

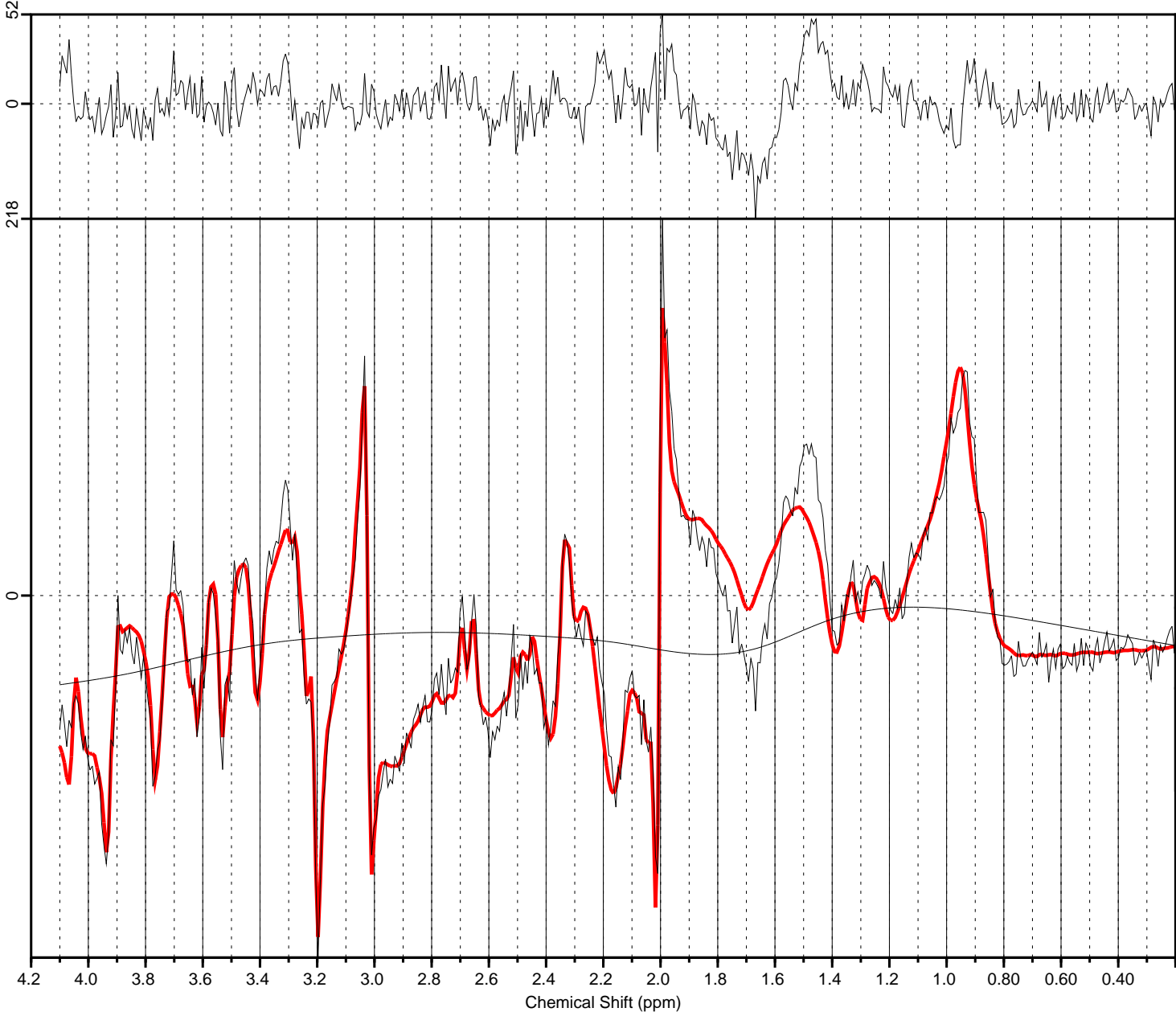
Slice_N1@18_19 28-Feb-2025 11:38:35

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Ref.: Magn. Reson. Med. 30:672-679 (1993).

Fri Feb 28 11:38:38 2025



Conc.	%SD	/Cr+PCr	Metabolite
3.25E-04	4%	3.2E-04	Mac
3.464	26%	3.363	Cr
4.776	18%	4.637	PCr
7.956	11%	7.724	Ins
10.407	7%	10.103	NAA
6.859	12%	6.659	Tau
0.867	96%	0.841	PCho
0.874	90%	0.849	GPC
10.808	10%	10.493	Glu
3.432	23%	3.332	Gln
0.000	999%	0.000	Ala
1.726	67%	1.676	Asc
1.907	46%	1.852	Asp
0.540	88%	0.525	GABA
0.404	149%	0.392	Glc
1.357	32%	1.317	GSH
1.846	36%	1.792	Lac
0.988	55%	0.959	NAAG
1.023	106%	0.994	PE
1.741	16%	1.690	GPC+PCho
11.395	7%	11.063	NAA+NAAG
14.241	9%	13.825	Glu+Gln
8.240	9%	8.000	Cr+PCr
11.395	7%	11.063	NAA+NAAG
8.240	9%	8.000	Cr+PCr
14.241	9%	13.825	Glu+Gln

DIAGNOSTICS	
1 info	STARTV 20
Doing Water-Scaling	

MISCELLANEOUS OUTPUT	
FWHM = 0.016 ppm	S/N = 5
Data shift = 0.024 ppm	
Ph: -88 deg	0.0 deg/ppm

INPUT CHANGES	
hwdwat= 0.5	
wconst= 44444.	
ppmst= 4.1	
ppmend= 0.2	
nunfil= 768	
nomit= 15	
conrel=8	
namrel='Cr+PCr'	
neach= 999	
hzpppm= 400.216	

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DIAGNOSTICS				
1	info	STARTV	20	
Doing Water-Scaling				
MISCELLANEOUS OUTPUT				
FWHM = 0.016 ppm S/N = 5				
Data shift = 0.024 ppm				
Ph: -88 deg 0.0 deg/ppm				
INPUT CHANGES				
hwdwat= 0.5				
wconc= 44444.				
ppmst= 4.1				
ppmend= 0.2				
nunfil= 768				
nomit= 15				
conrel=8				
namrel='Cr+PCr'				
				neach= 999
				hzpppm= 400.216
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				DMRSI\Slice_N1\Data\Slice_N1@18_19.RAW'
				filps= 'Z:\Brayan\Data Processing\28022025_2DFID
				MRSI\Slice_N1\Data\Slice_N1@18_19.ps'
				filh2o= 'Z:\Brayan\Data Processing\28022025_2DFI
				DMRSI\Slice_N1\Data\Slice_N1@18_19w.RAW'
				filbas= 'Z:\Brayan\Basis_Sets\TE=1300microsec\9.
				4T_29102024\9_4T_SIM_MRSI_Brayan_TE=1300micros
				ec_9_4T_TRUEBasis29102024.BASIS'
				filcoo= 'Z:\Brayan\Data Processing\28022025_2DFI
				DMRSI\Slice_N1\Data\Slice_N1@18_19.coord'
				filtab= 'Z:\Brayan\Data Processing\28022025_2DFI
				DMRSI\Slice_N1\Data\tables\Slice_N1@18_19.tabl
				e'
				ltable= 7
				lcoord=9
				nlhmet = 3
				wsppm = 0.0
				wsmet = 'Cr'
				dorefs = F
				dows= T
				dgppmx= 5
				dgppmn= -5
				sddegp= 0
				degppm= 0
				dkntmn= 0.25
				deltat= 2e-04
				chomit= '-CrCH2' 'Gua' 'Ser' 'Lip13a' 'Lip13b' 'Lip09' 'MM09' 'Lip20' 'MM20' 'MM12' 'MM14' 'MM17' 'Ace' 'Cit' 'bHB'
				chcomb= 'GPC+PCho' 'NAA+NAAG' 'Glu+Gln' 'Cr+PCr'
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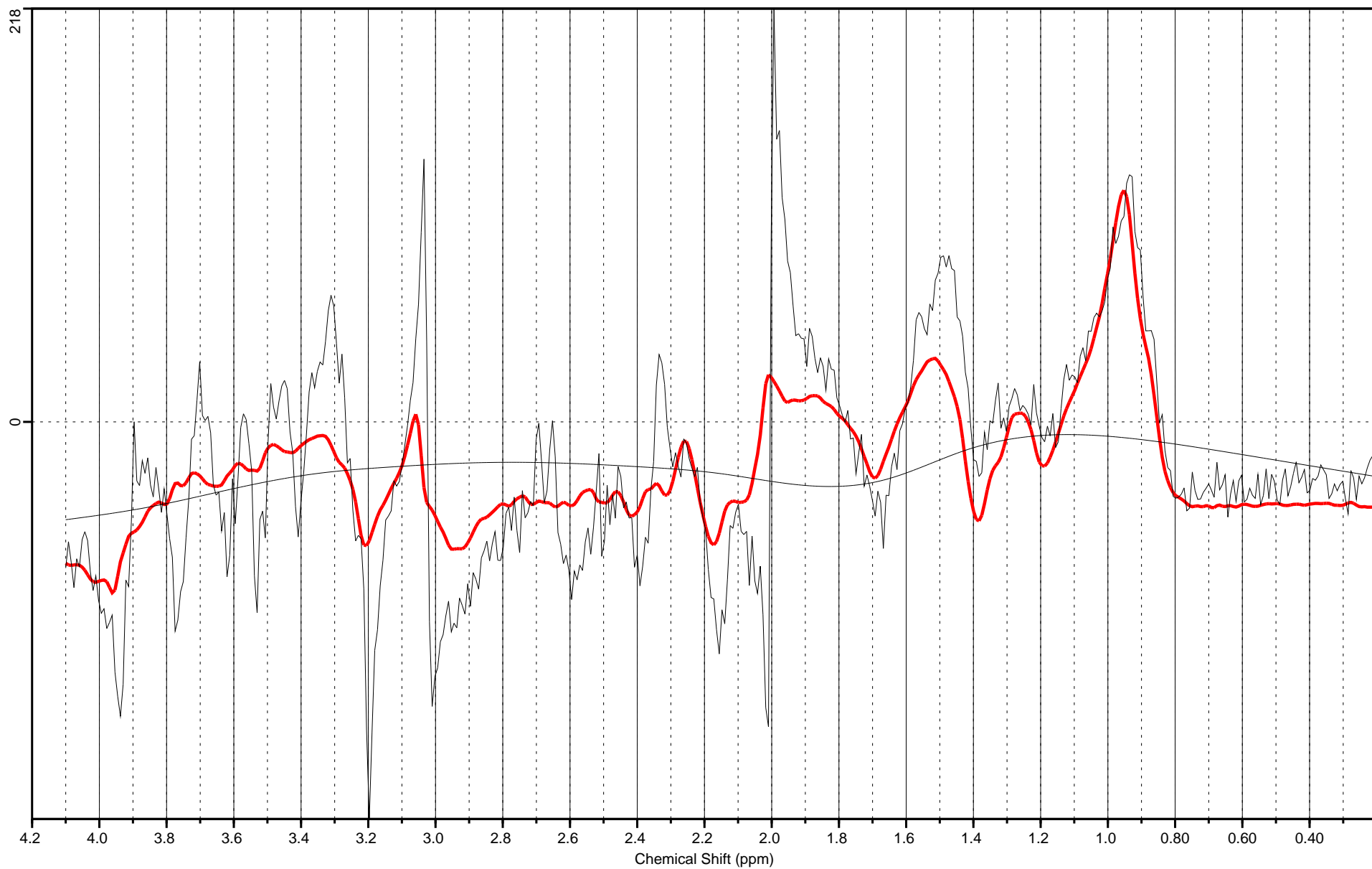
Mac Conc. = 3.25E-04

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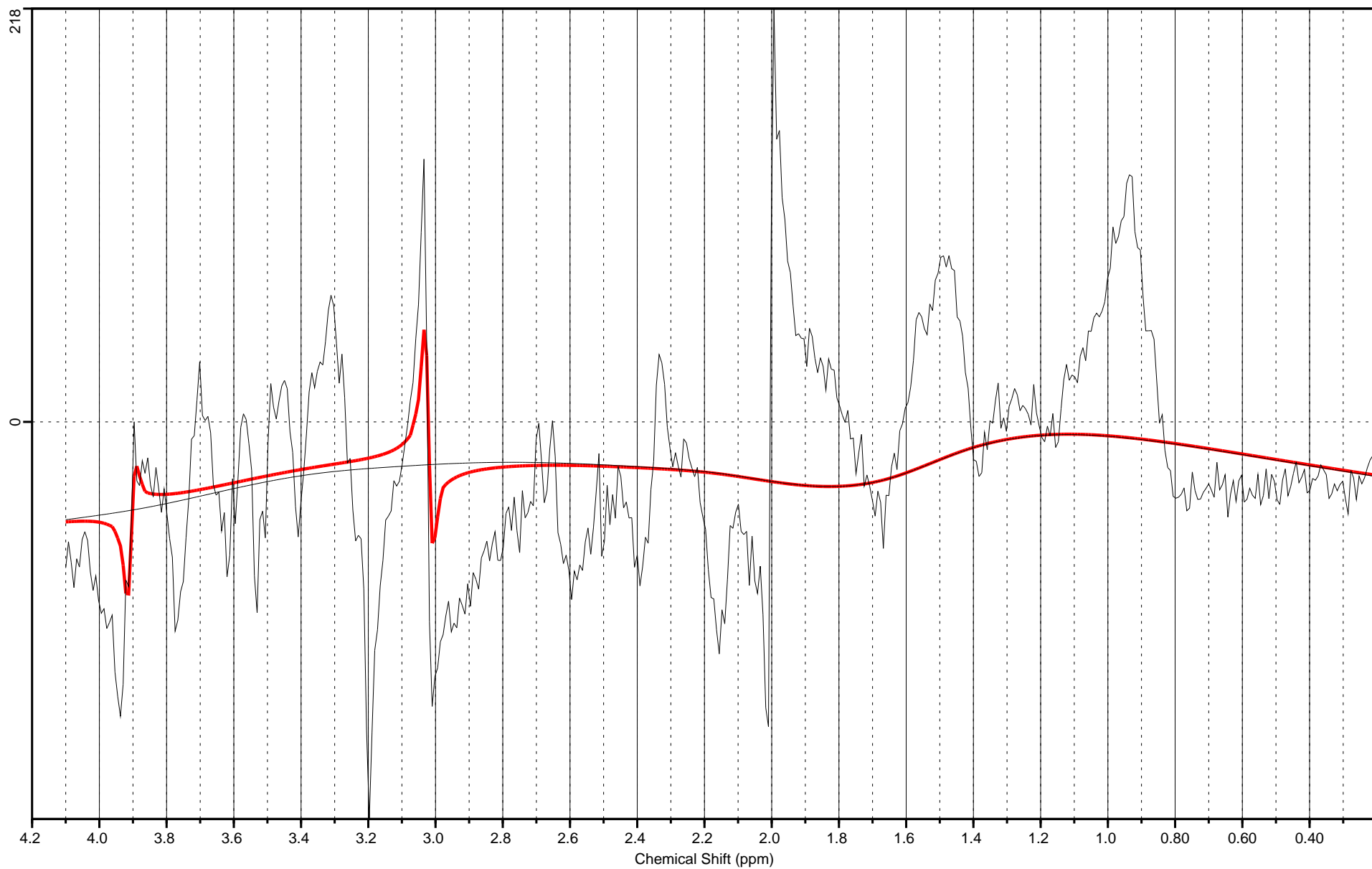
Cr Conc. = 3.46E+00

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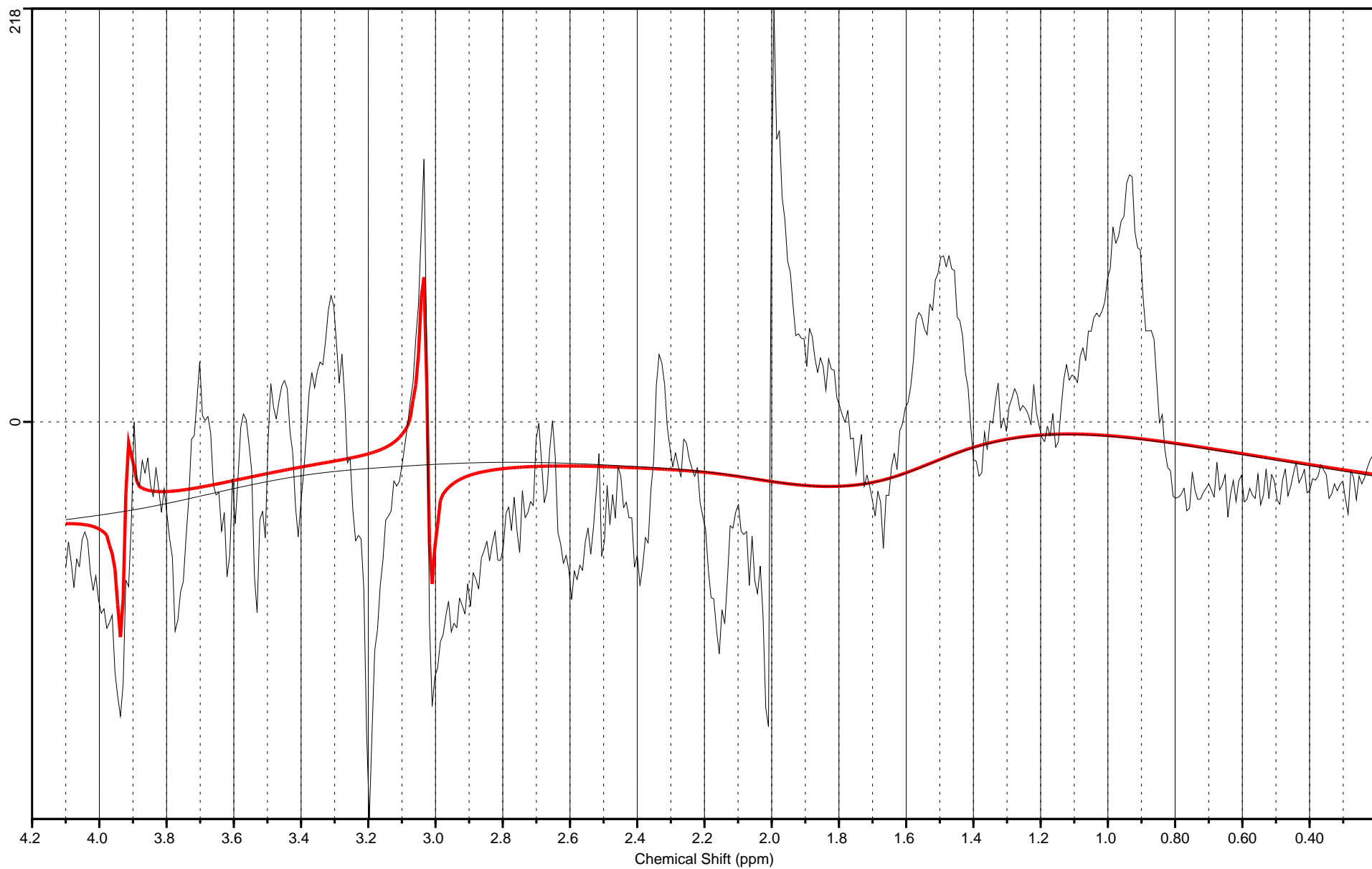
PCr Conc. = 4.78E+00

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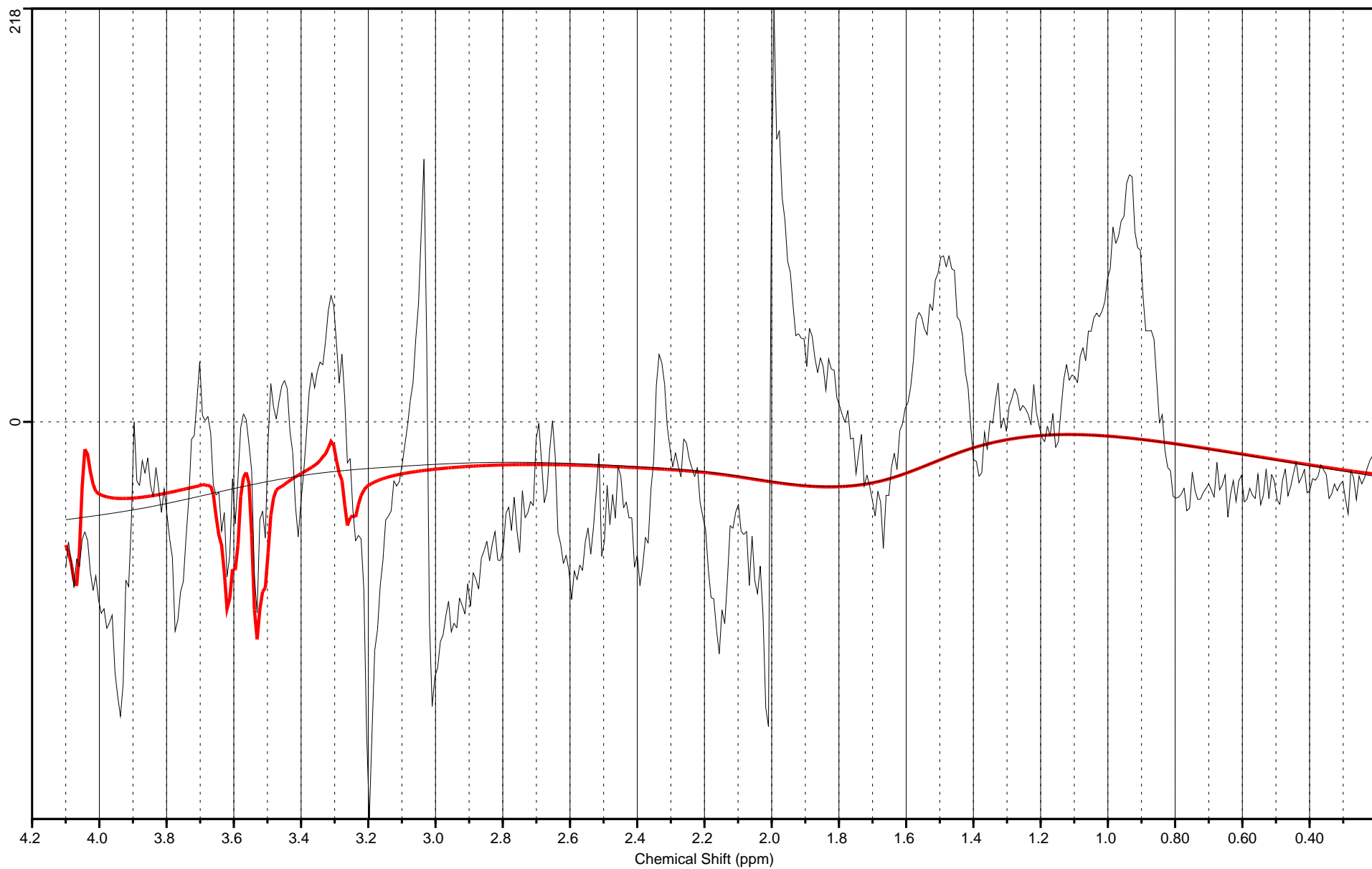
Ins Conc. = 7.96E+00

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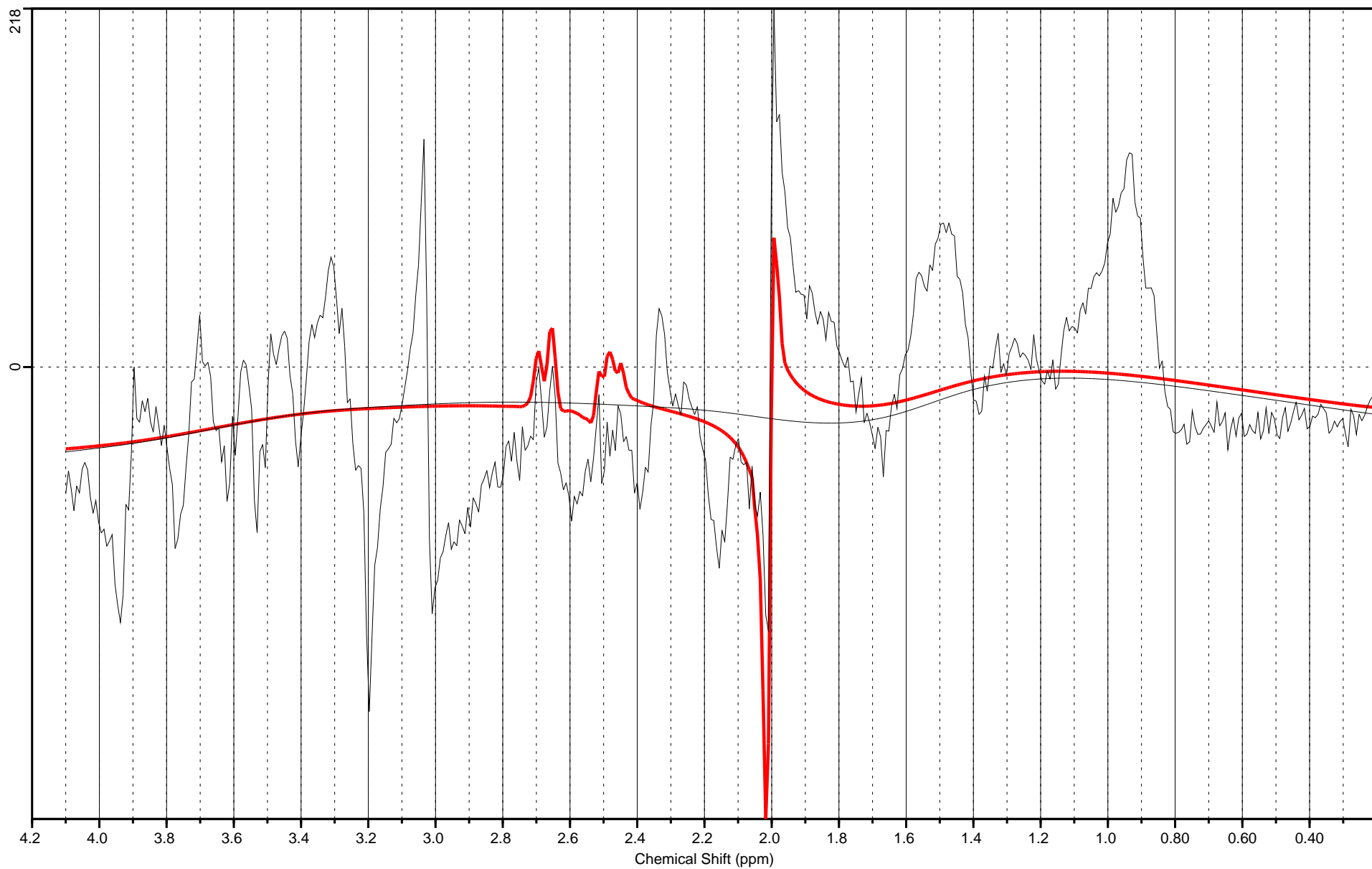
NAA Conc. = 1.04E+01

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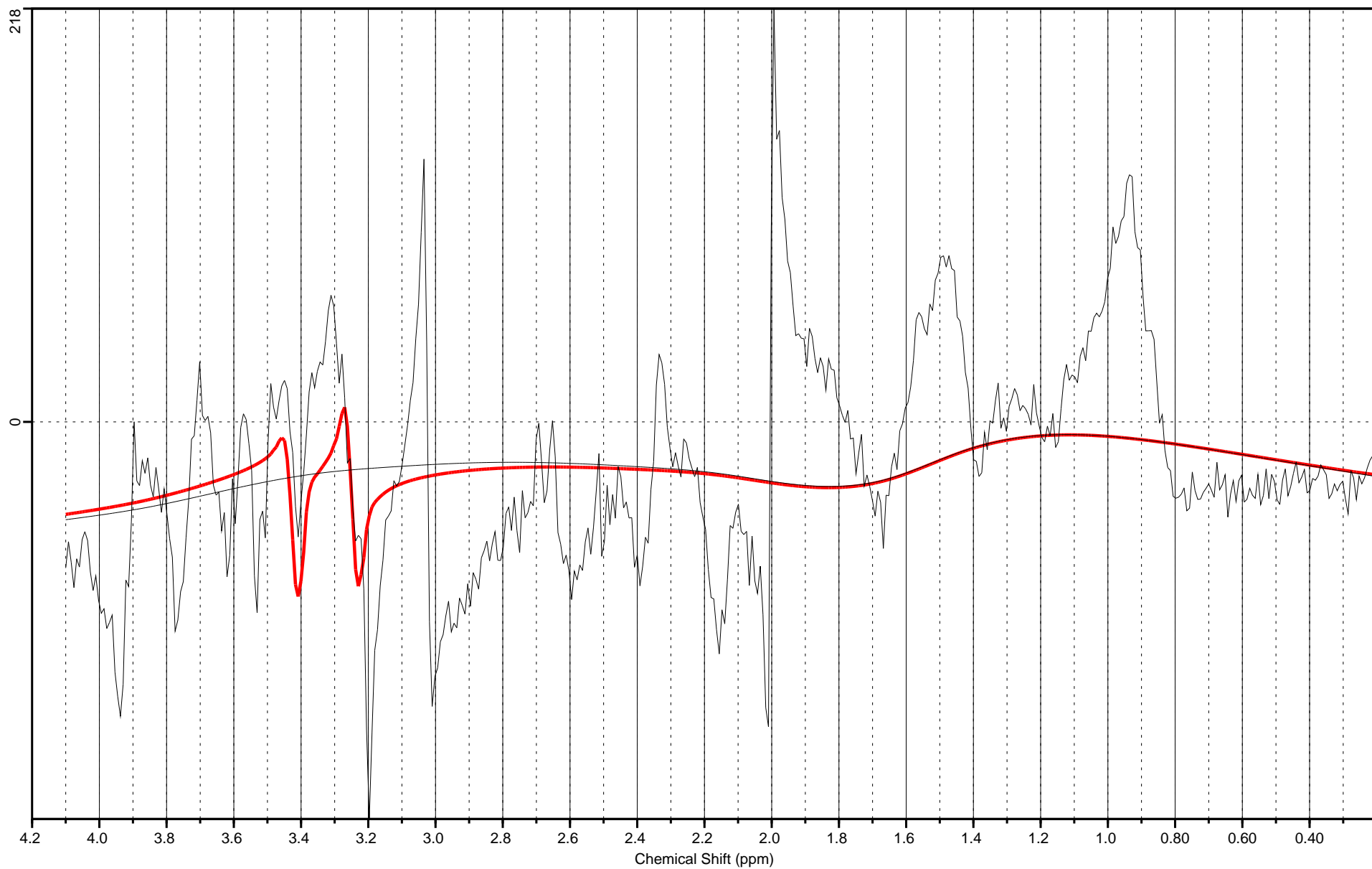
Tau Conc. = 6.86E+00

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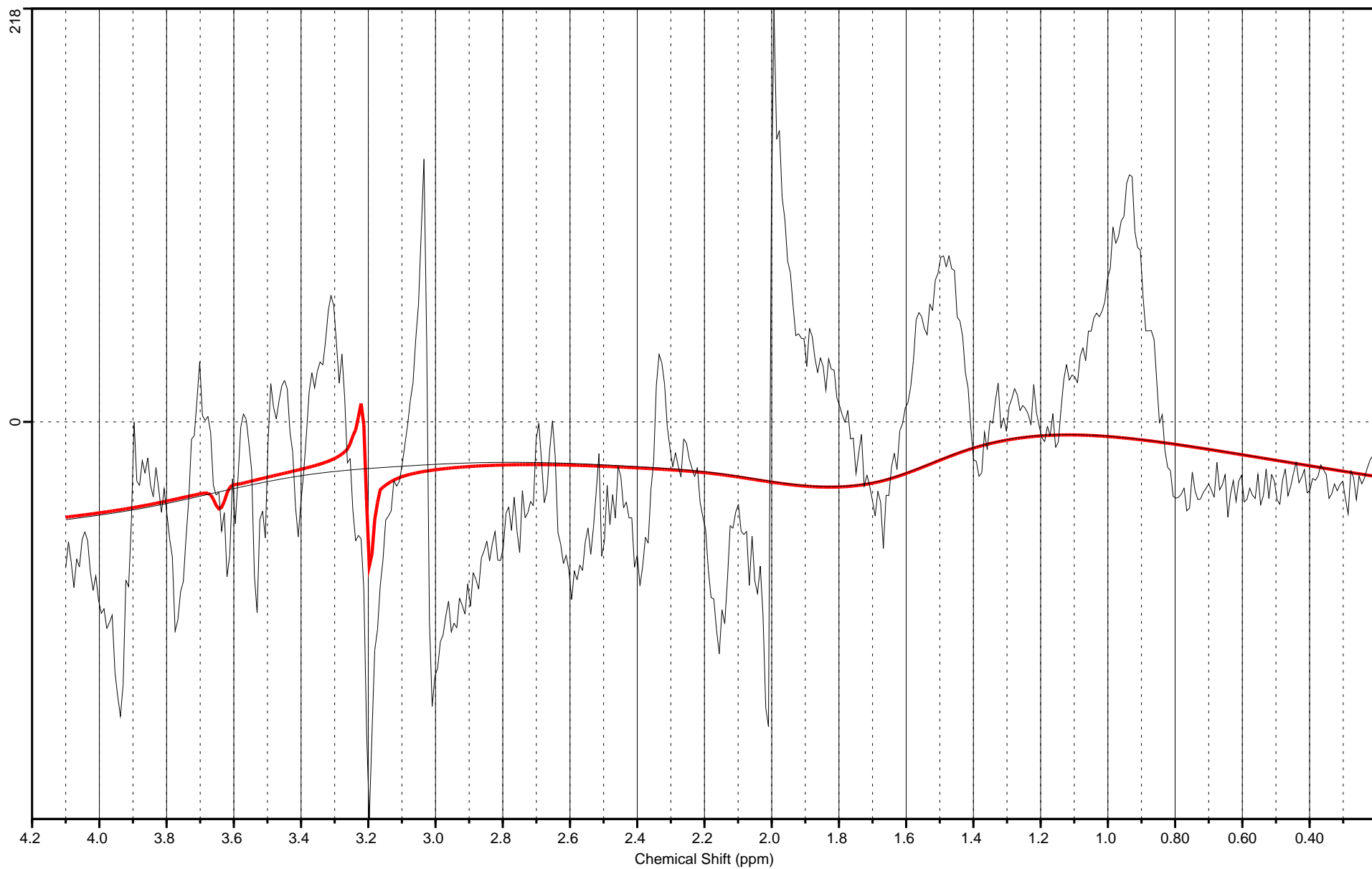
PCho Conc. = 8.67E-01

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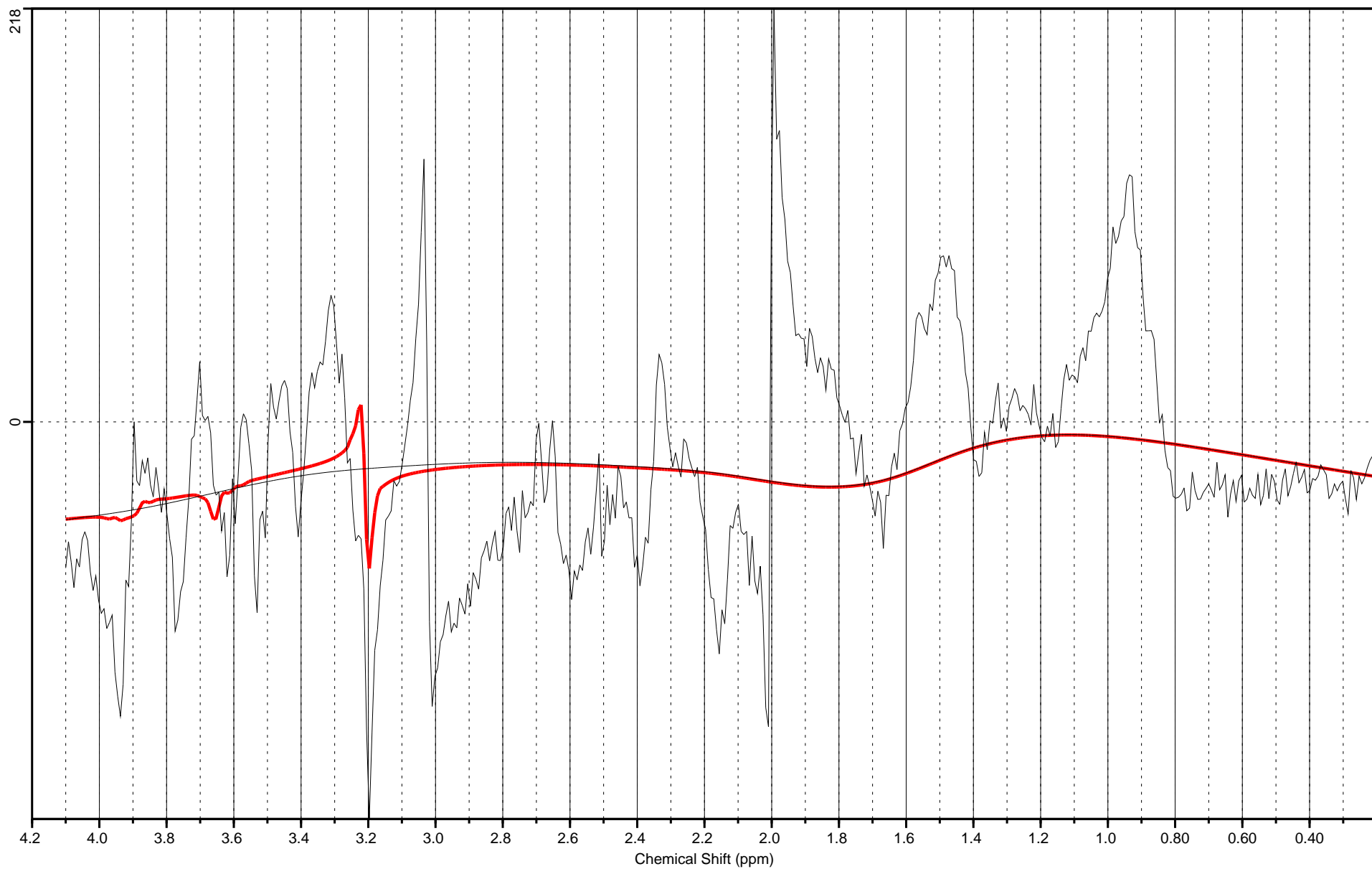
GPC Conc. = 8.74E-01

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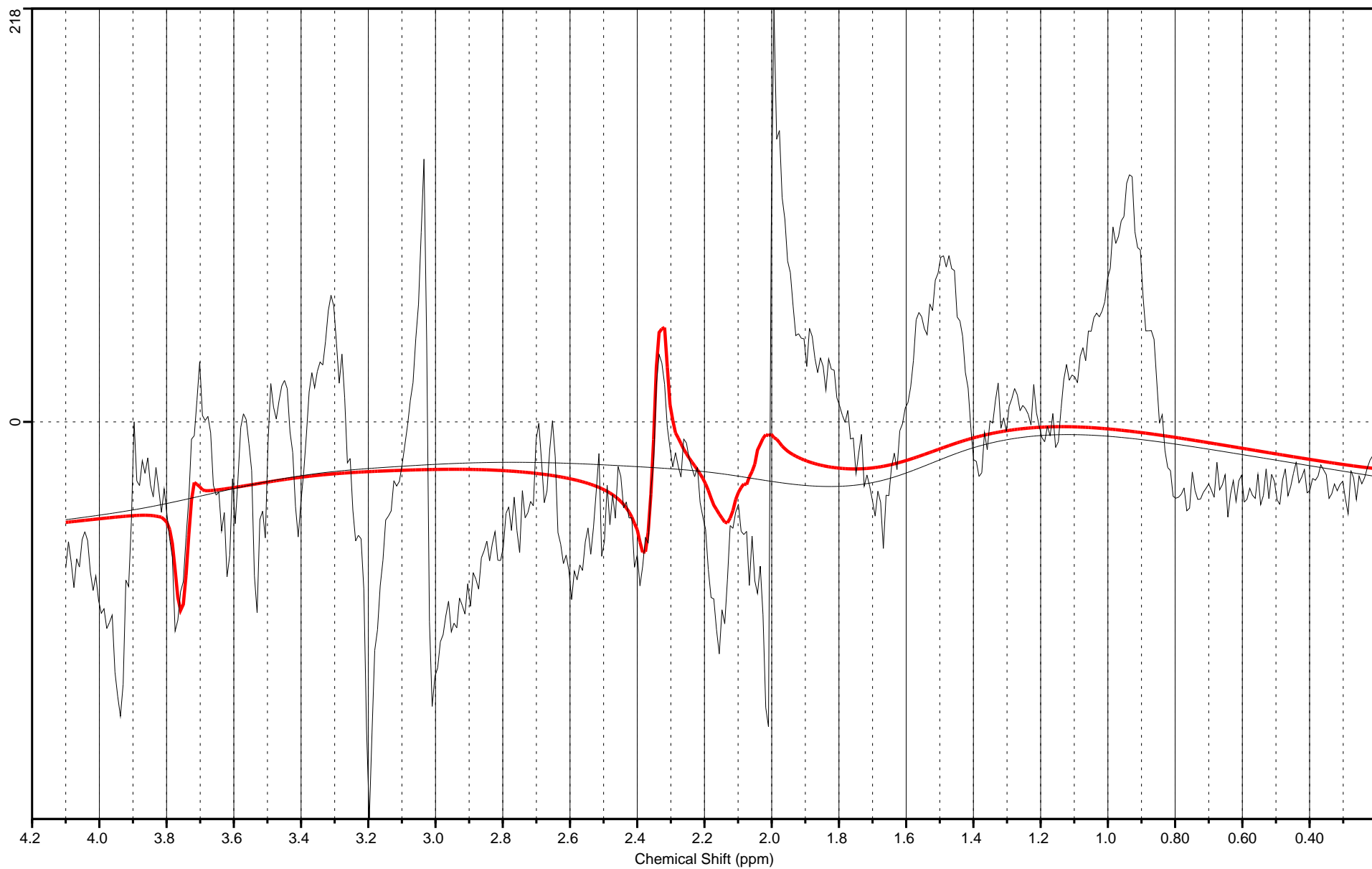
Glu Conc. = 1.08E+01

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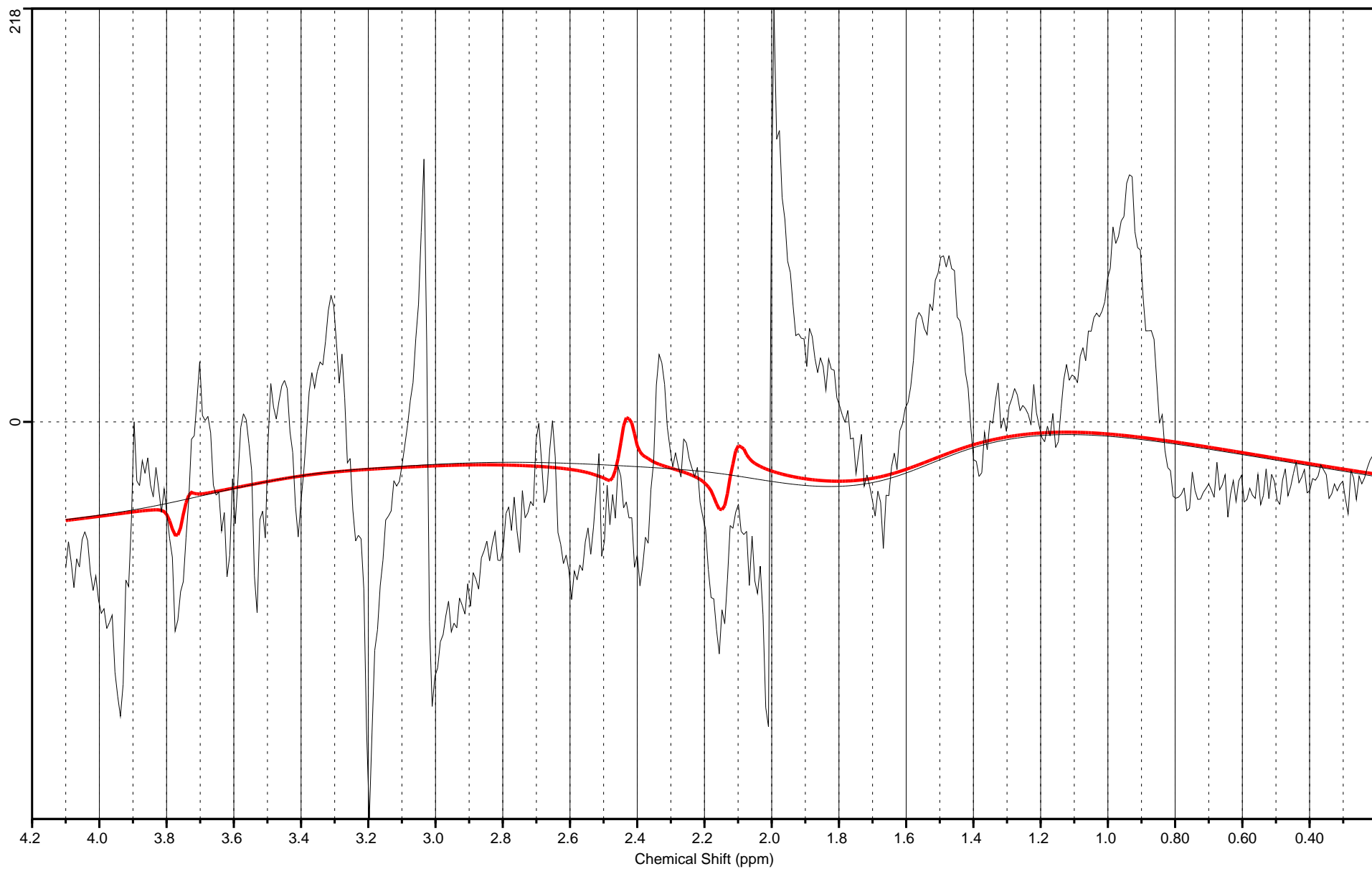
Gln Conc. = 3.43E+00

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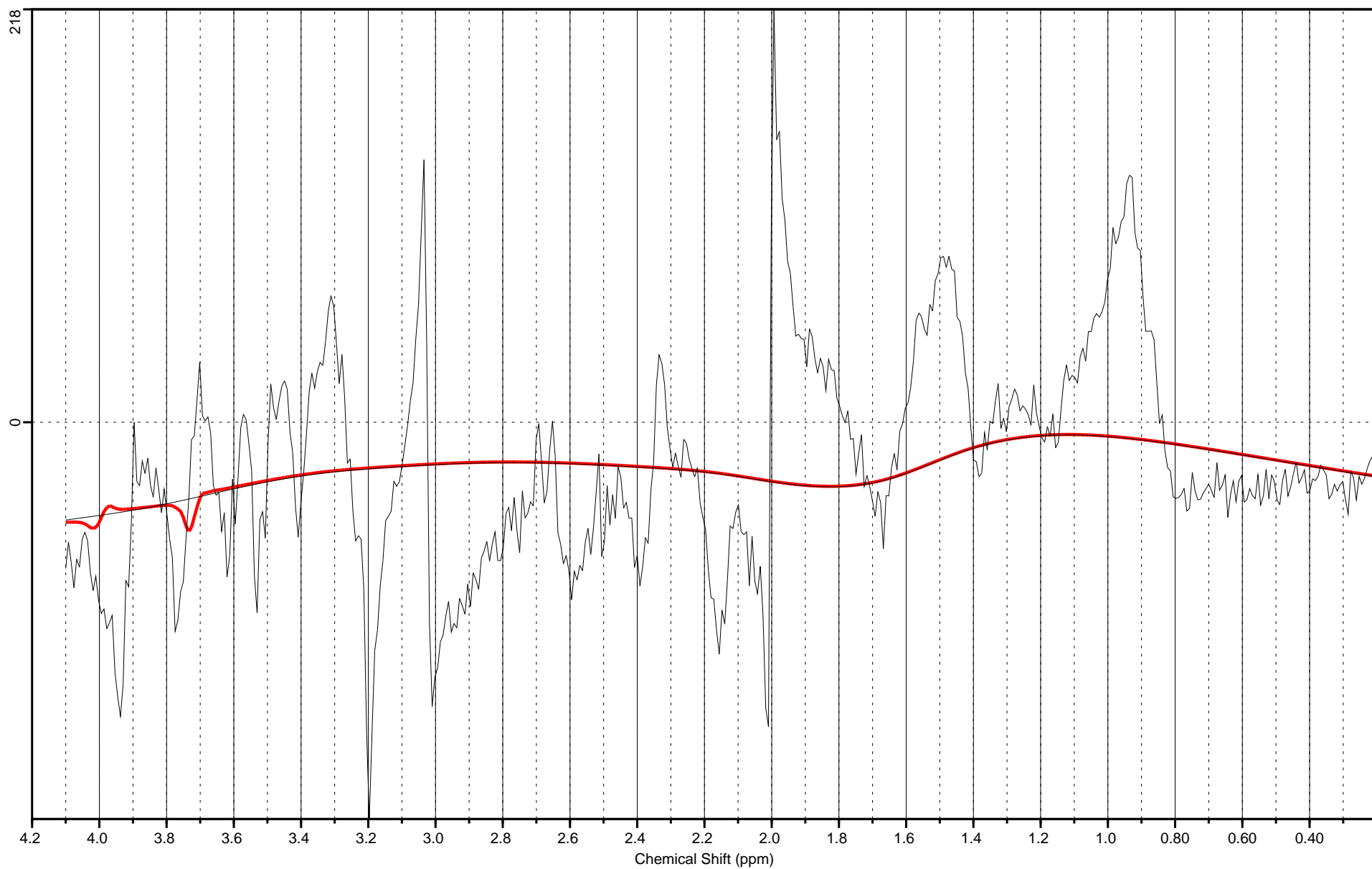
Asc Conc. = 1.73E+00

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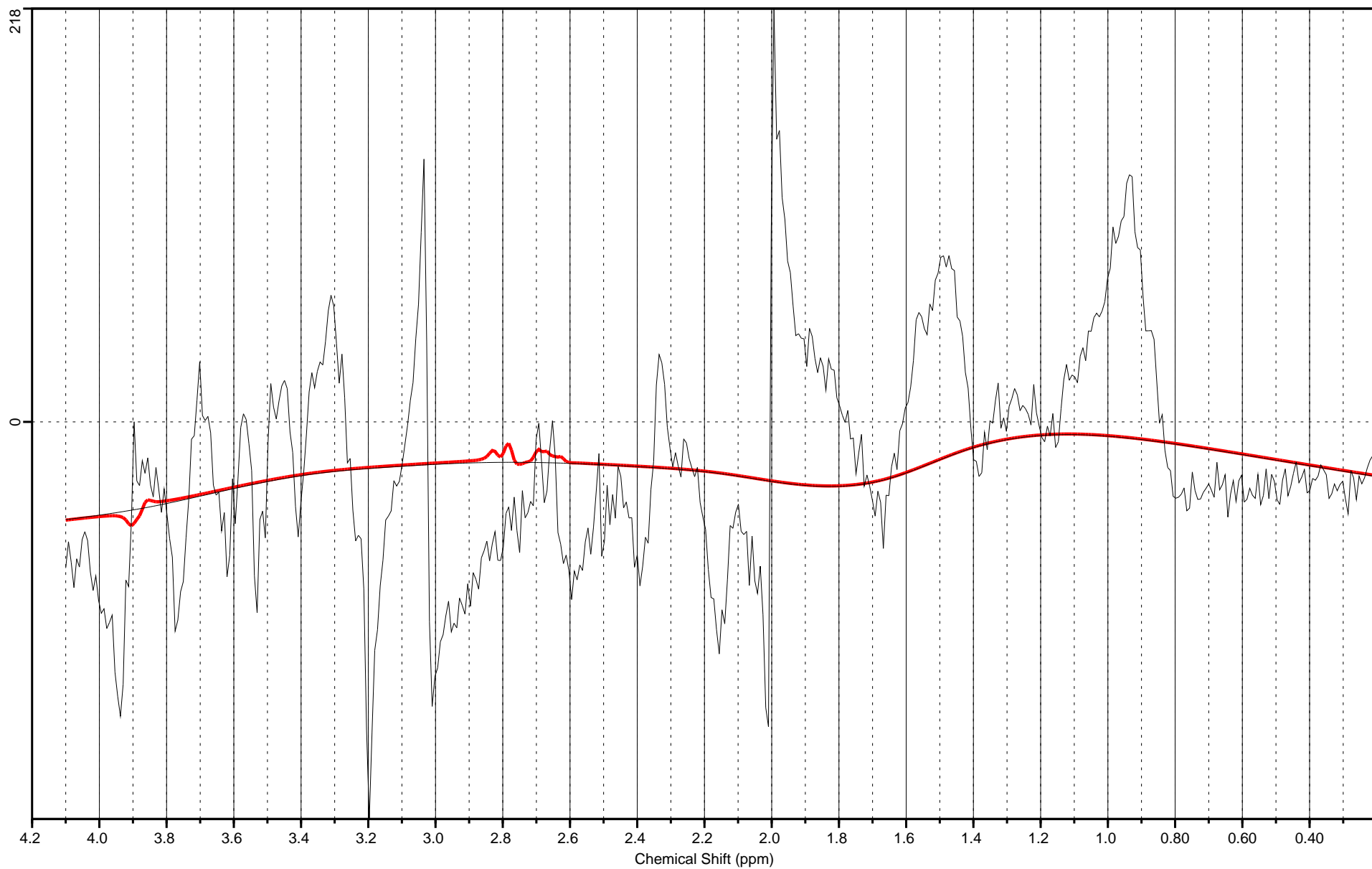
Asp Conc. = 1.91E+00

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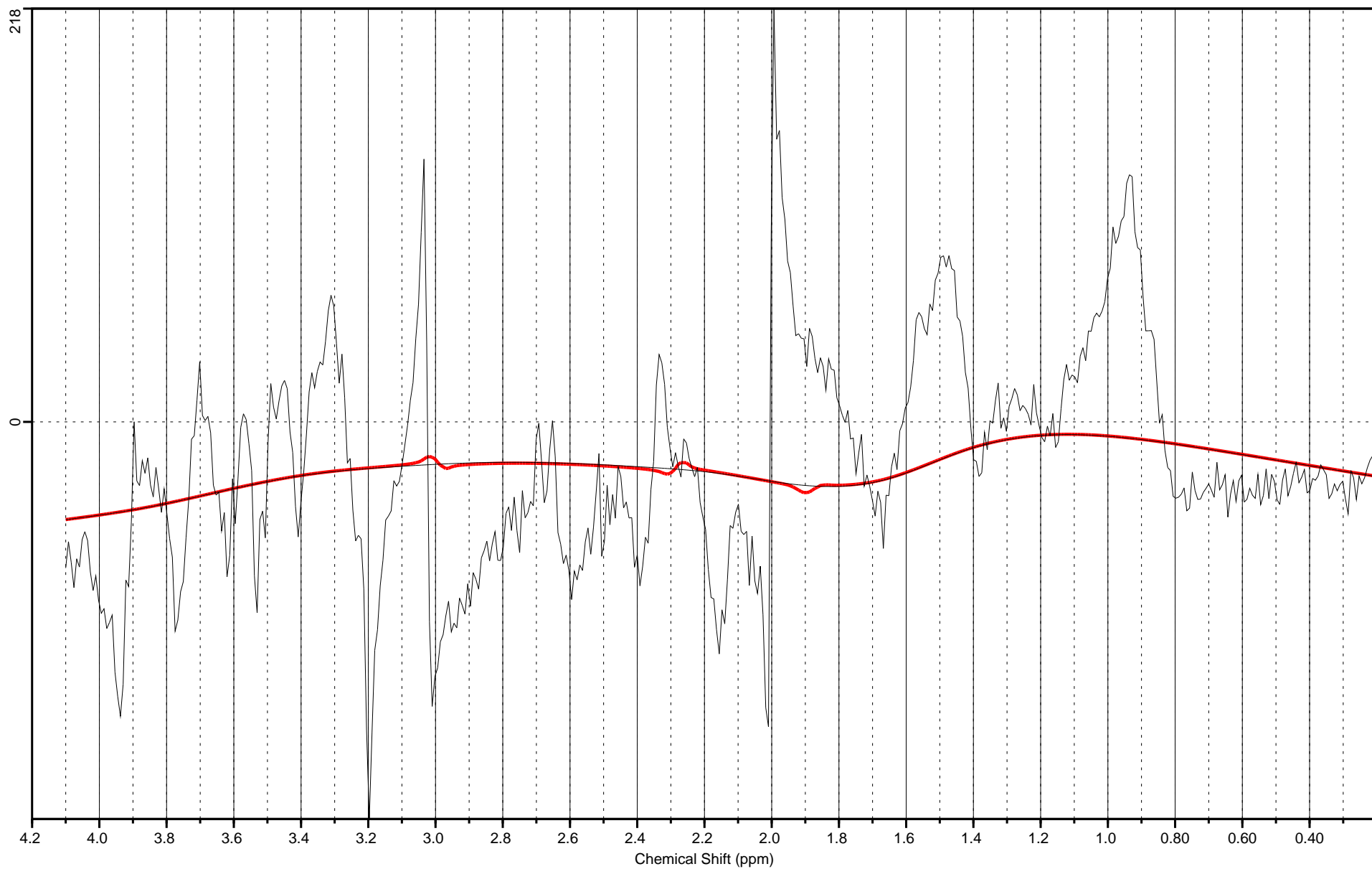
GABA Conc. = 5.40E-01

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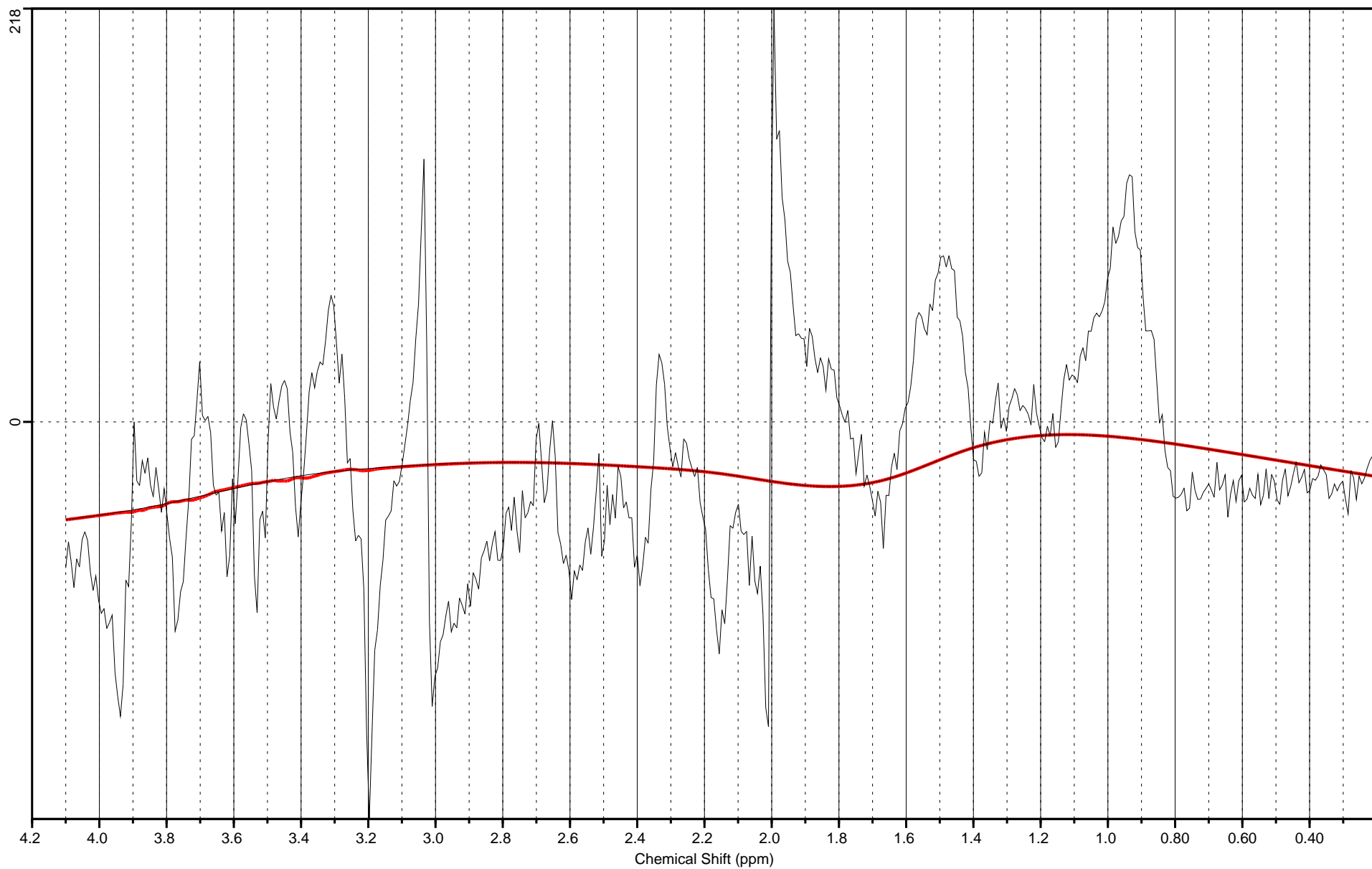
Glc Conc. = 4.04E-01

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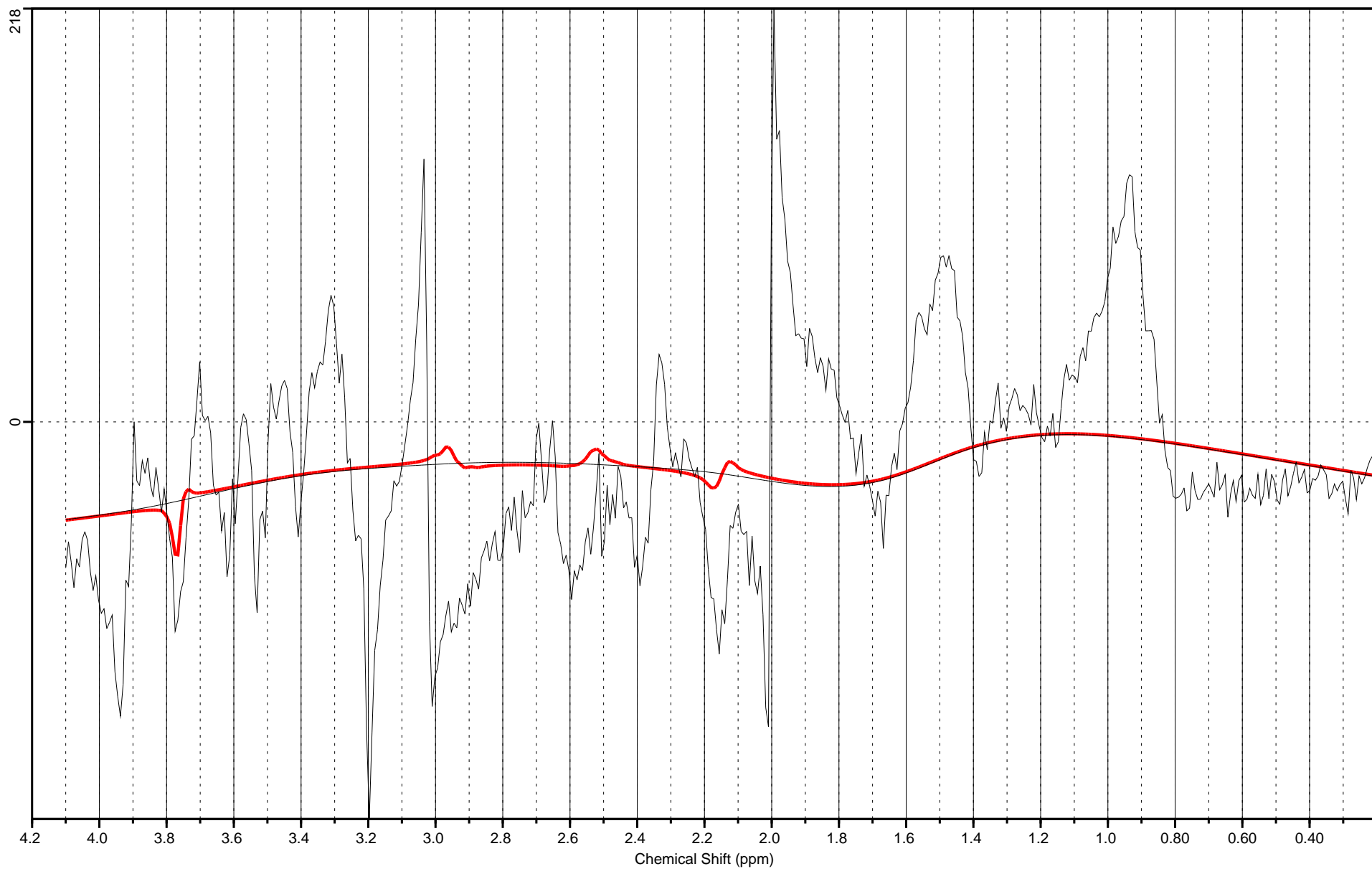
GSH Conc. = 1.36E+00

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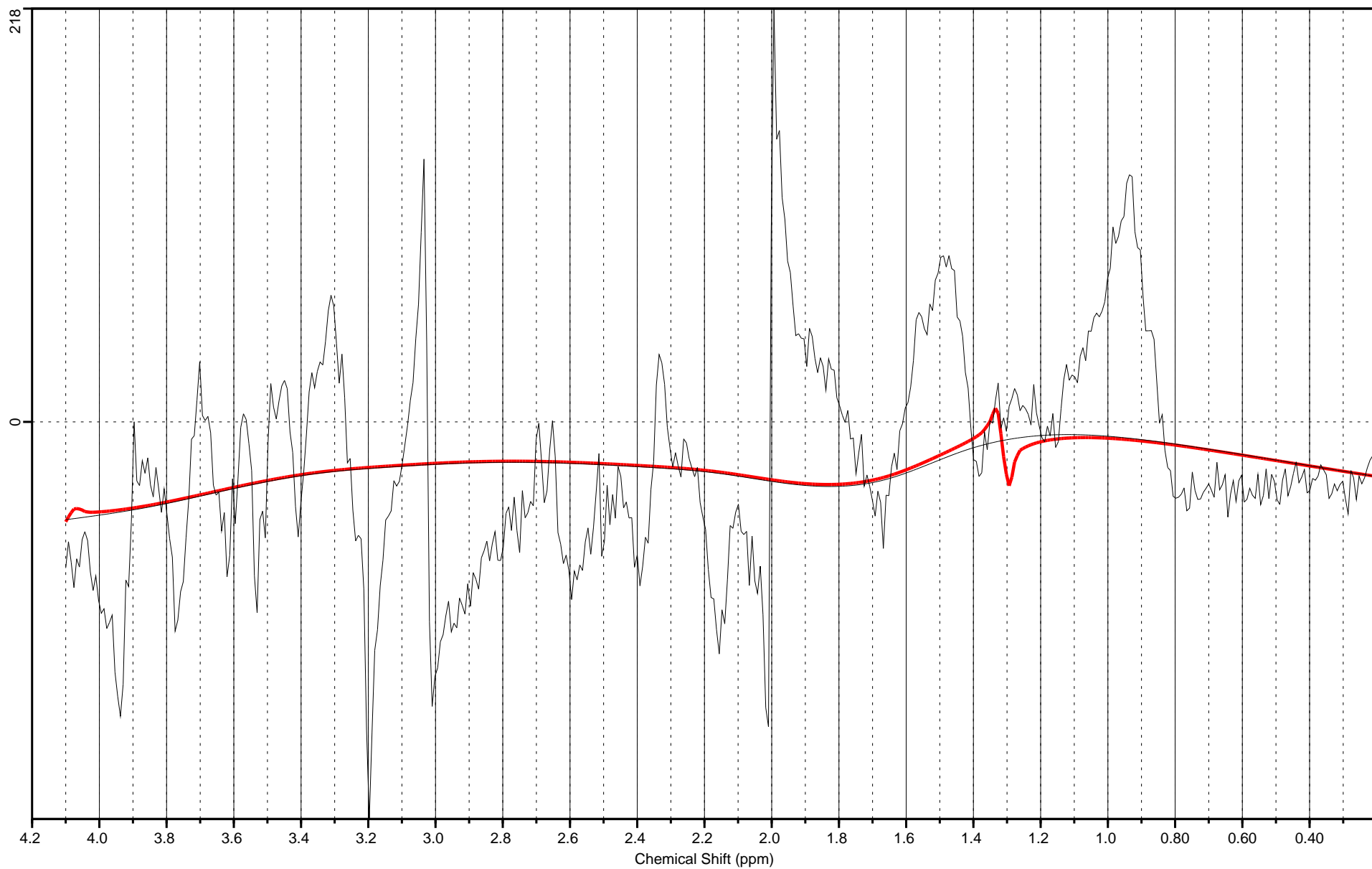
Lac Conc. = 1.85E+00

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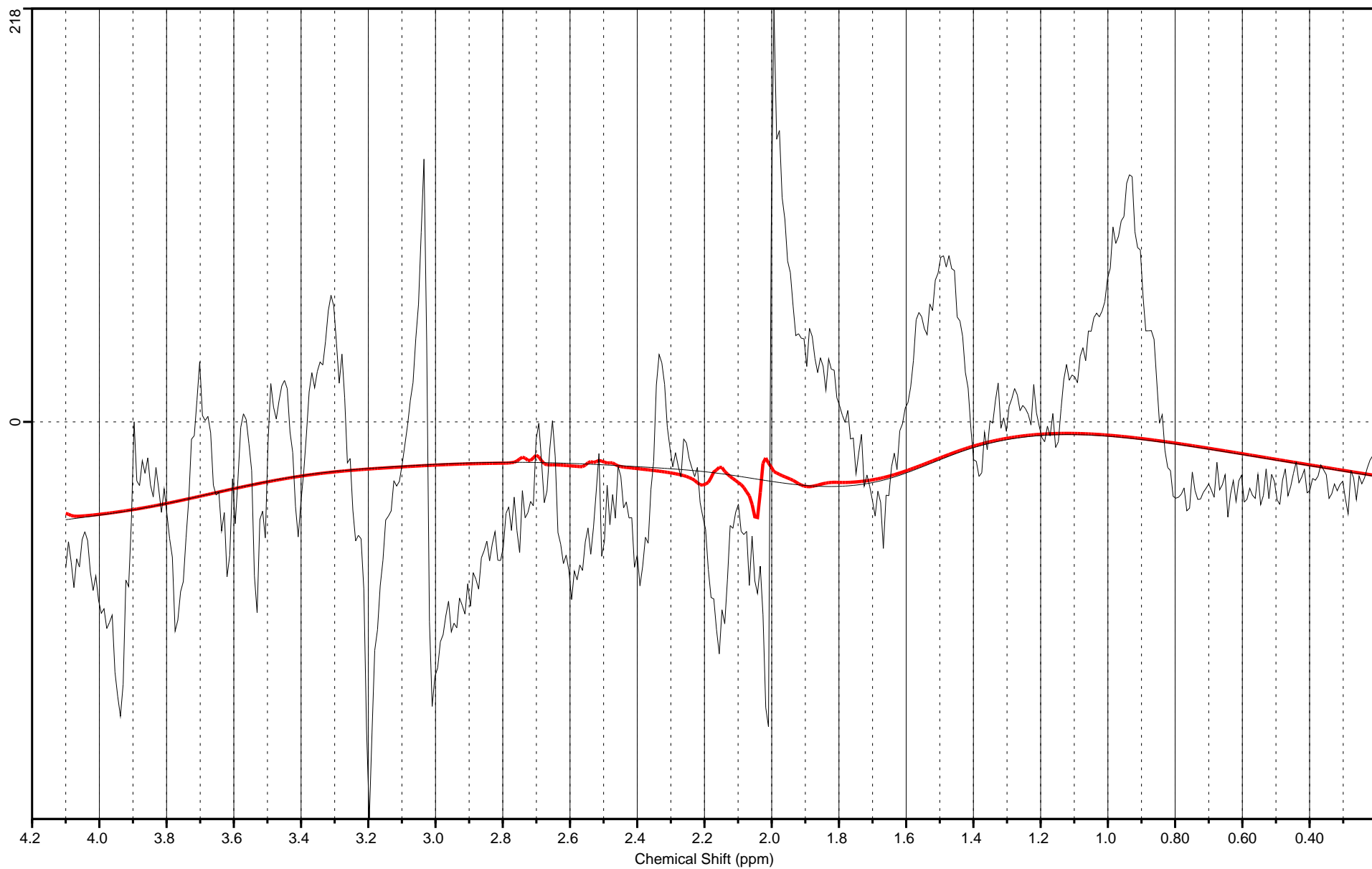
NAAG Conc. = 9.88E-01

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PE Conc. = 1.02E+00

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