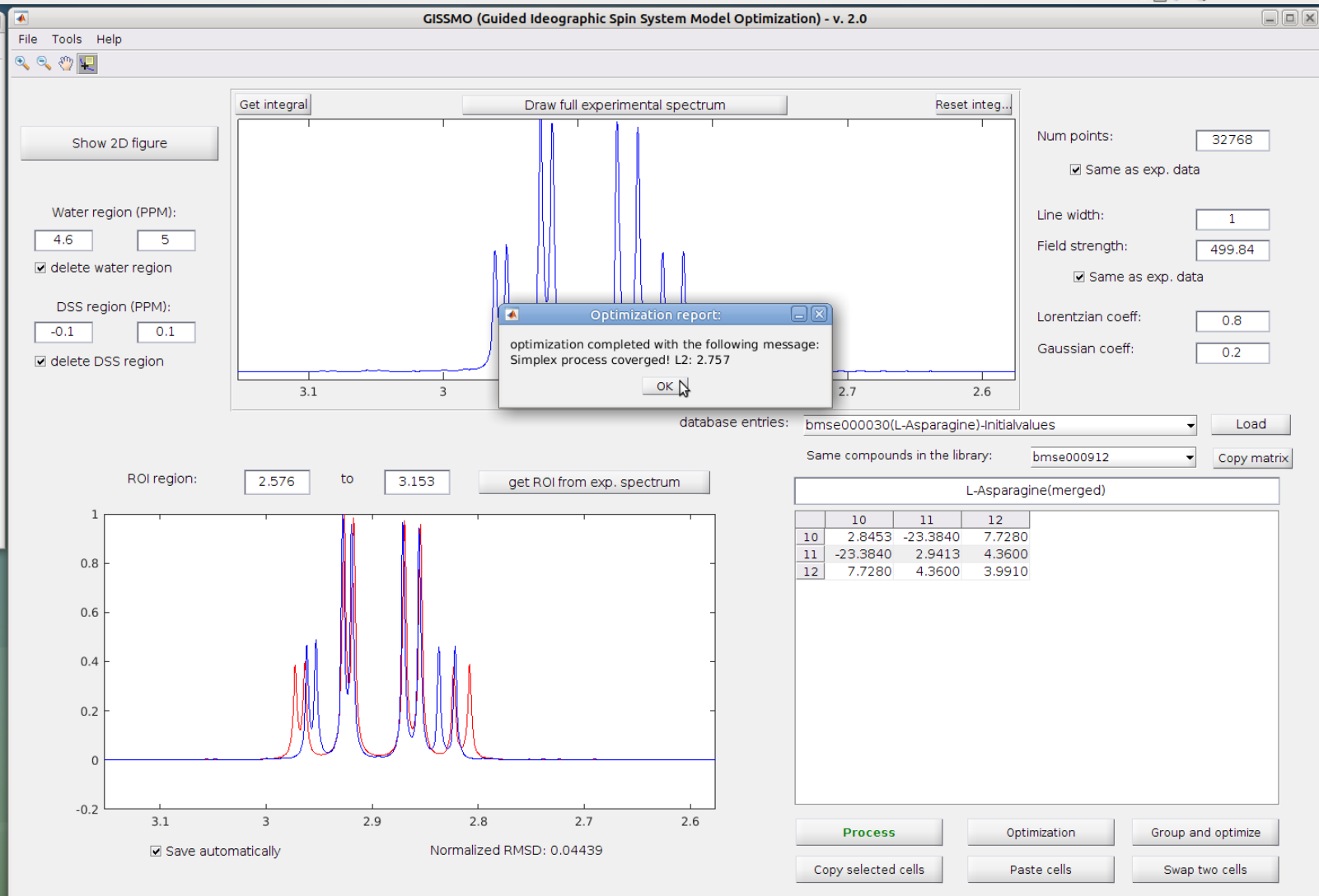
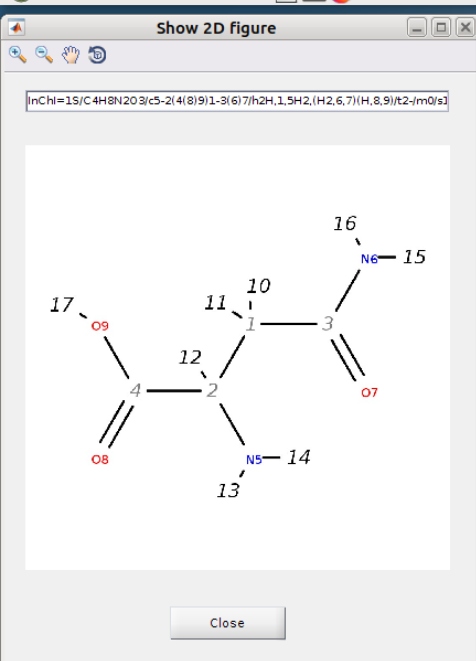


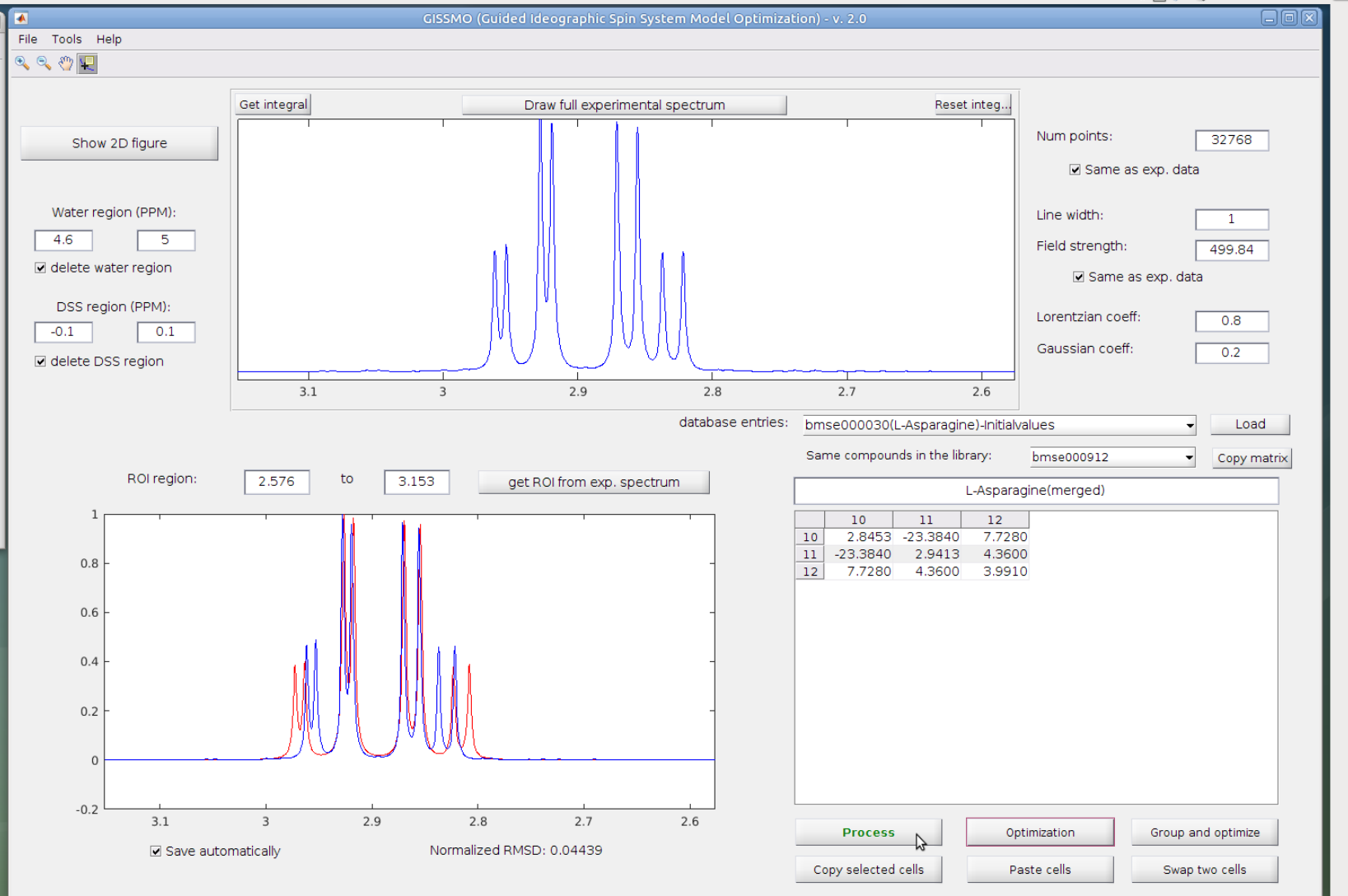
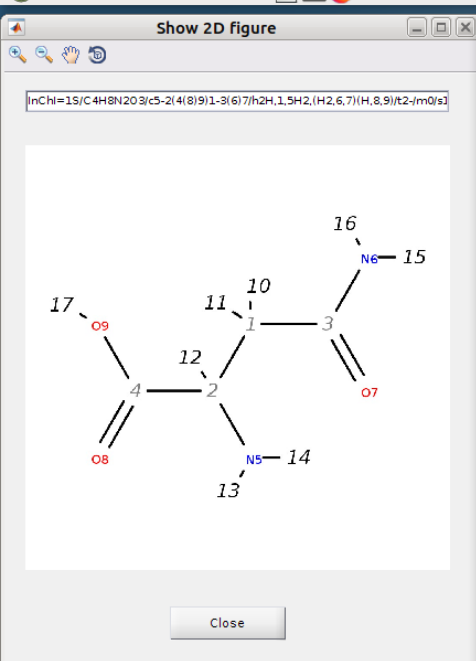


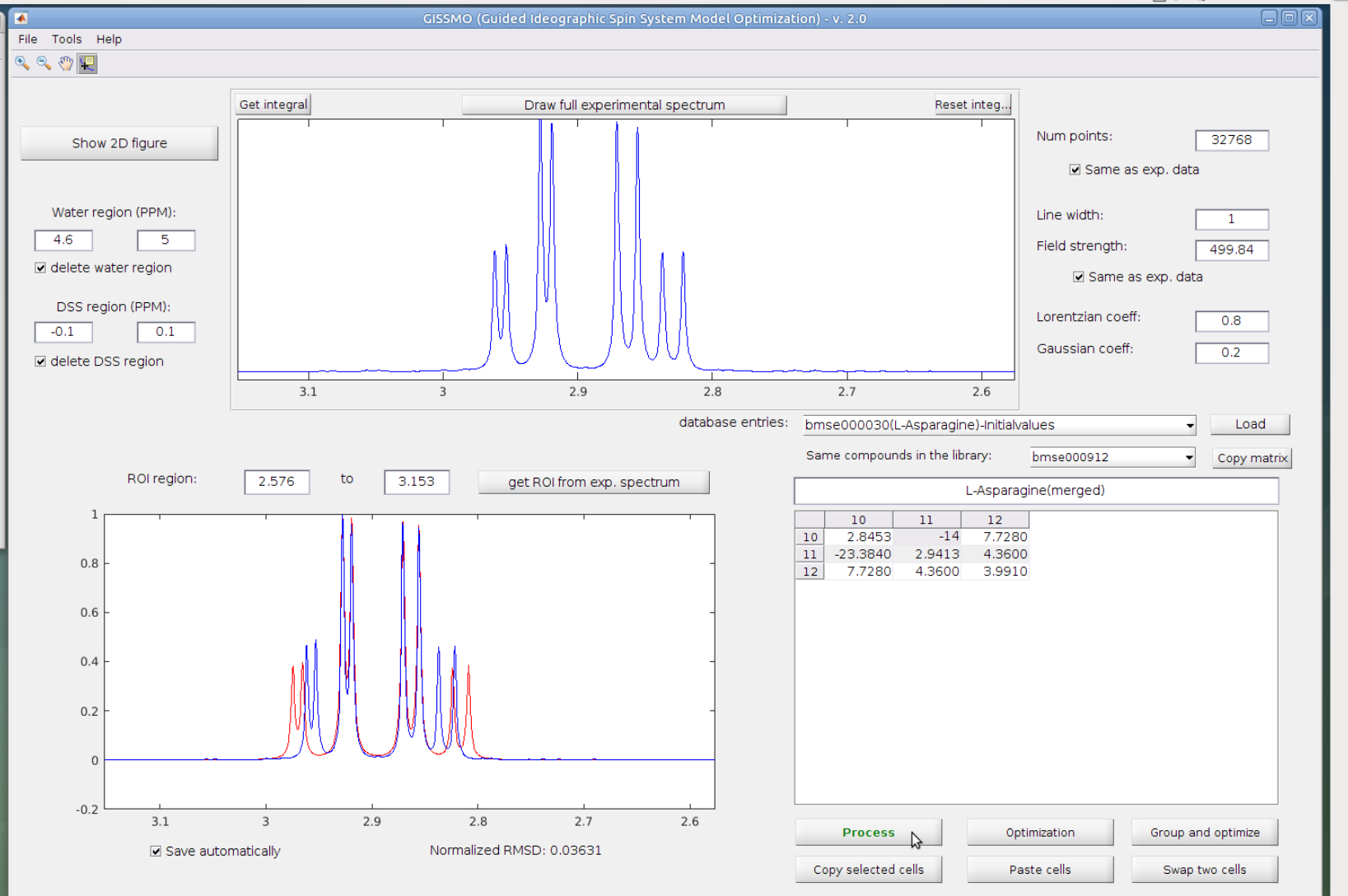
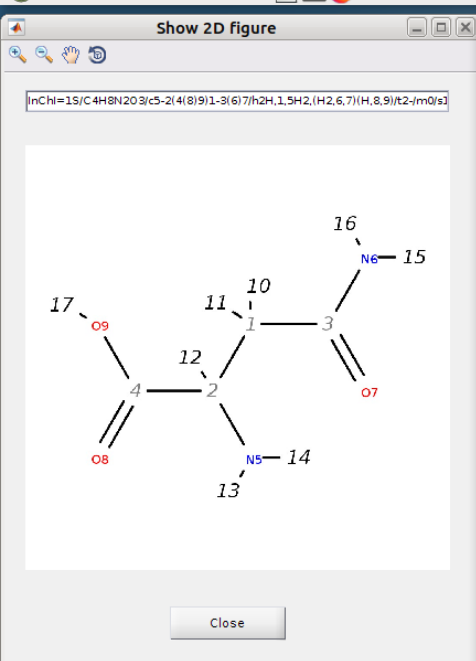


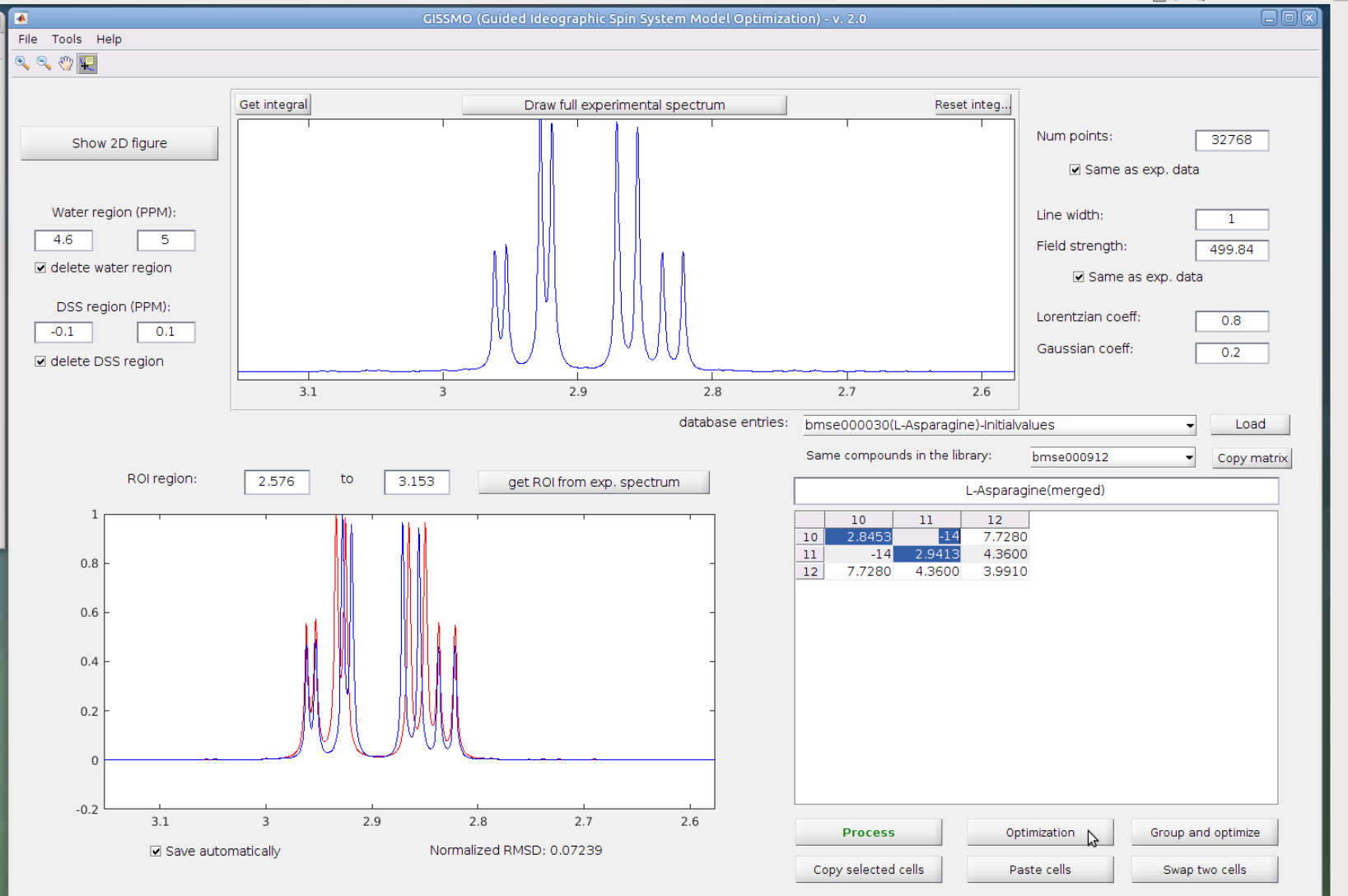
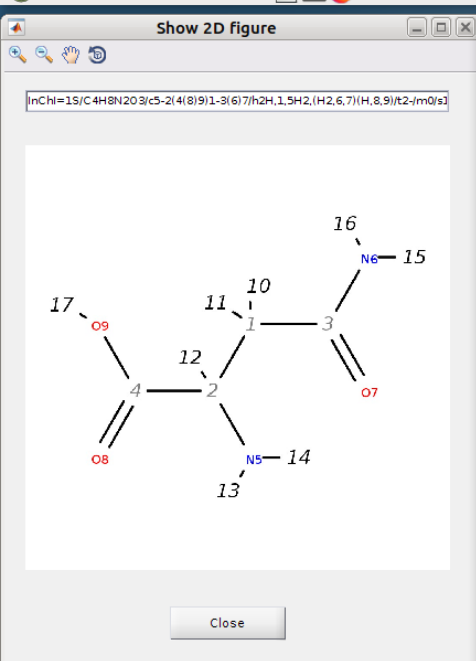


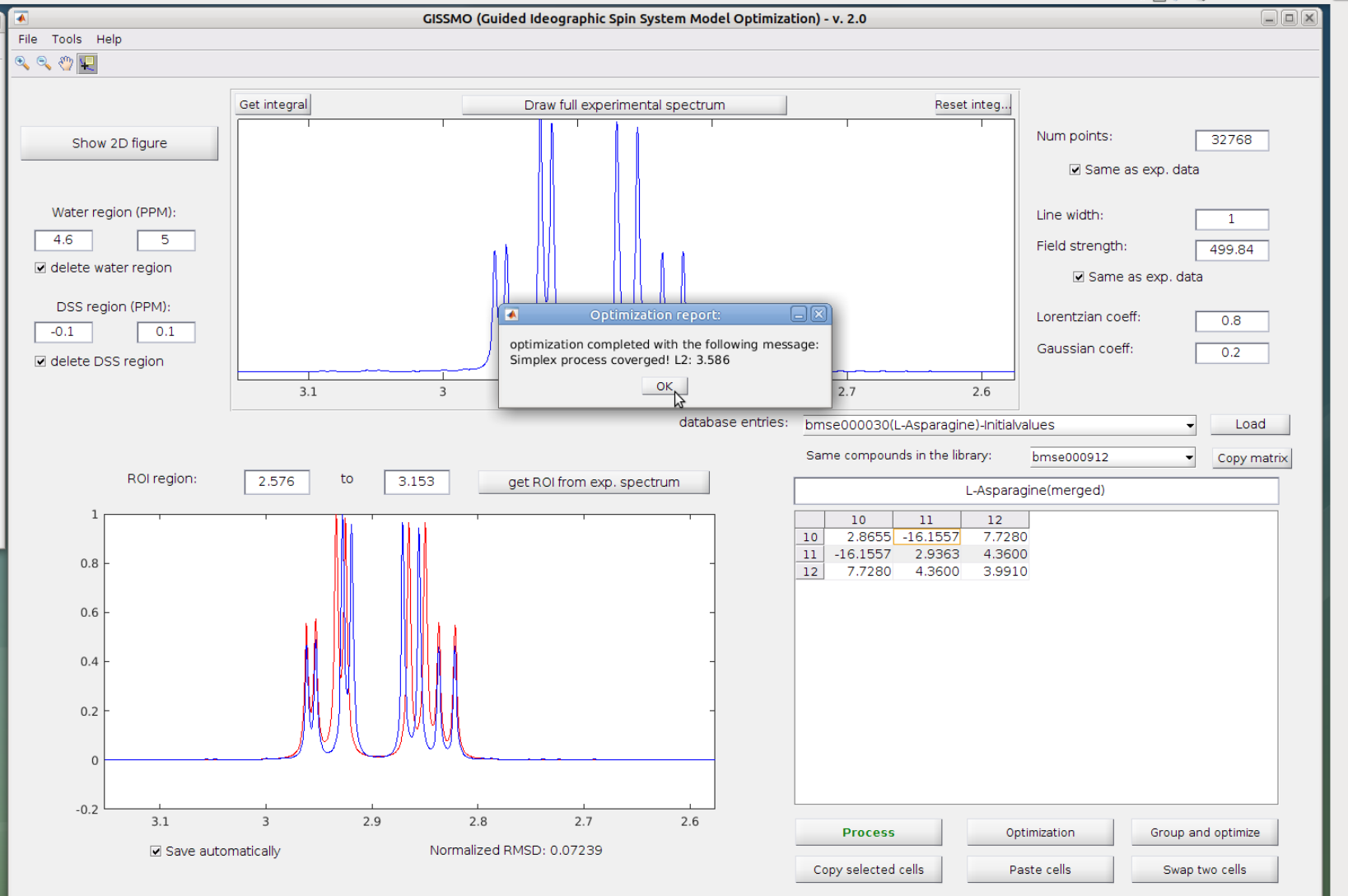
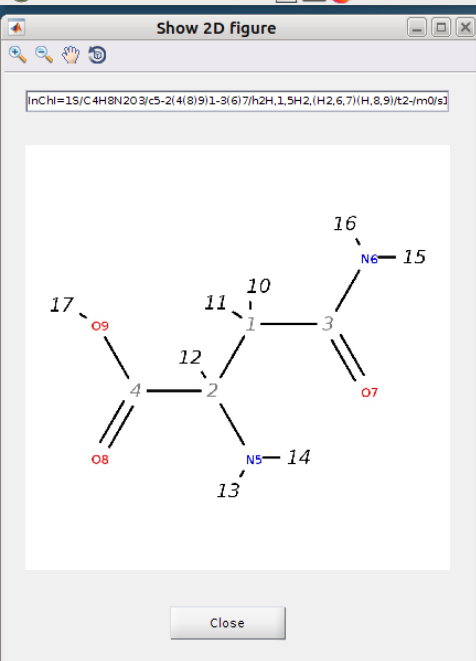


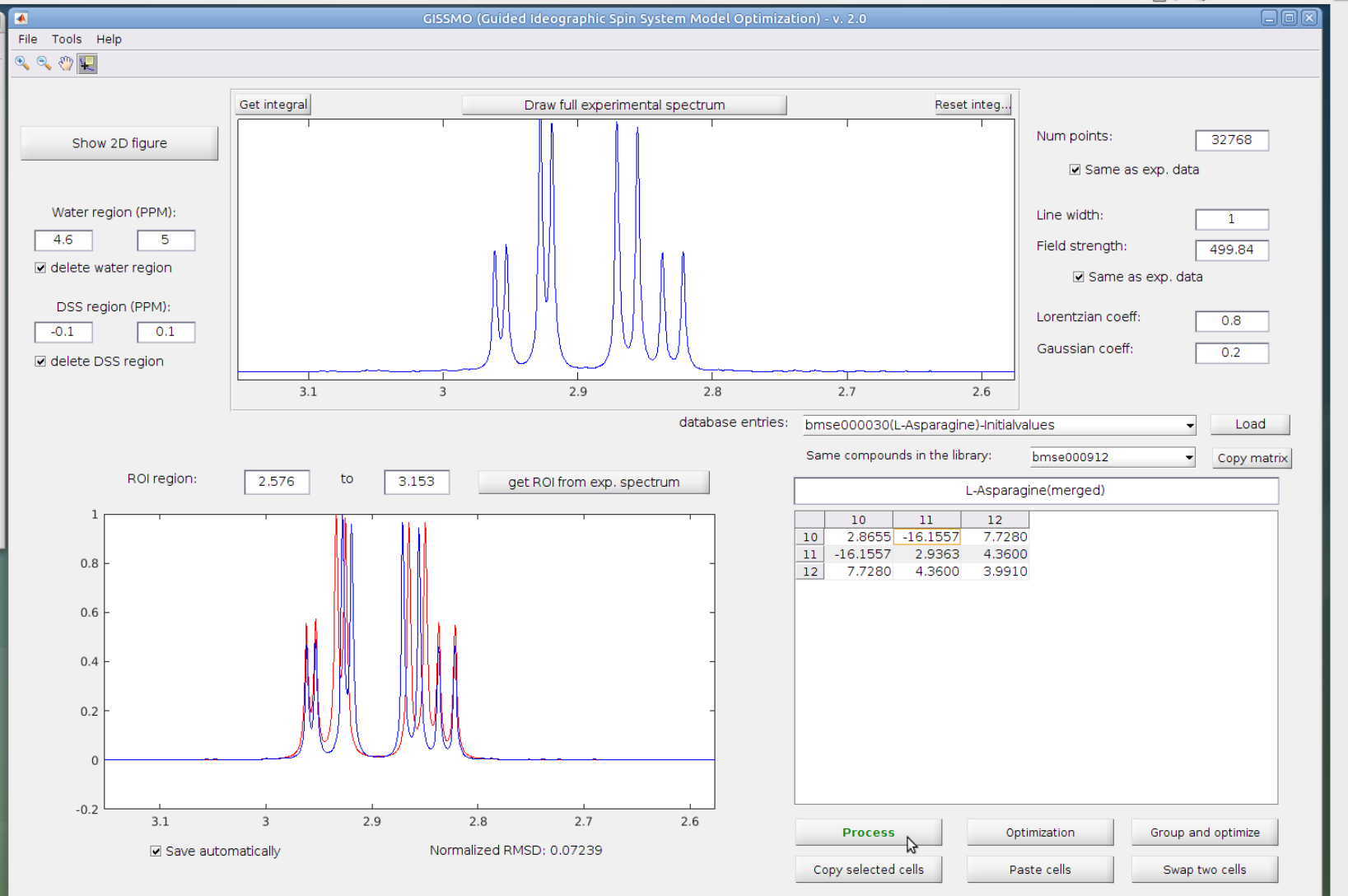
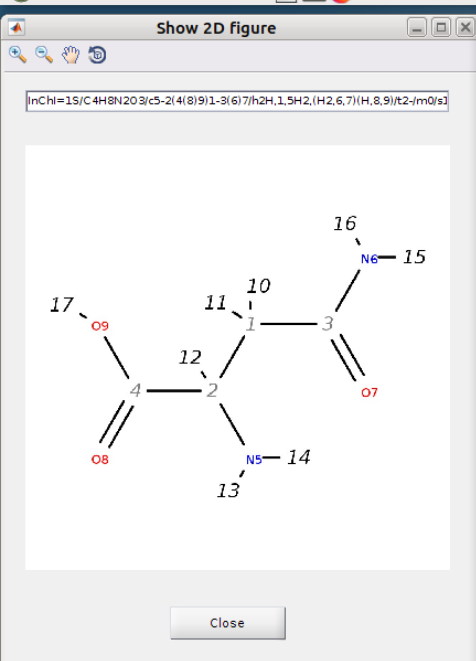


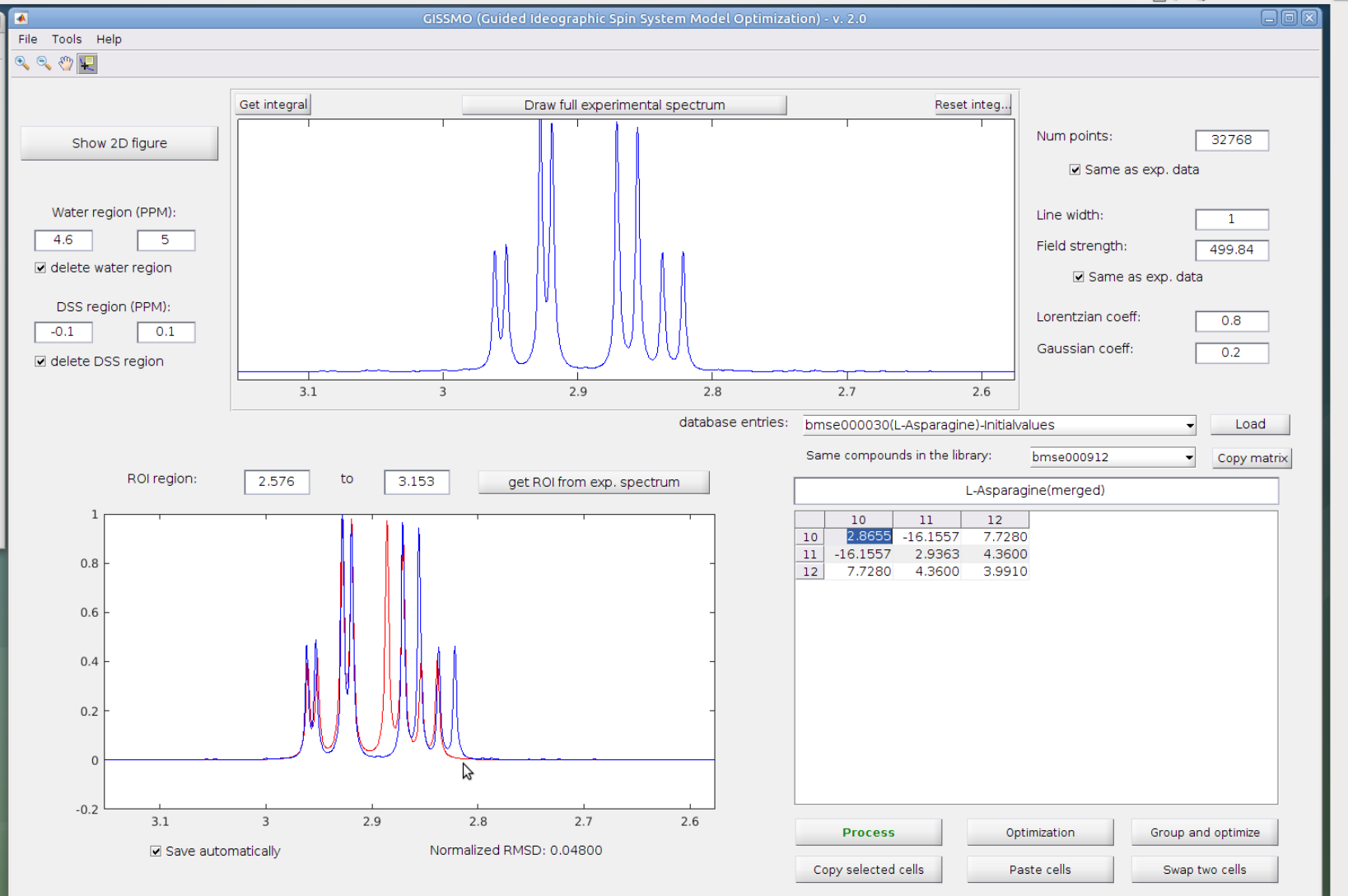
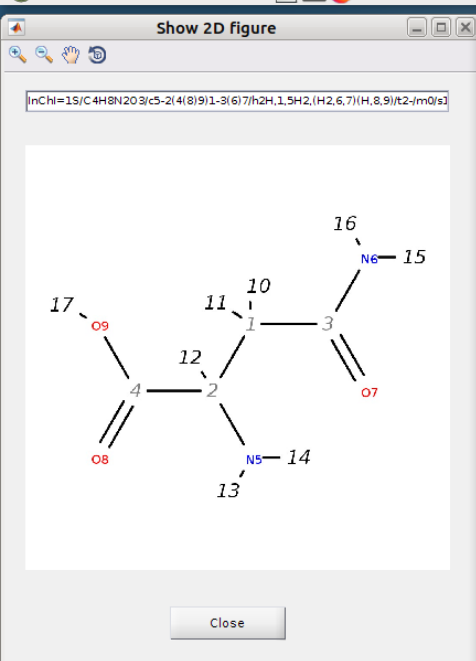


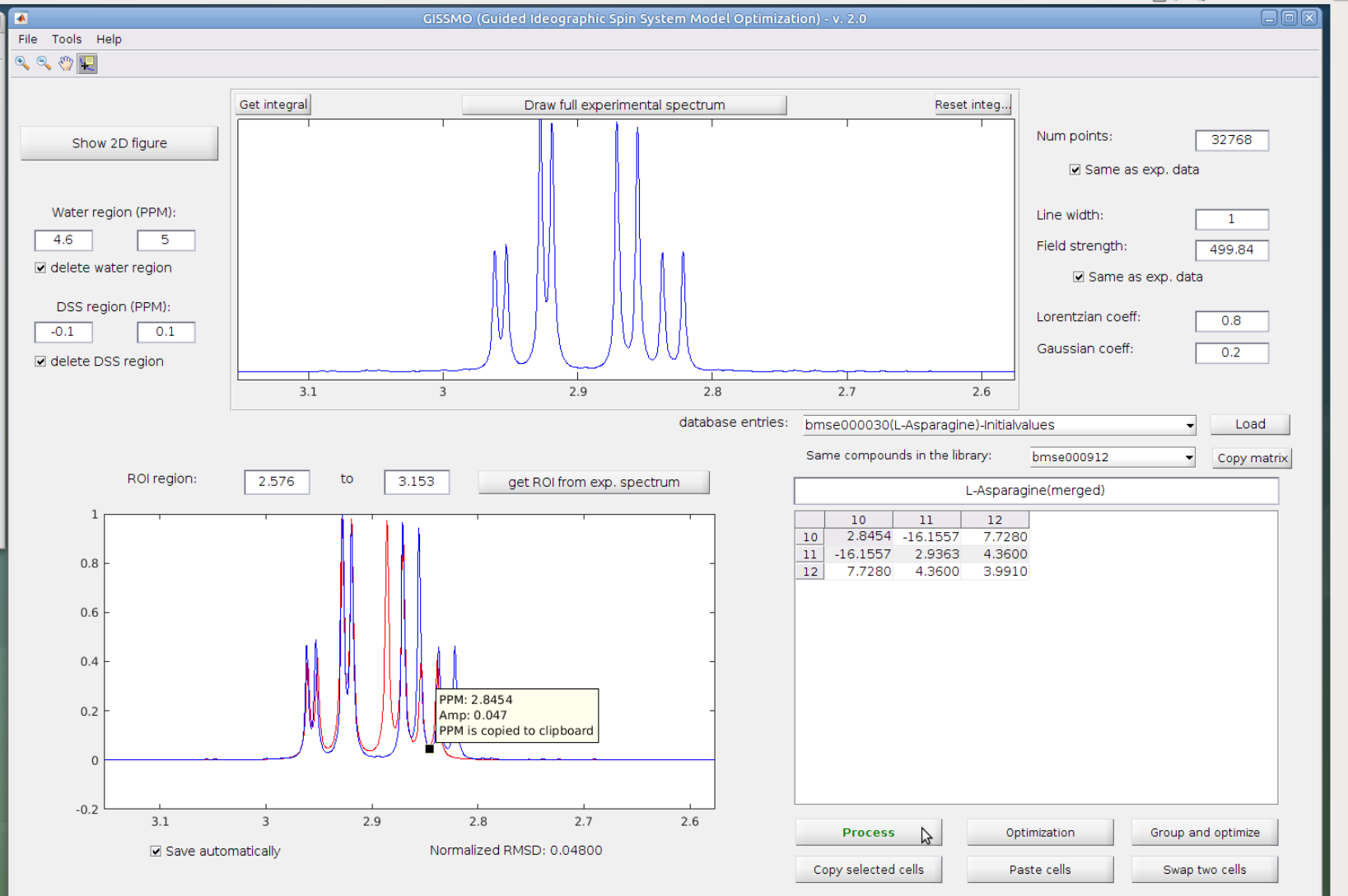
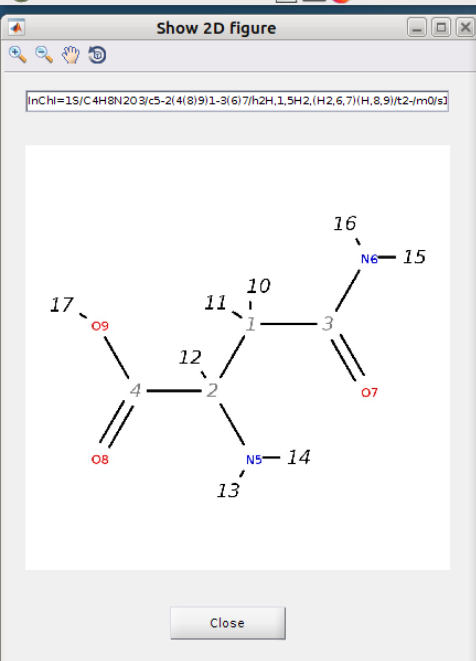


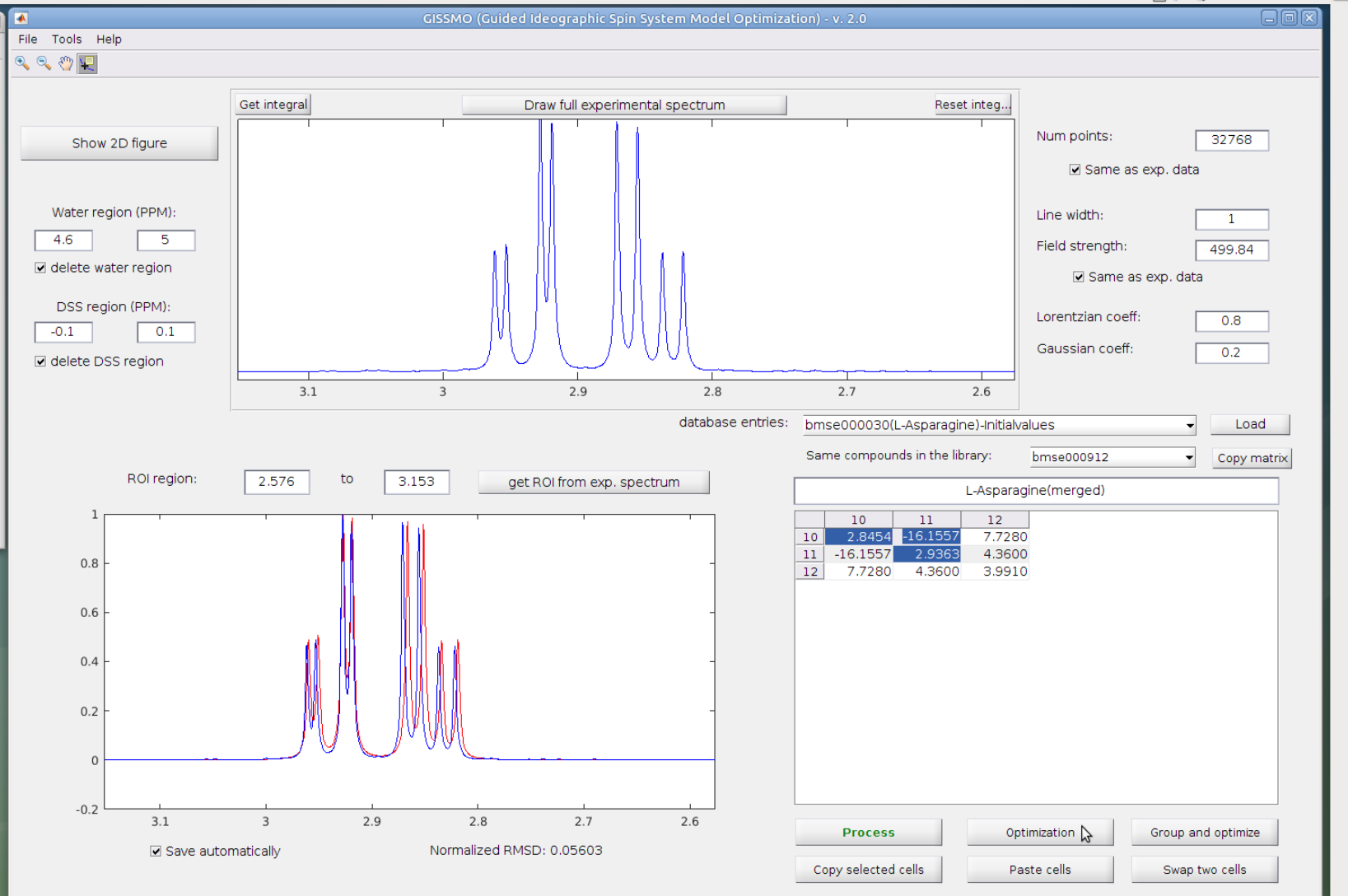
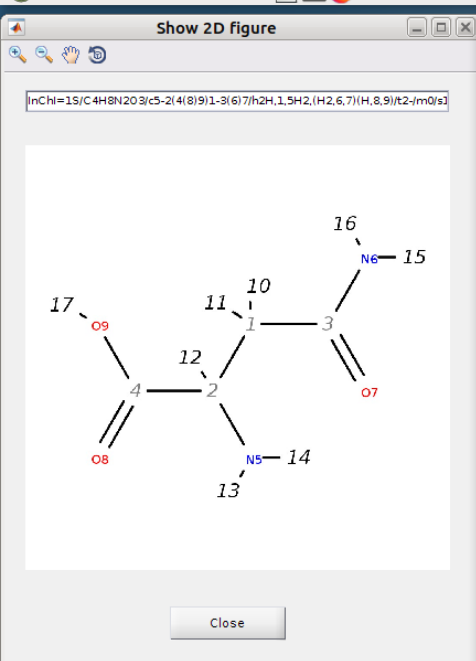


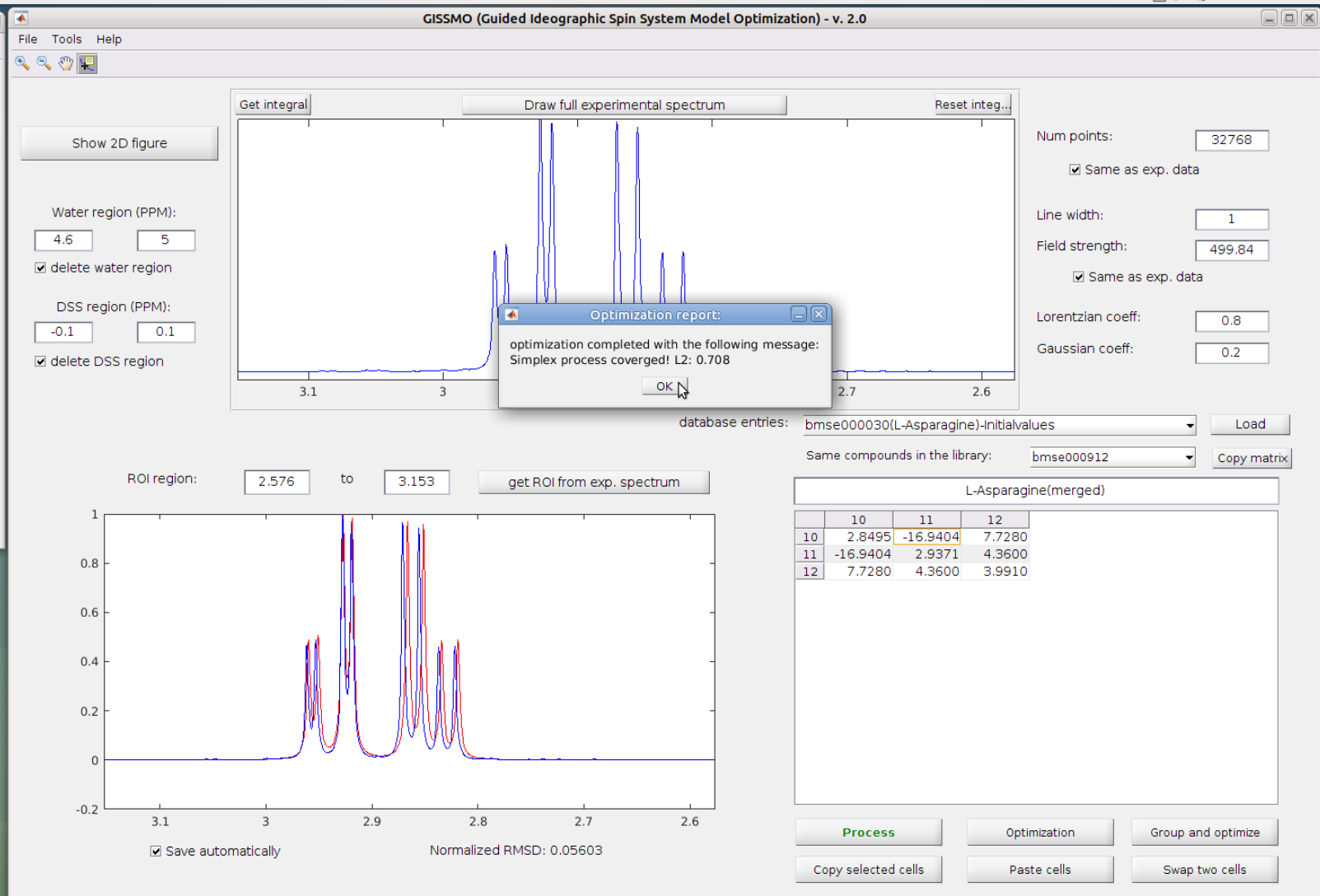
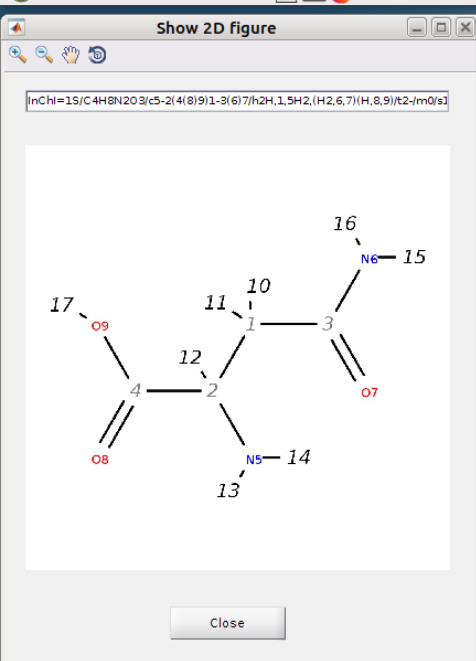


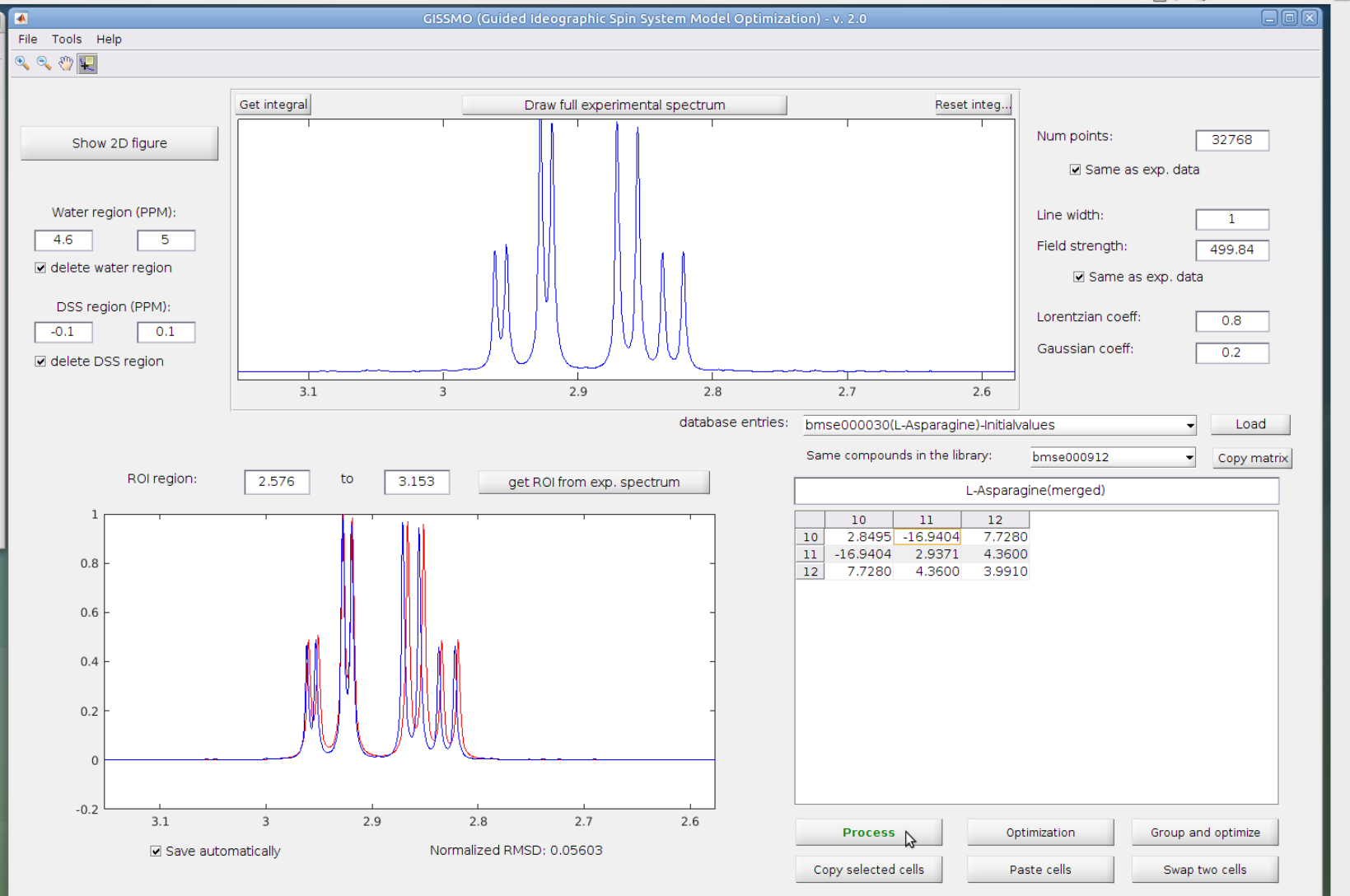
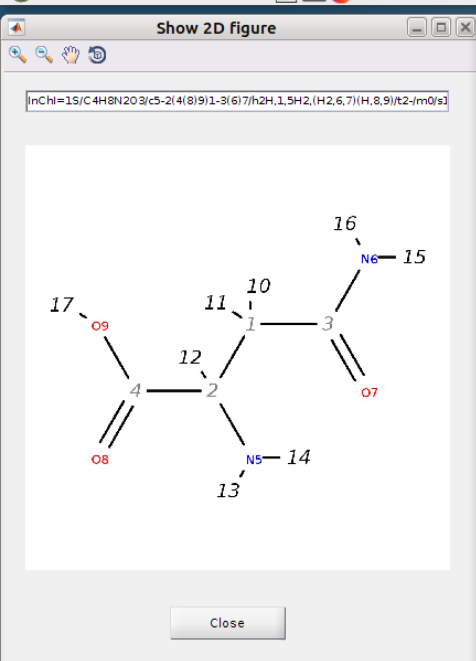


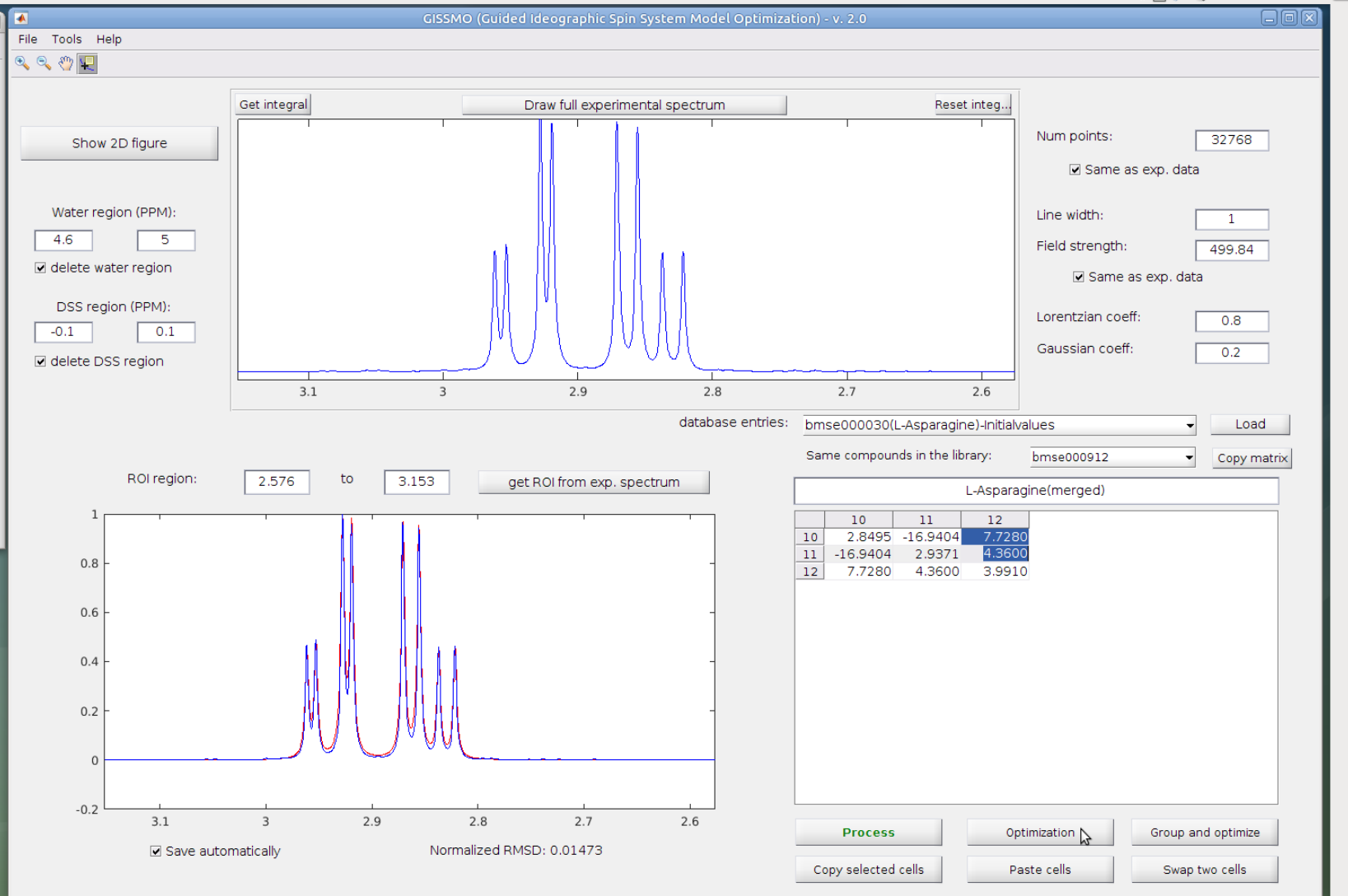
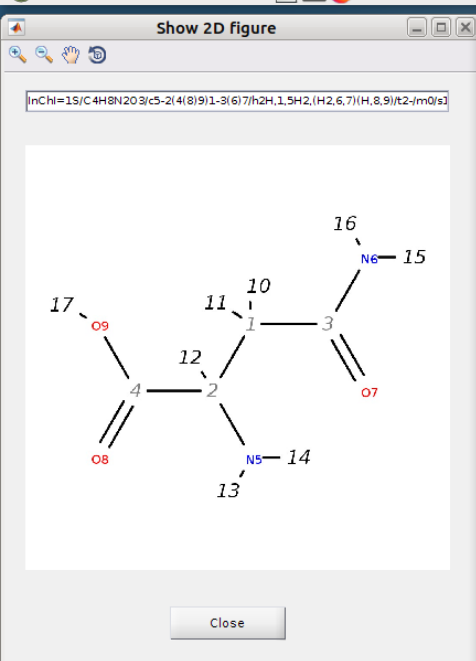


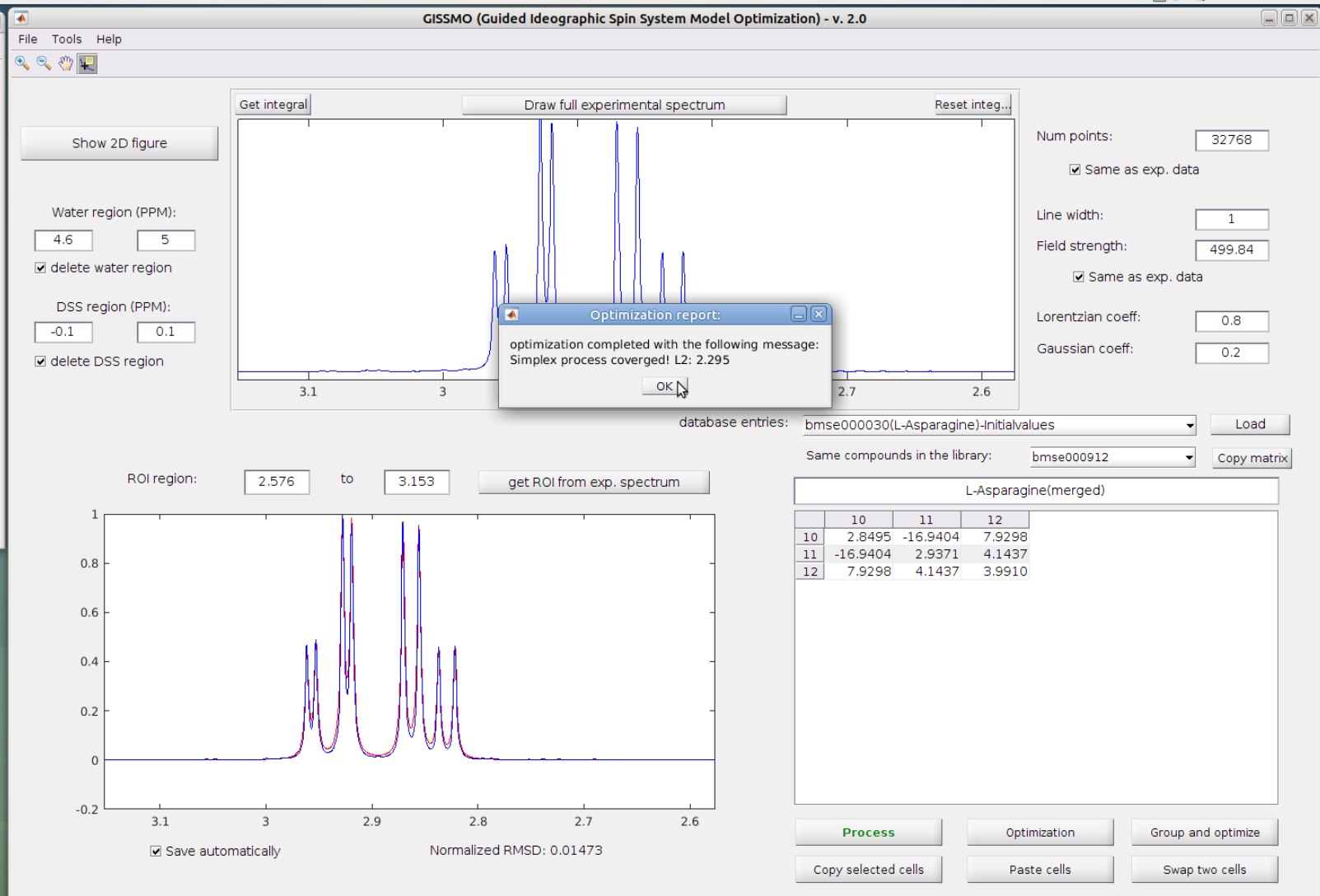
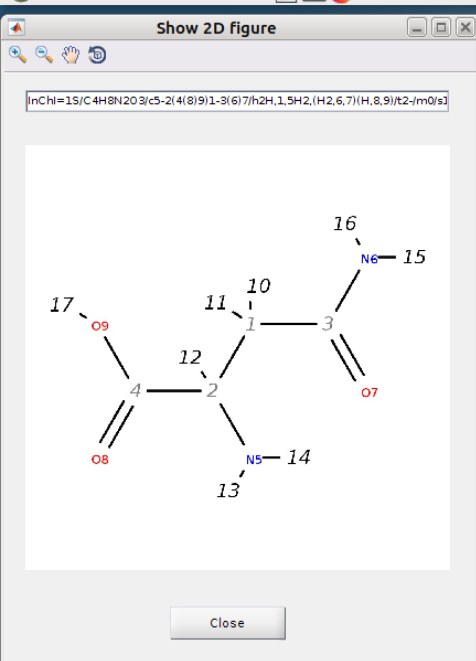


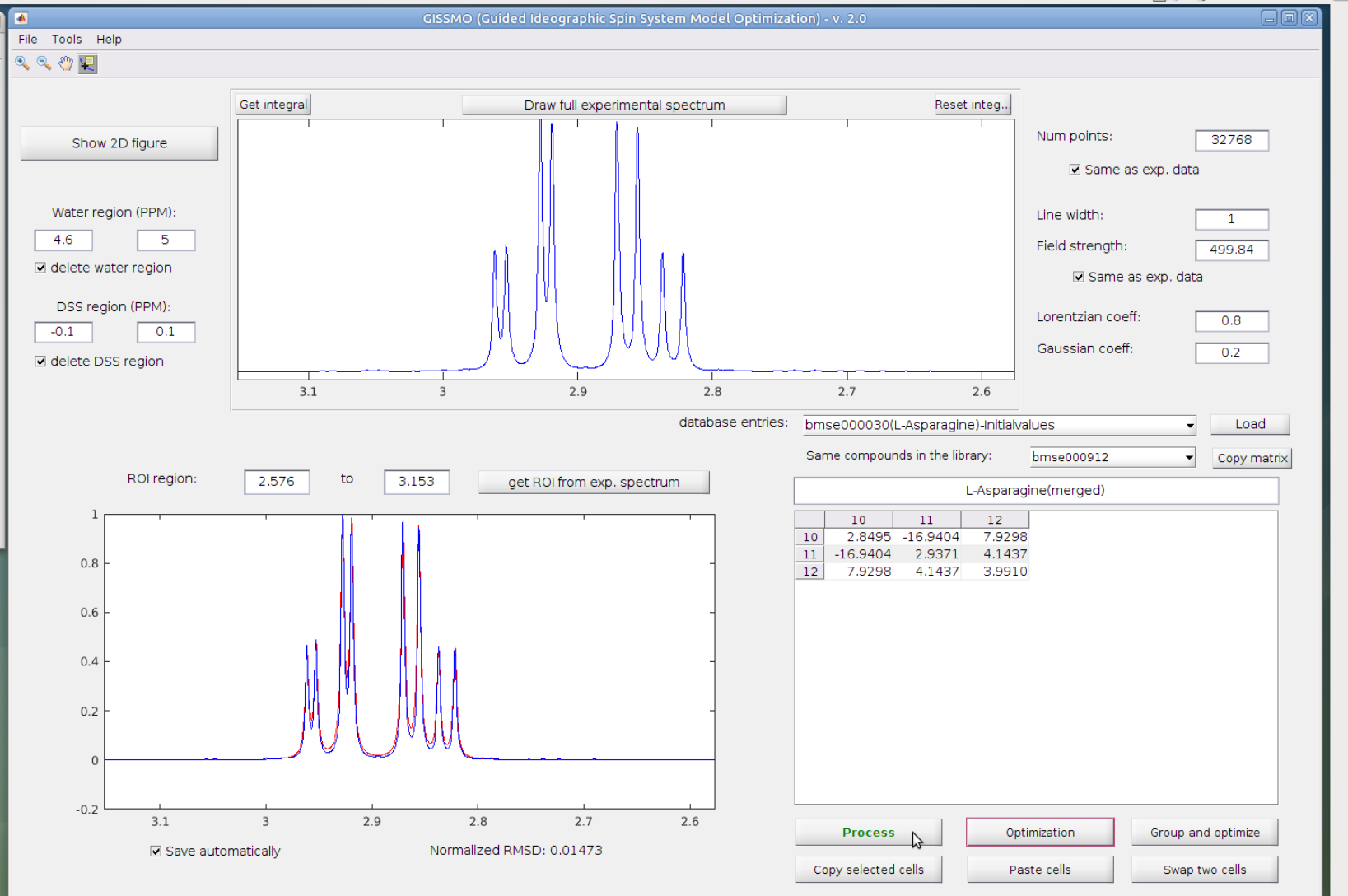
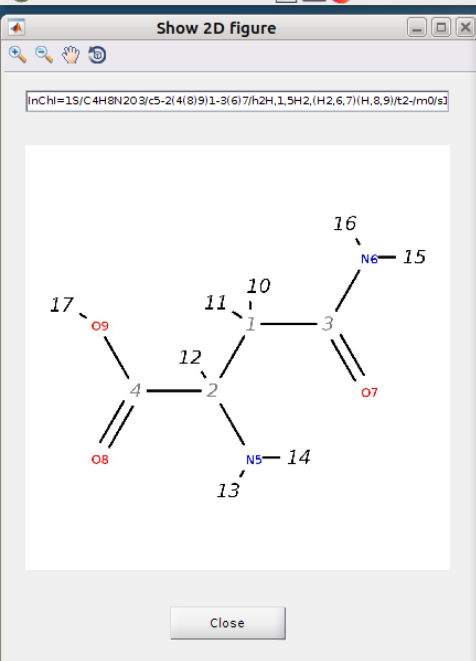


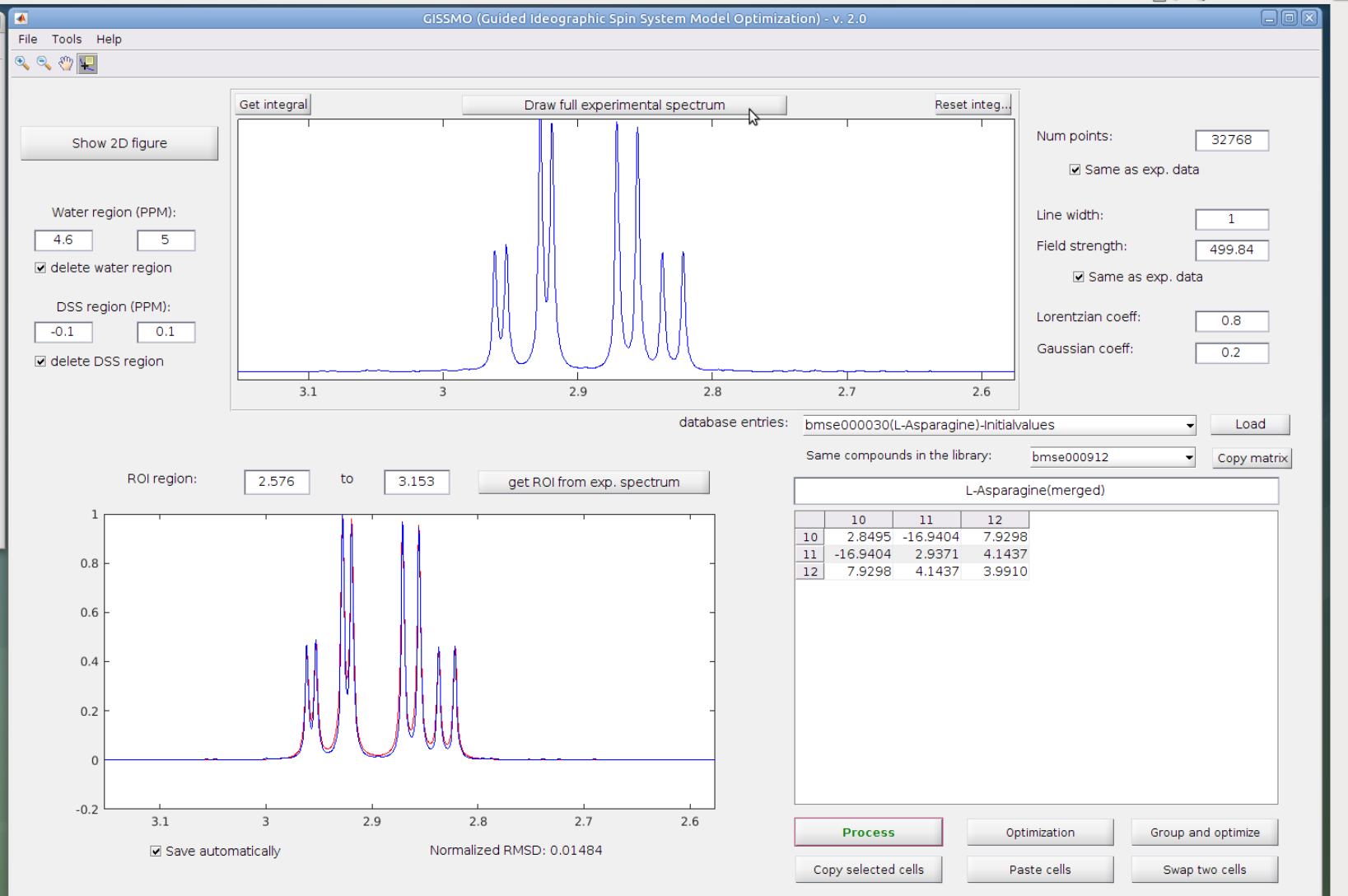
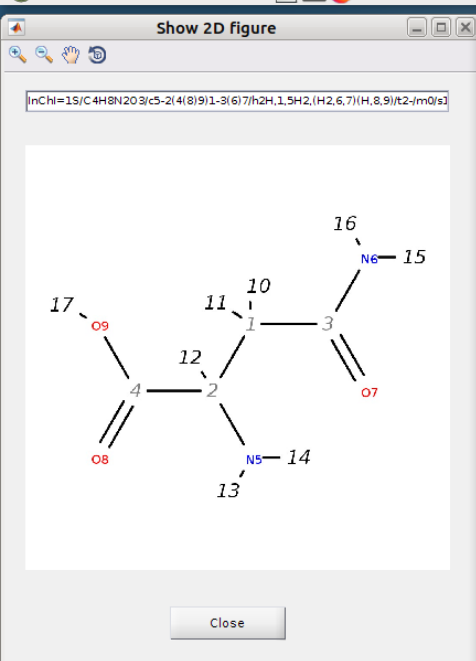


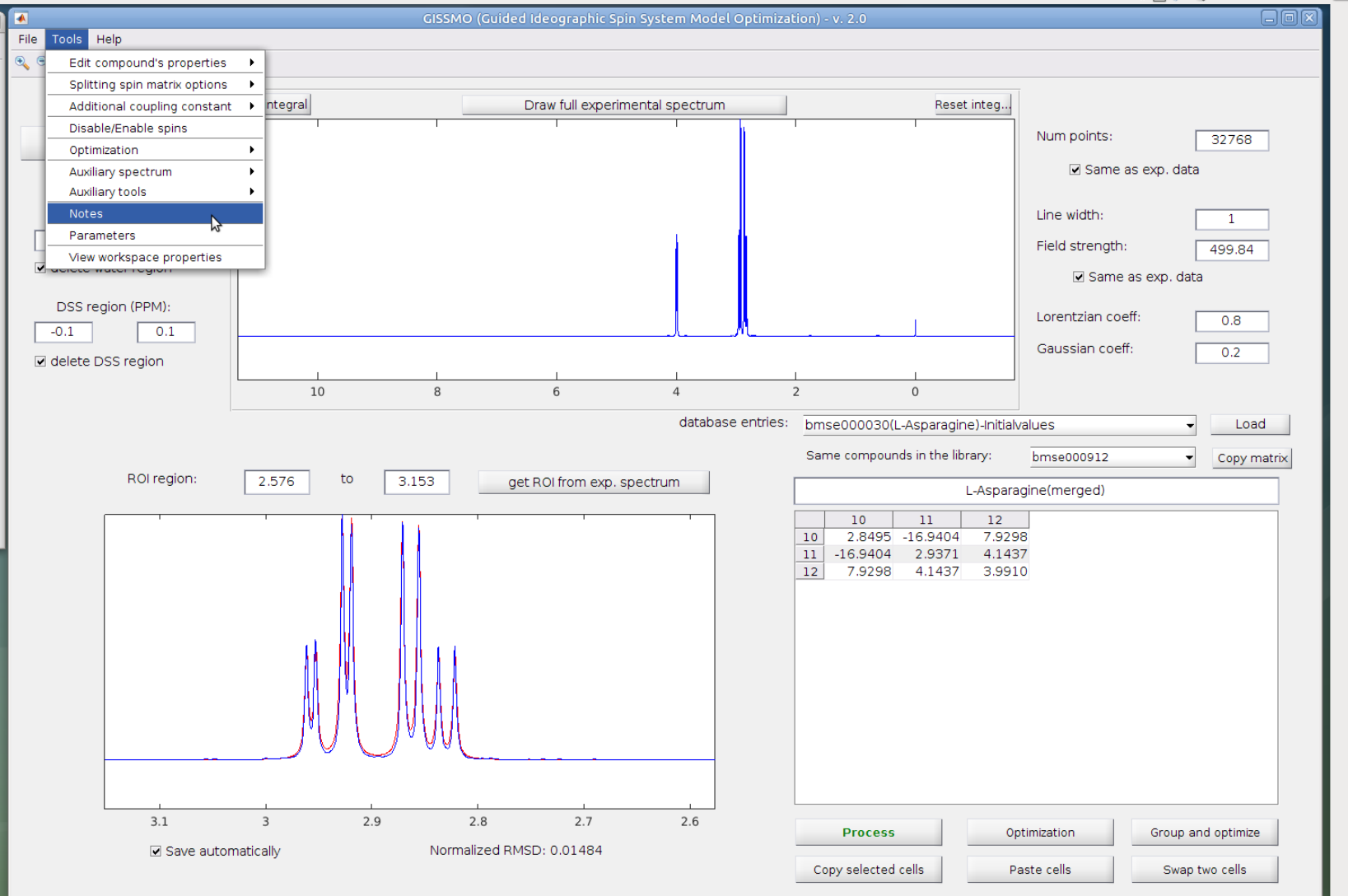
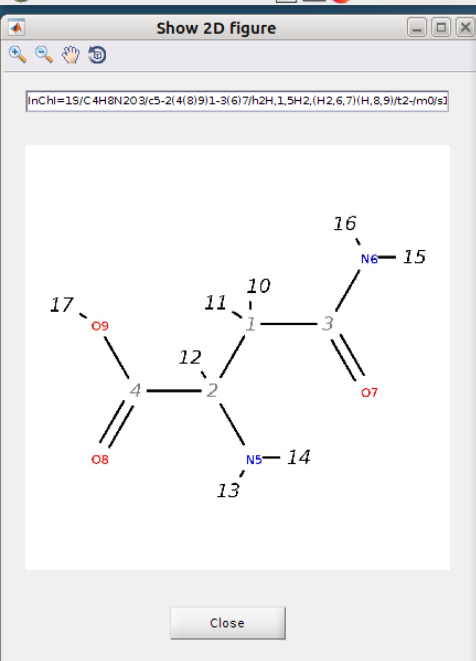


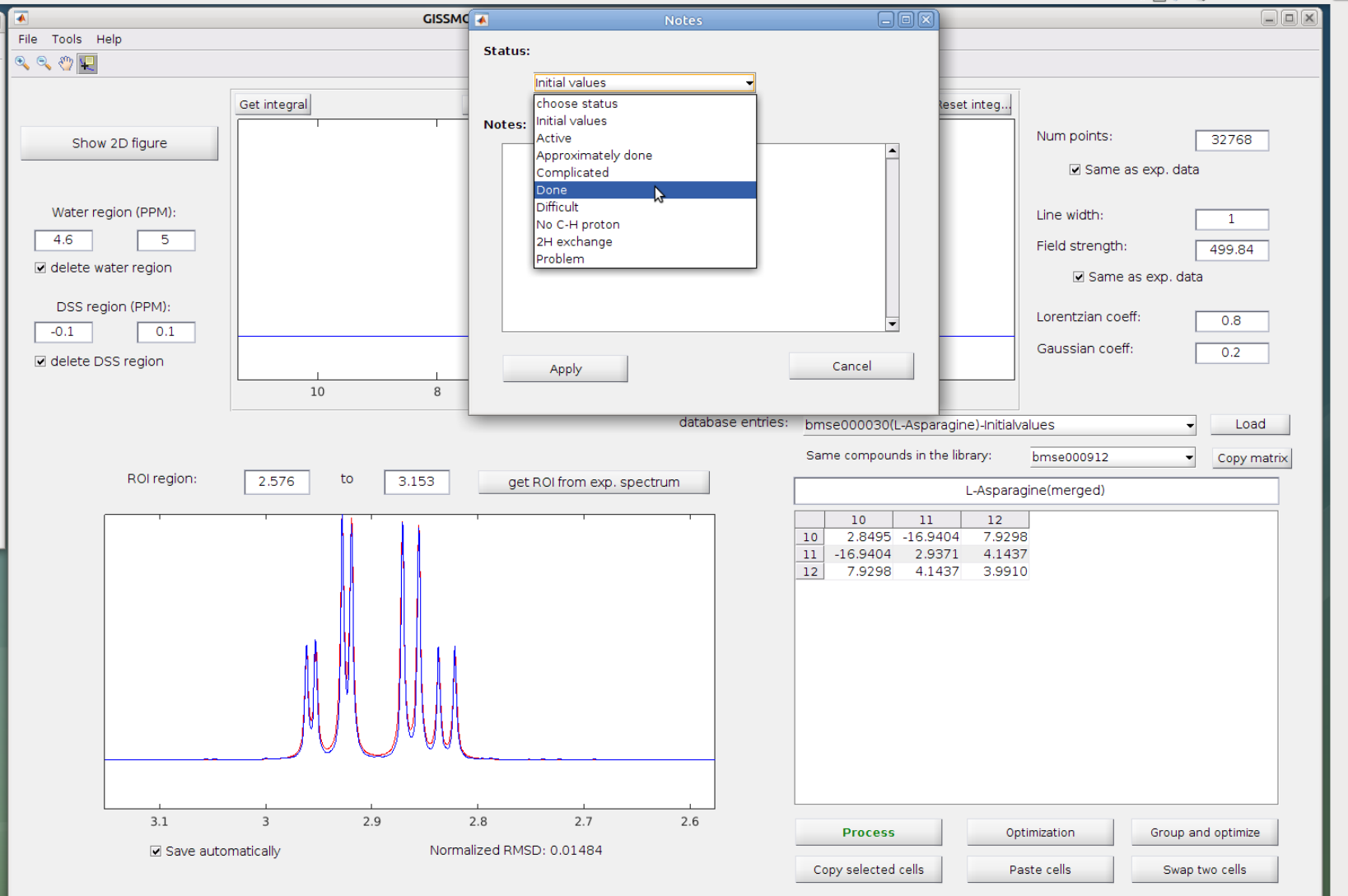
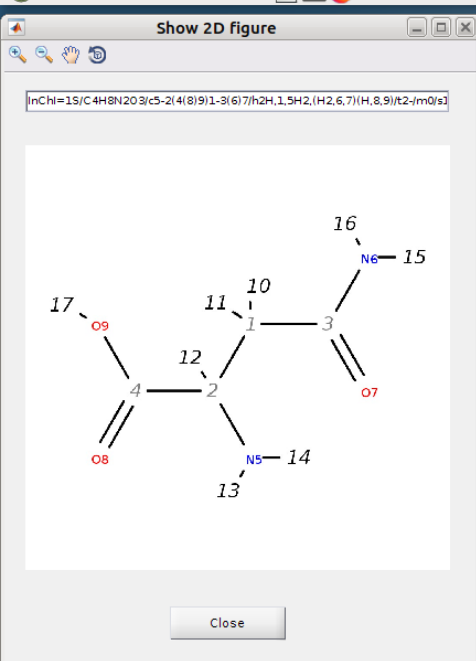


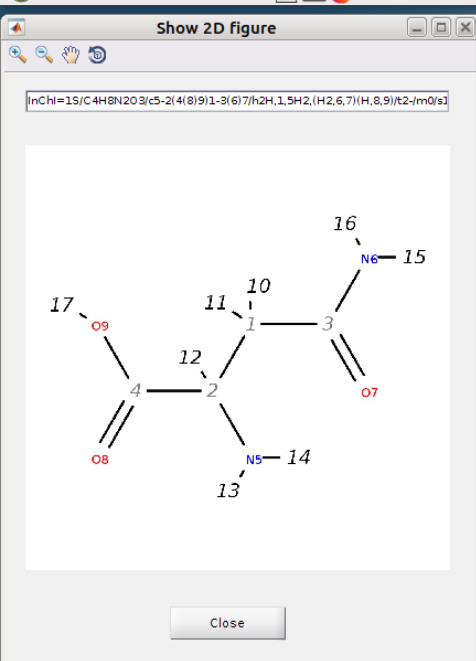












GISSMO

File Tools Help

Get integral

Show 2D figure

Water region (PPM):
4.6 5
☒ delete water region

DSS region (PPM):
-0.1 0.1
☒ delete DSS region

Notes

Status: Done

Notes:
add any notes

Apply Cancel

Num points: 32768
☒ Same as exp. data

Line width: 1

Field strength: 499.84
☒ Same as exp. data

Lorentzian coeff: 0.8

Gaussian coeff: 0.2

database entries: bmse000030(L-Asparagine)-Initialvalues Load

Same compounds in the library: bmse000912 Copy matrix

L-Asparagine(merged)

	10	11	12
10	2.8495	-16.9404	7.9298
11	-16.9404	2.9371	4.1437
12	7.9298	4.1437	3.9910

Process Optimization Group and optimize

Copy selected cells Paste cells Swap two cells

ROI region: 2.576 to 3.153 get ROI from exp. spectrum

Save automatically

