

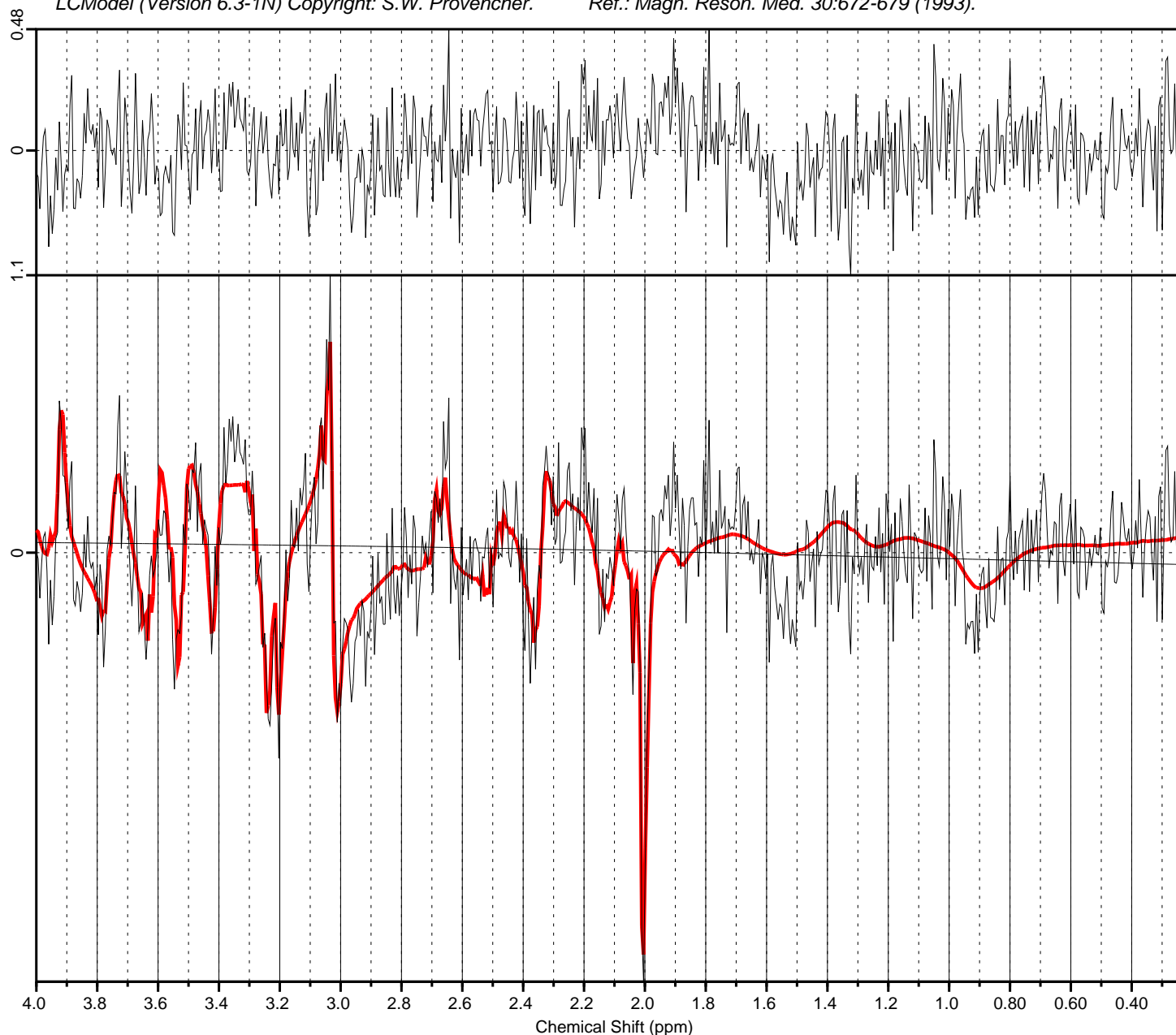
Slice_N1@19_17 02-Jun-2023 14:36:57

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Conc.	%SD	/Cr+PCr	Metabolite
1.60E-06	18%	2.2E-04	Mac
2.92E-02	27%	4.013	Cr
2.91E-02	26%	3.987	PCr
7.44E-02	10%	10.206	Ins
8.02E-02	7%	11.010	NAA
6.10E-02	11%	8.363	Tau
1.04E-02	17%	1.431	PCho
0.000	999%	0.000	GPC
6.92E-02	12%	9.498	Glu
1.29E-02	45%	1.769	Gln
0.000	999%	0.000	Ala
2.61E-02	45%	3.580	Asc
7.23E-03	87%	0.992	Asp
1.03E-02	33%	1.410	GABA
5.61E-03	73%	0.770	Glc
8.28E-03	37%	1.136	GSH
7.71E-04	815%	0.106	Lac
0.000	999%	0.000	NAAG
0.000	999%	0.000	PE
1.04E-02	17%	1.431	GPC+PCho
8.02E-02	7%	11.010	NAA+NAAG
8.21E-02	12%	11.267	Glu+Gln
5.83E-02	9%	8.000	Cr+PCr
8.02E-02	7%	11.010	NAA+NAAG
5.83E-02	9%	8.000	Cr+PCr
8.21E-02	12%	11.267	Glu+Gln

DIAGNOSTICS

3 info's RFALSI 11
 1 info RFALSI 4
 1 info FINOUT 9
 Doing Water-Scaling

MISCELLANEOUS OUTPUT

FWHM = 0.023 ppm S/N = 5
 Data shift = 0.017 ppm
 Ph: -25 deg -17.4 deg/ppm

INPUT CHANGES

hwdwat= 0.5
 wconc= 810.
 ppmst= 4.0
 ppmend= 0.2
 nunfil= 1024
 nomit= 15
 conrel=8
 namrel='Cr+PCr'

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5.83E-02	9%	8.000	Cr+PCr	
8.21E-02	12%	11.267	Glu+Gln	
DIAGNOSTICS				
3	info's	RFALSI	11	
1	info	RFALSI	4	
1	info	FINOUT	9	
Doing Water-Scaling				
MISCELLANEOUS OUTPUT				
FWHM = 0.023 ppm S/N = 5				
Data shift = 0.017 ppm				
Ph: -25 deg -17.4 deg/ppm				
INPUT CHANGES				
hwdwat= 0.5				
wconc= 810.				
ppmst= 4.0				
ppmend= 0.2				
nunfil= 1024				
nomit= 15				

```

conrel=8
namrel='Cr+PCr'
neach= 999
hzpppm= 599.419
filraw= 'Z:\Brayan\Data Processing\31052022_NewB
asis_lavgT1\Slice_N1\Data\Slice_N1@19_17.RAW'
filps= 'Z:\Brayan\Data Processing\31052022_NewBa
sis_lavgT1\Slice_N1\Data\Slice_N1@19_17.ps'
filh2o= 'Z:\Brayan\Data Processing\31052022_NewB
asis_lavgT1\Slice_N1\Data\Slice_N1@19_17w.RAW'
filbas= 'Y:\TE=1300microsec_Basis_16052023\14T_S
IM_MRSI_Dunja_Brayan_TE=1300microsec_test.BASI
S'
filcoo= 'Z:\Brayan\Data Processing\31052022_NewB
asis_lavgT1\Slice_N1\Data\Slice_N1@19_17.coord
'
filtab= 'Z:\Brayan\Data Processing\31052022_NewB
asis_lavgT1\Slice_N1\Data\tables\Slice_N1@19_1
7.table'
ltable= 7
lcoord=9
dows= T
dkntmn= 0.25
deltat= 1.40e-04
chomit= '-CrCH2' 'Gua' 'Ser' 'Lip13a' 'Lip13b' '
Lip09' 'MM09' 'Lip20' 'MM20' 'MM12' 'MM14' 'MM
17' 'Ace' 'Cit' 'bHB'
chcomb= 'GPC+PCho' 'NAA+NAAG' 'Glu+Gln' 'Cr+PCr'
atth2o= 1.0
savdir= 'Z:\Brayan\Matlab Codes\LCModel\lcmodelem
odelfiles\saved'

```

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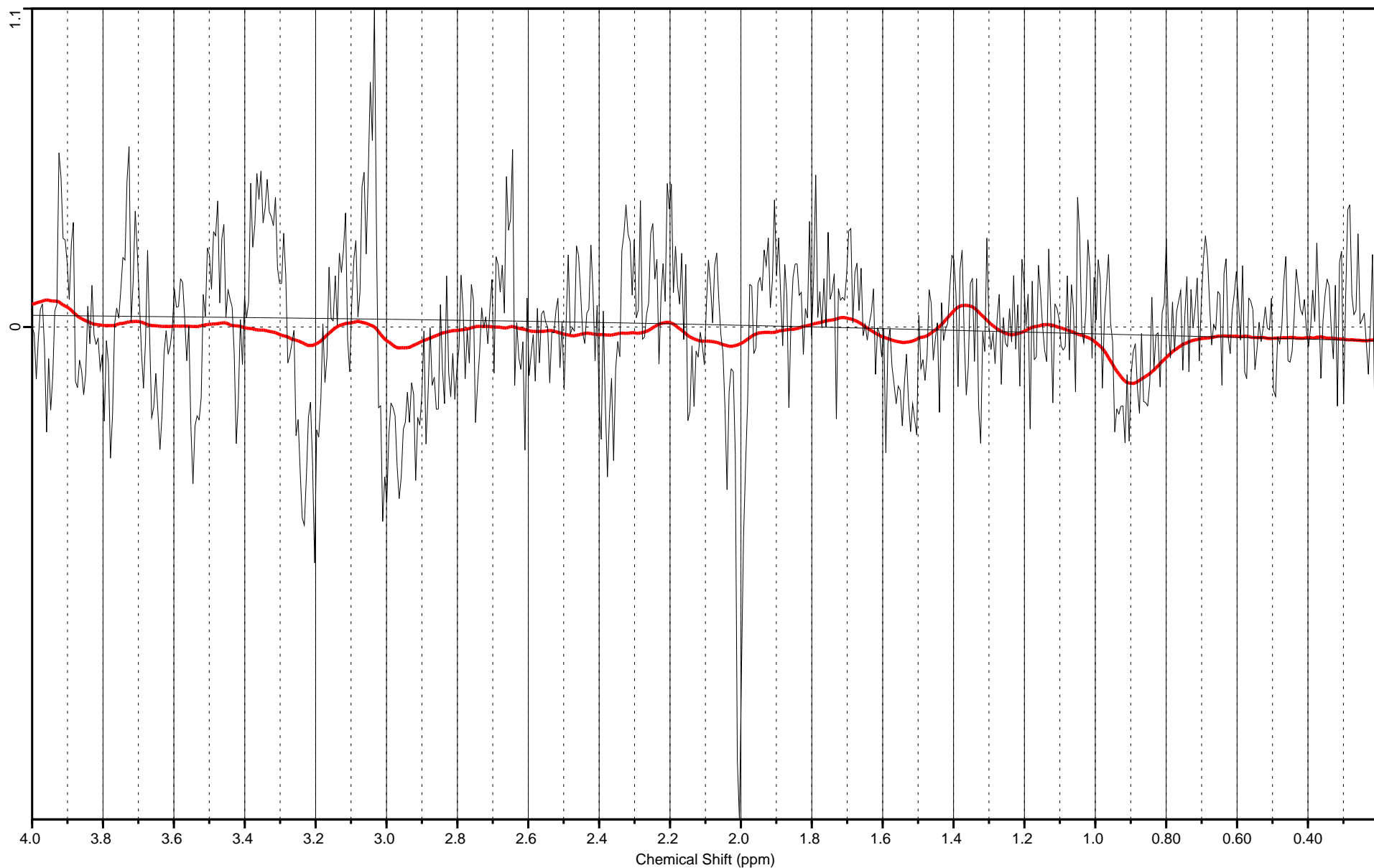
Mac Conc. = 1.60E-06

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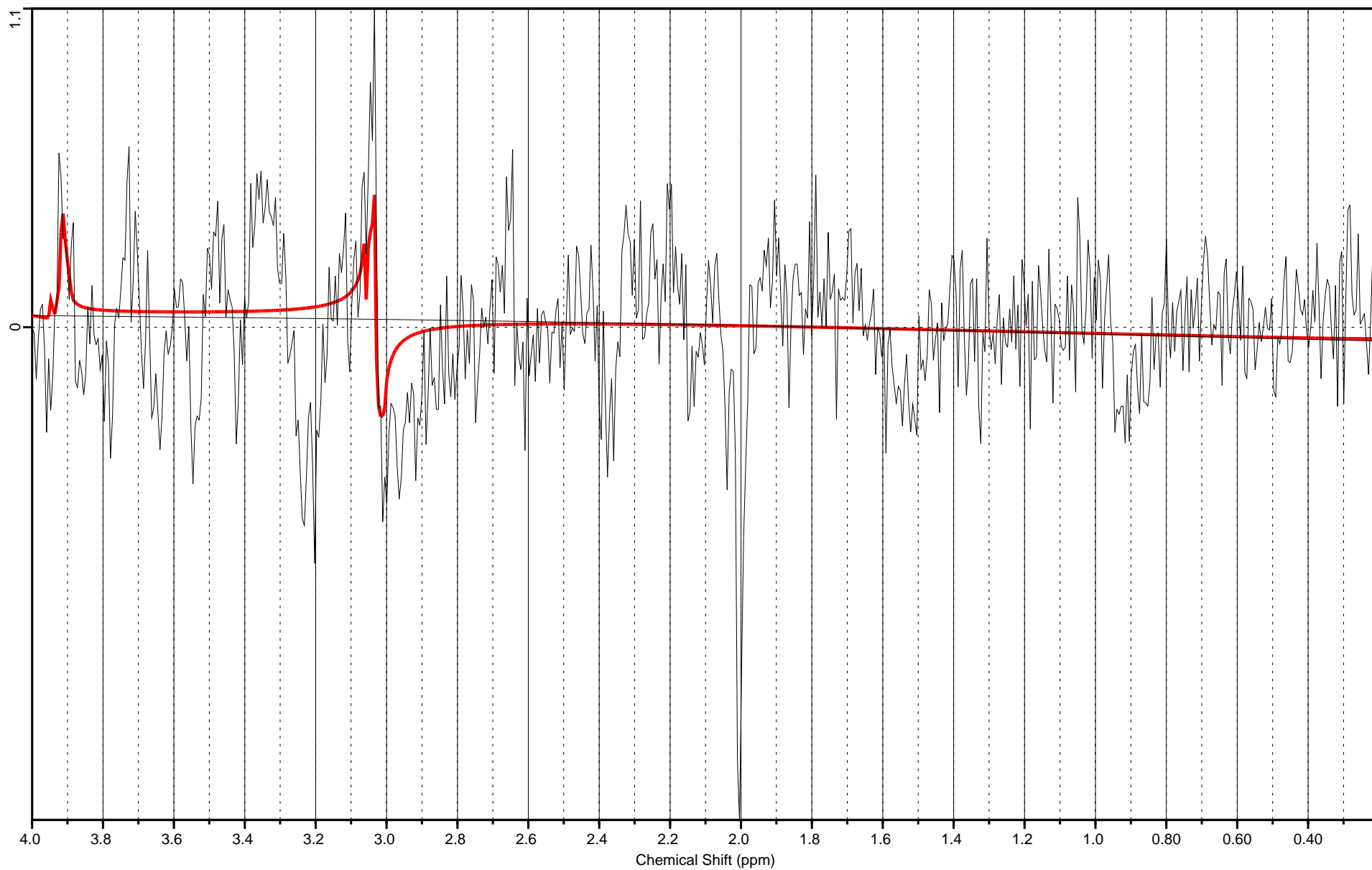
Cr Conc. = 2.92E-02

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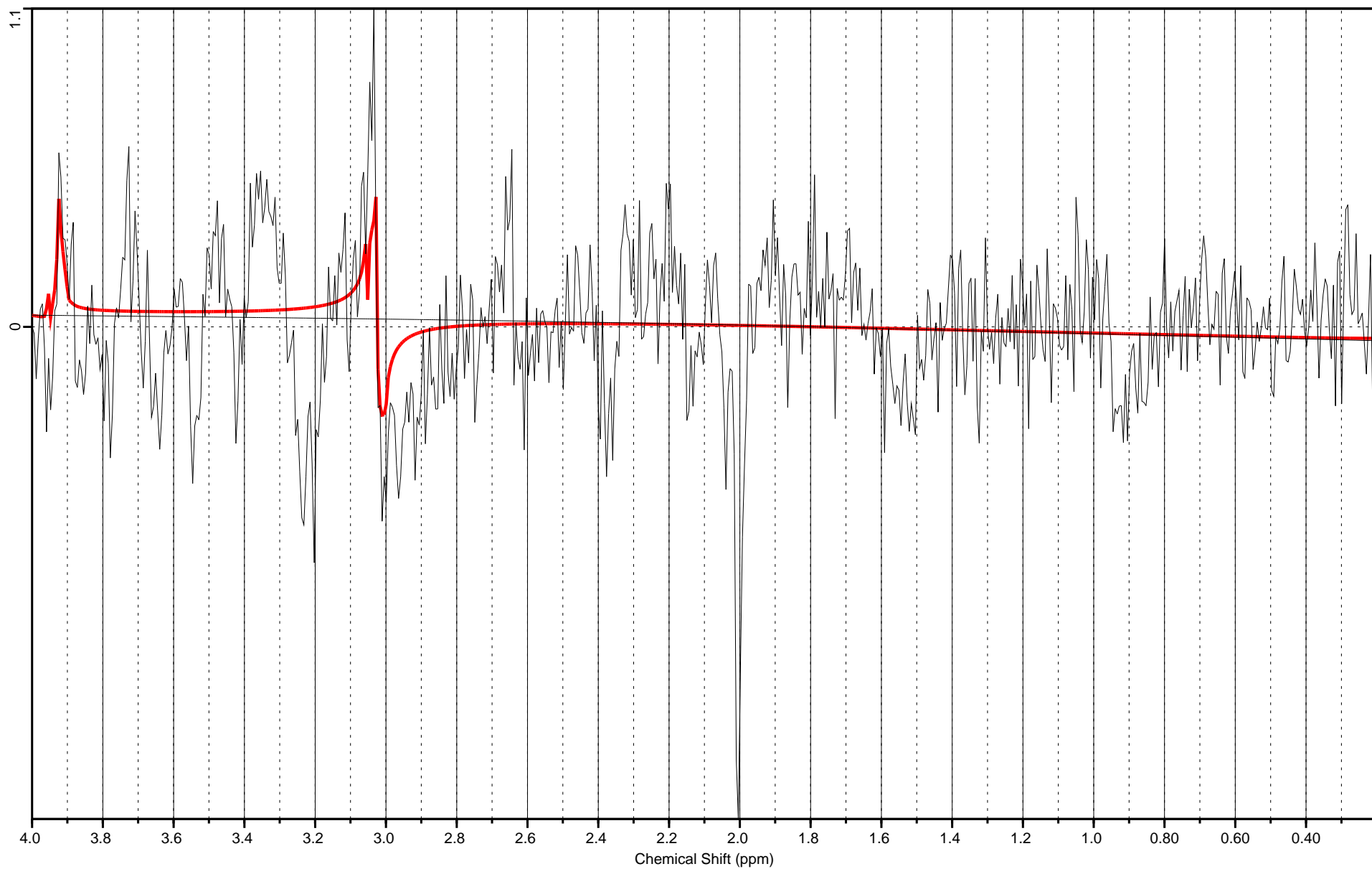
PCr Conc. = 2.91E-02

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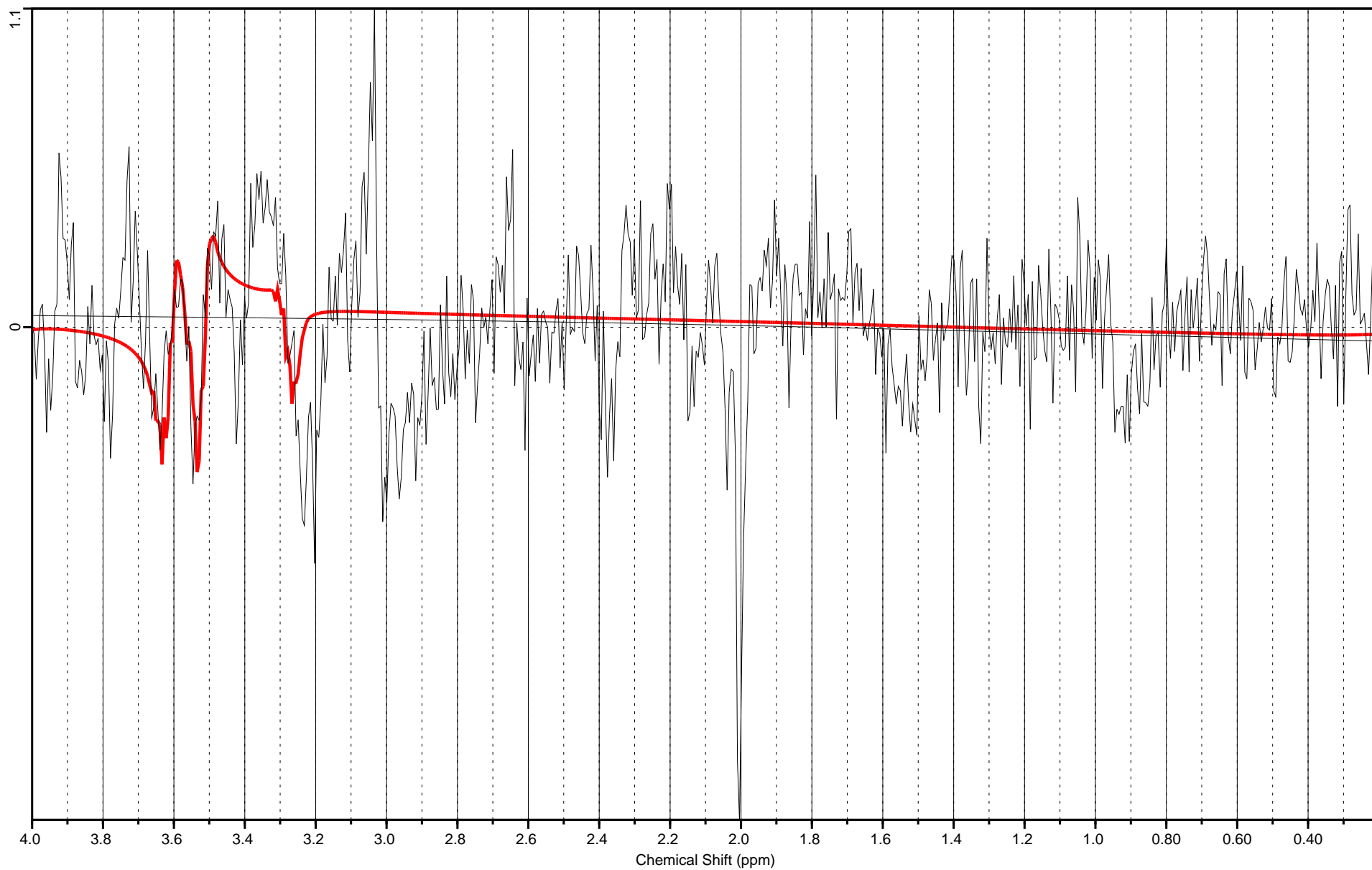
Ins Conc. = 7.44E-02

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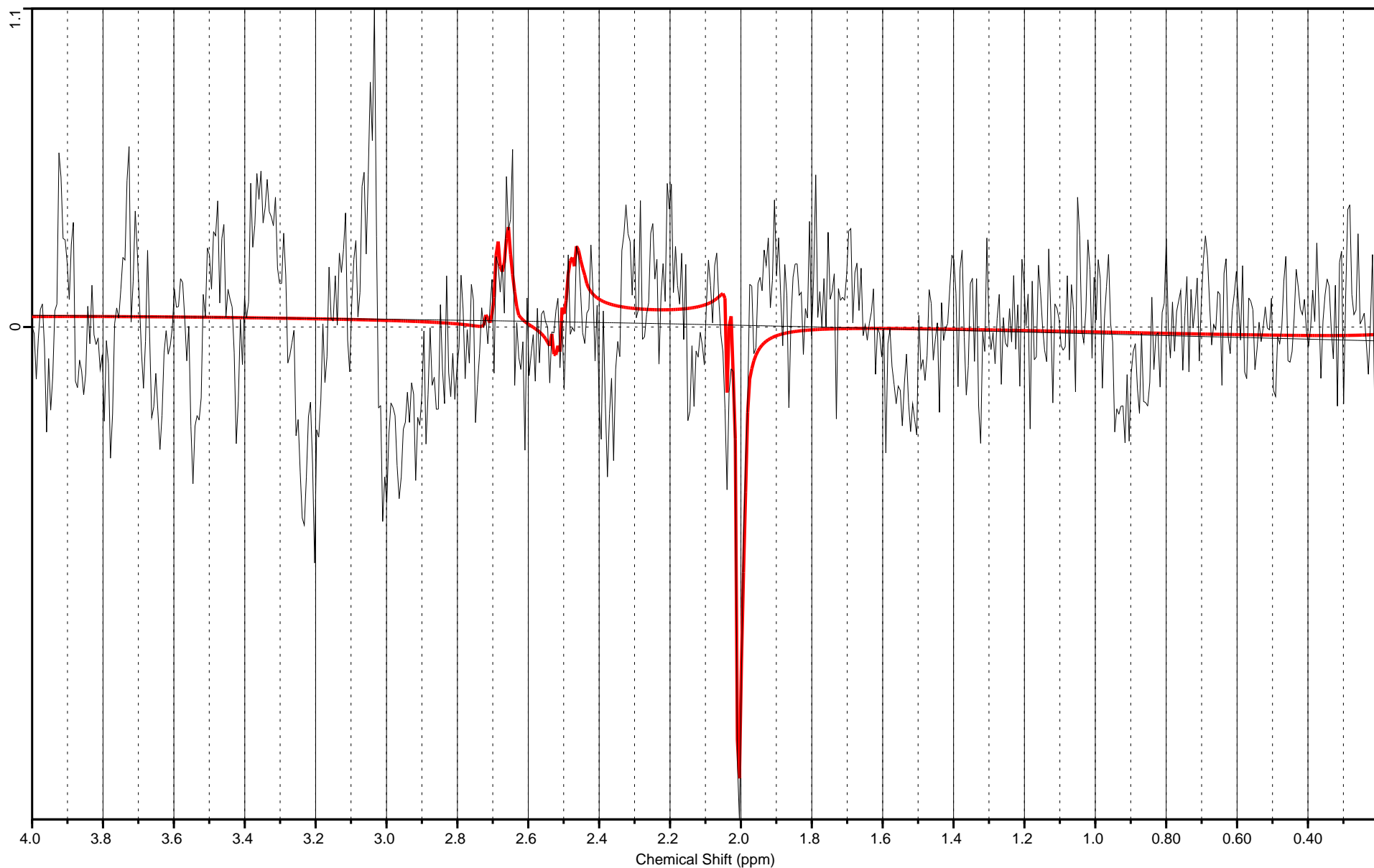
NAA Conc. = 8.02E-02

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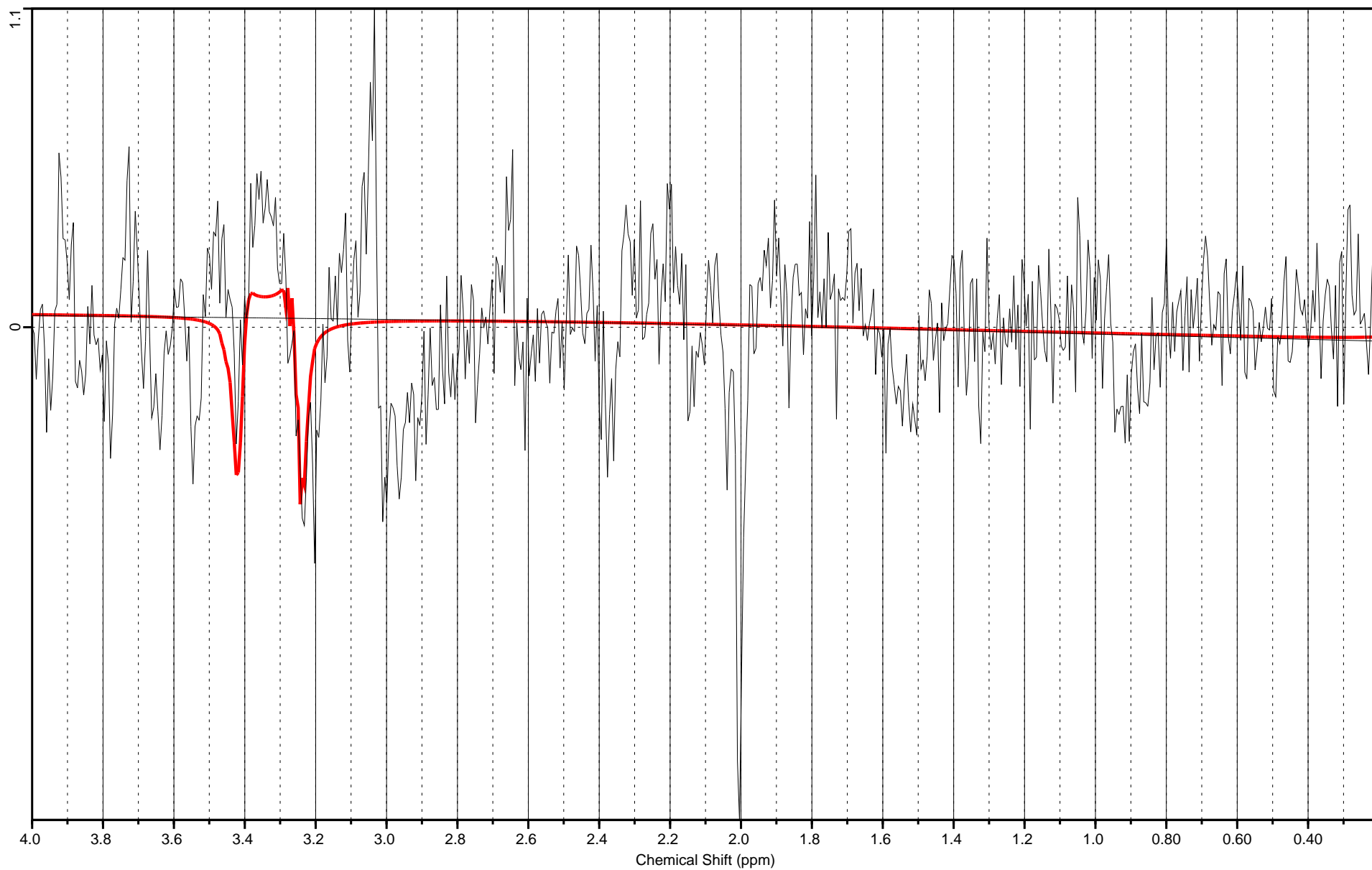
Tau Conc. = 6.10E-02

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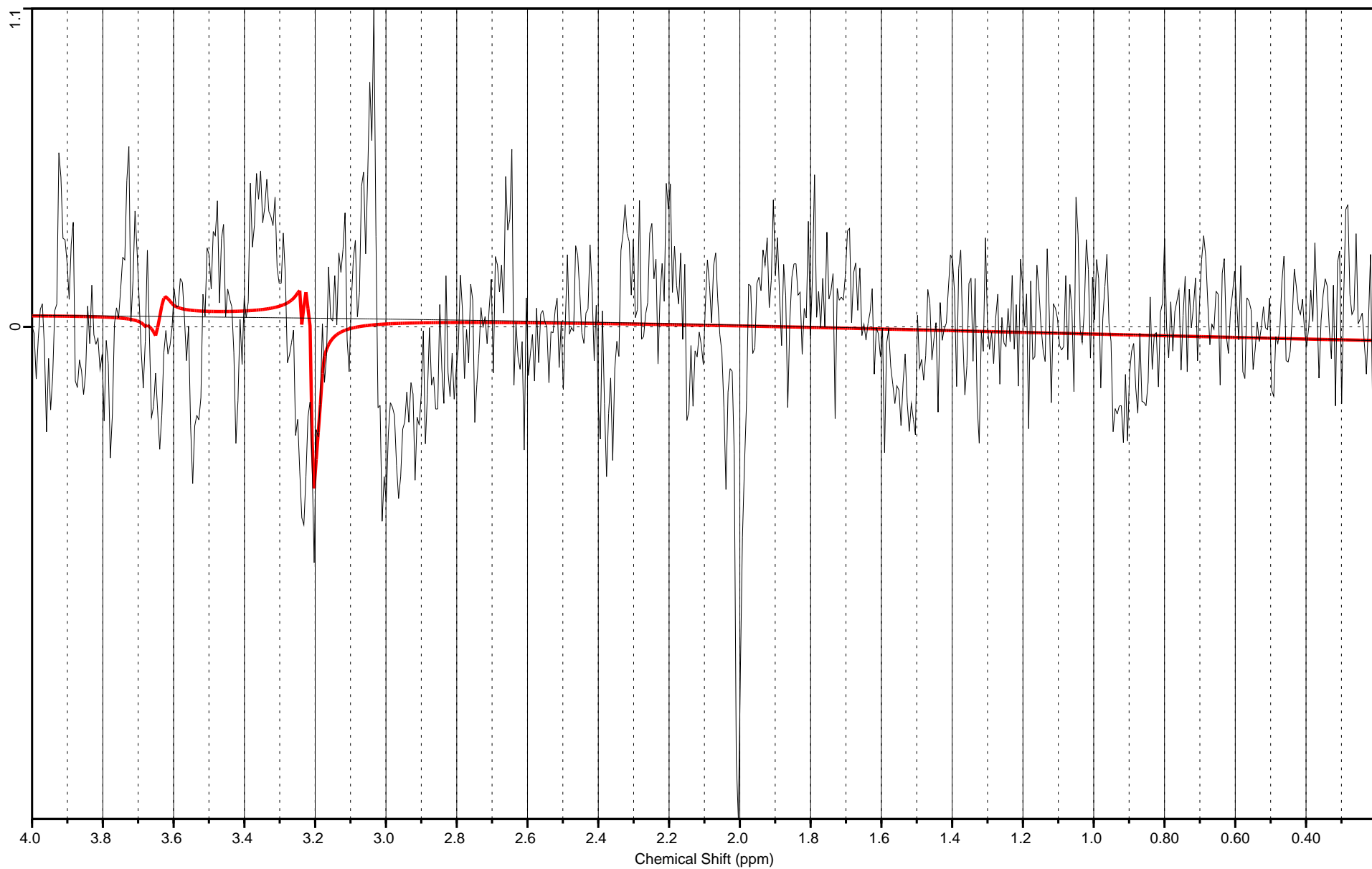
PCho Conc. = 1.04E-02

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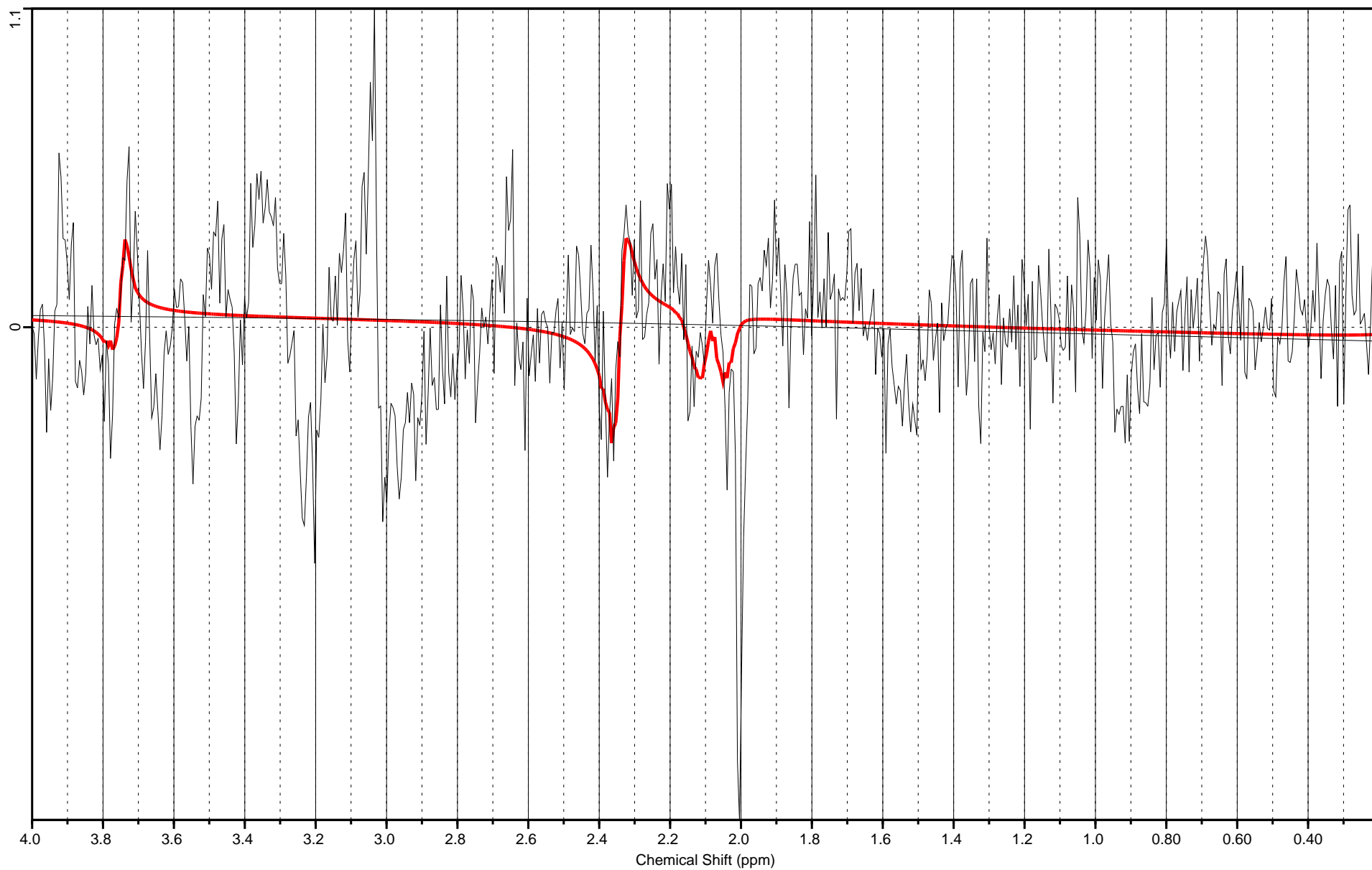
Glu Conc. = 6.92E-02

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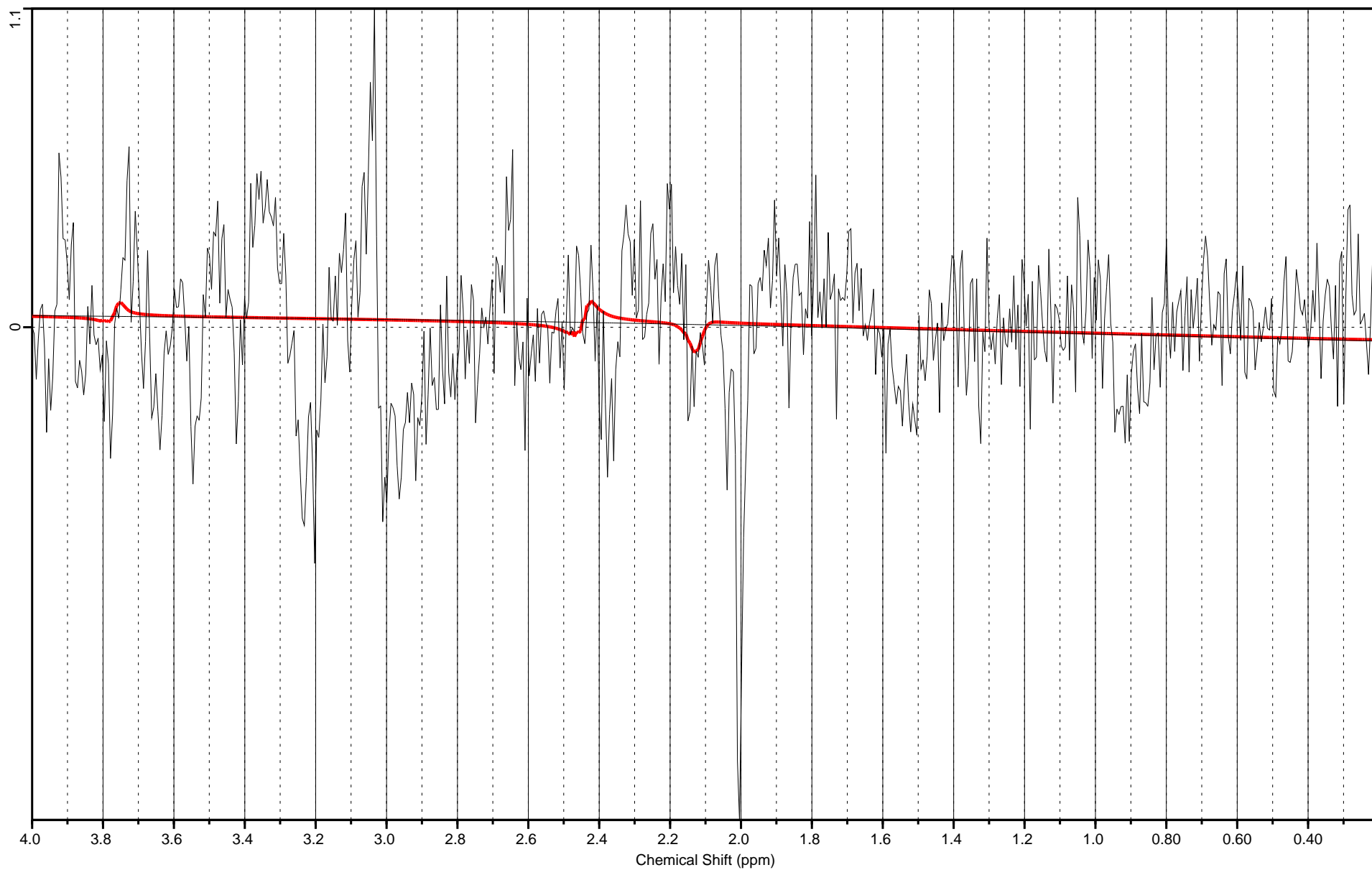
Gln Conc. = 1.29E-02

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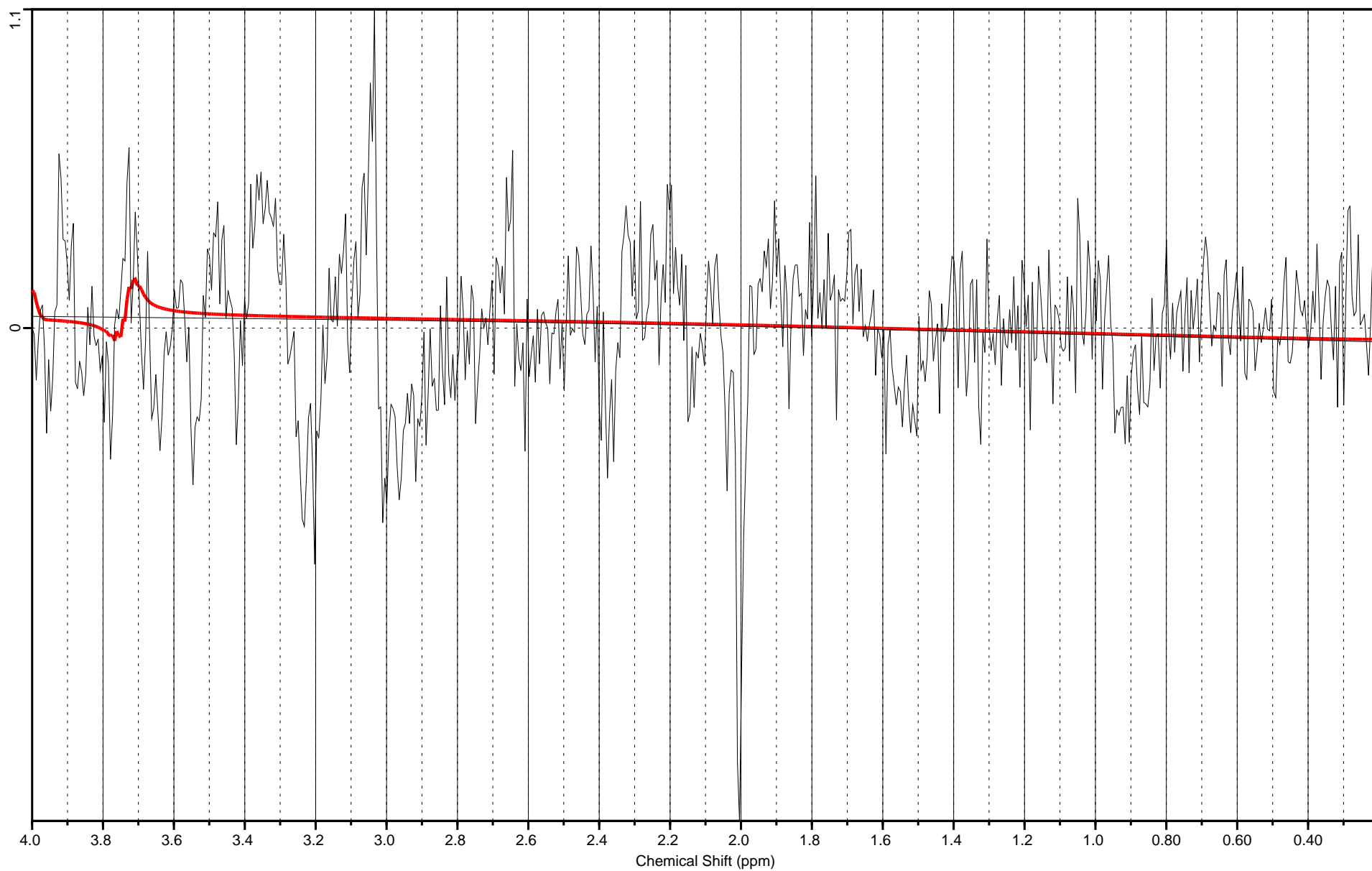
Asc Conc. = 2.61E-02

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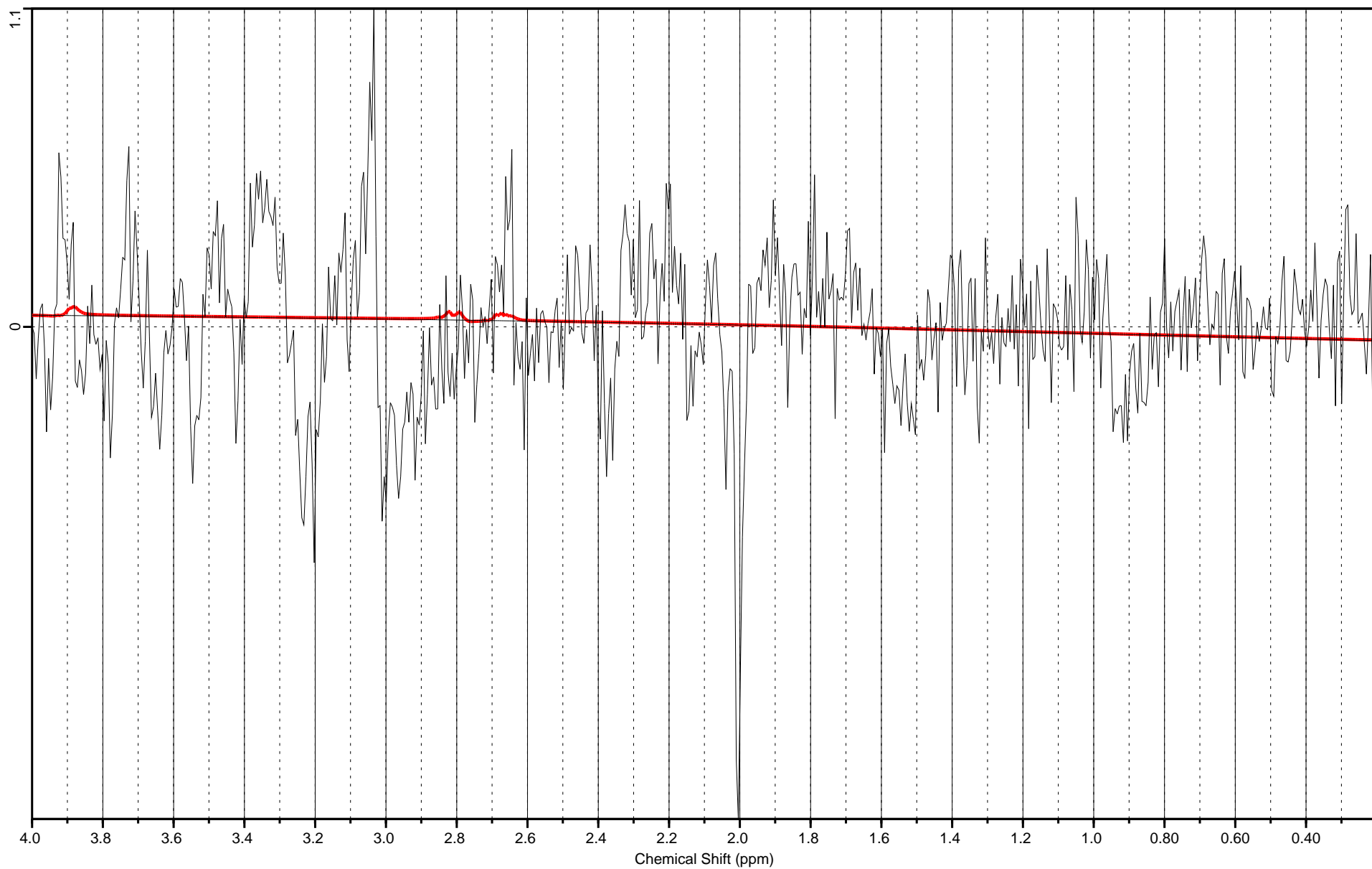
Asp Conc. = 7.23E-03

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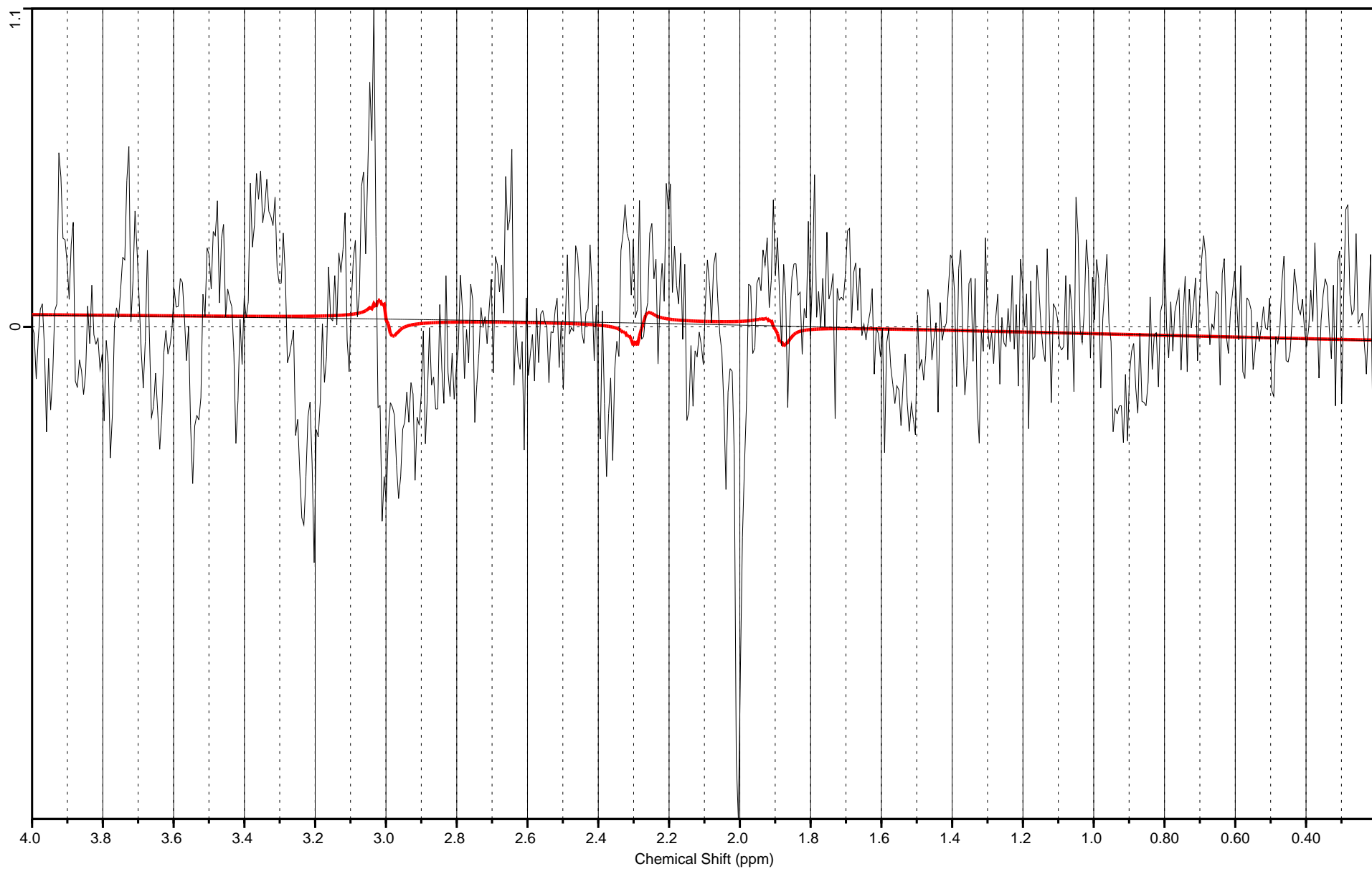
GABA Conc. = 1.03E-02

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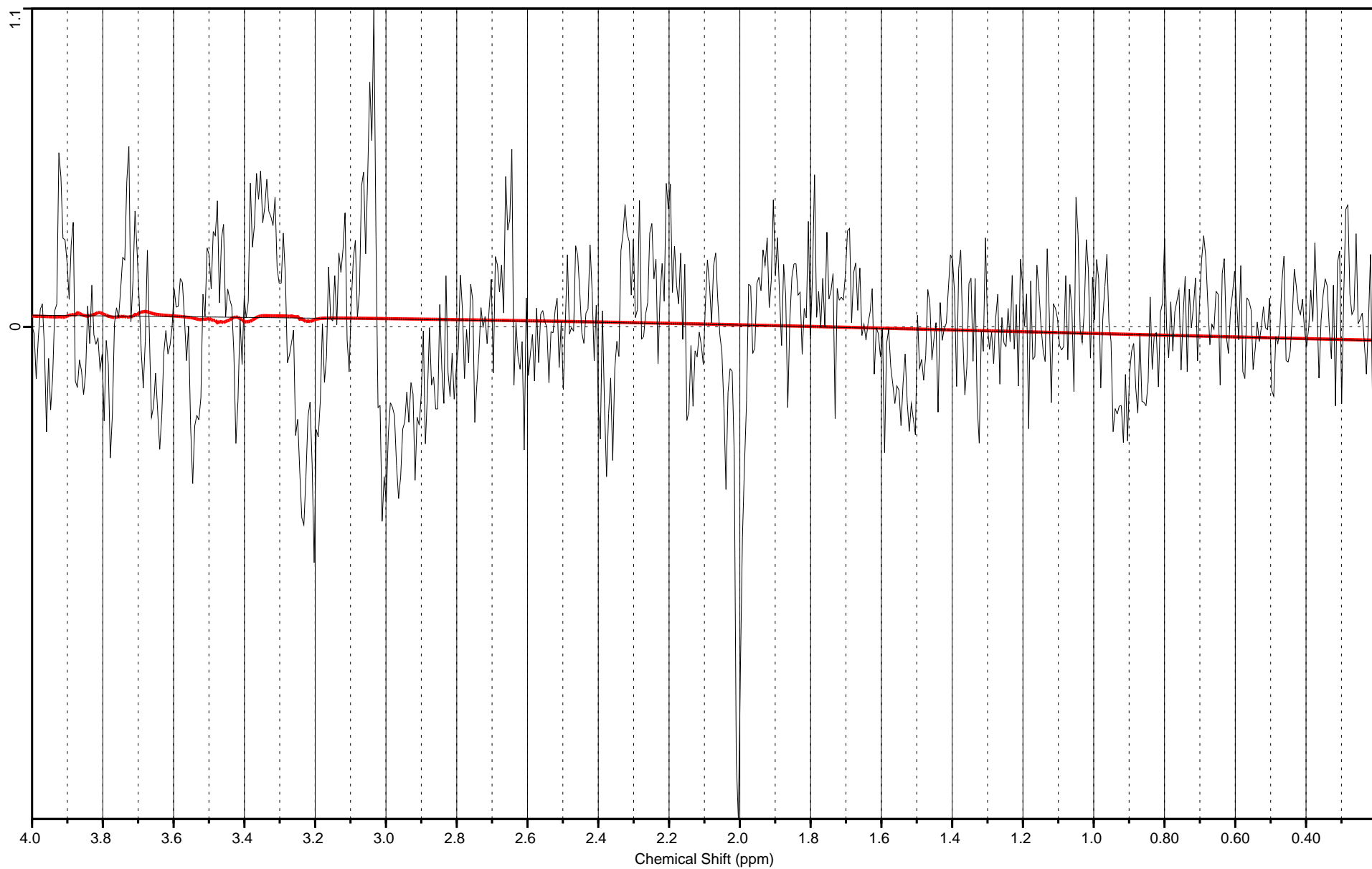
Glc Conc. = 5.61E-03

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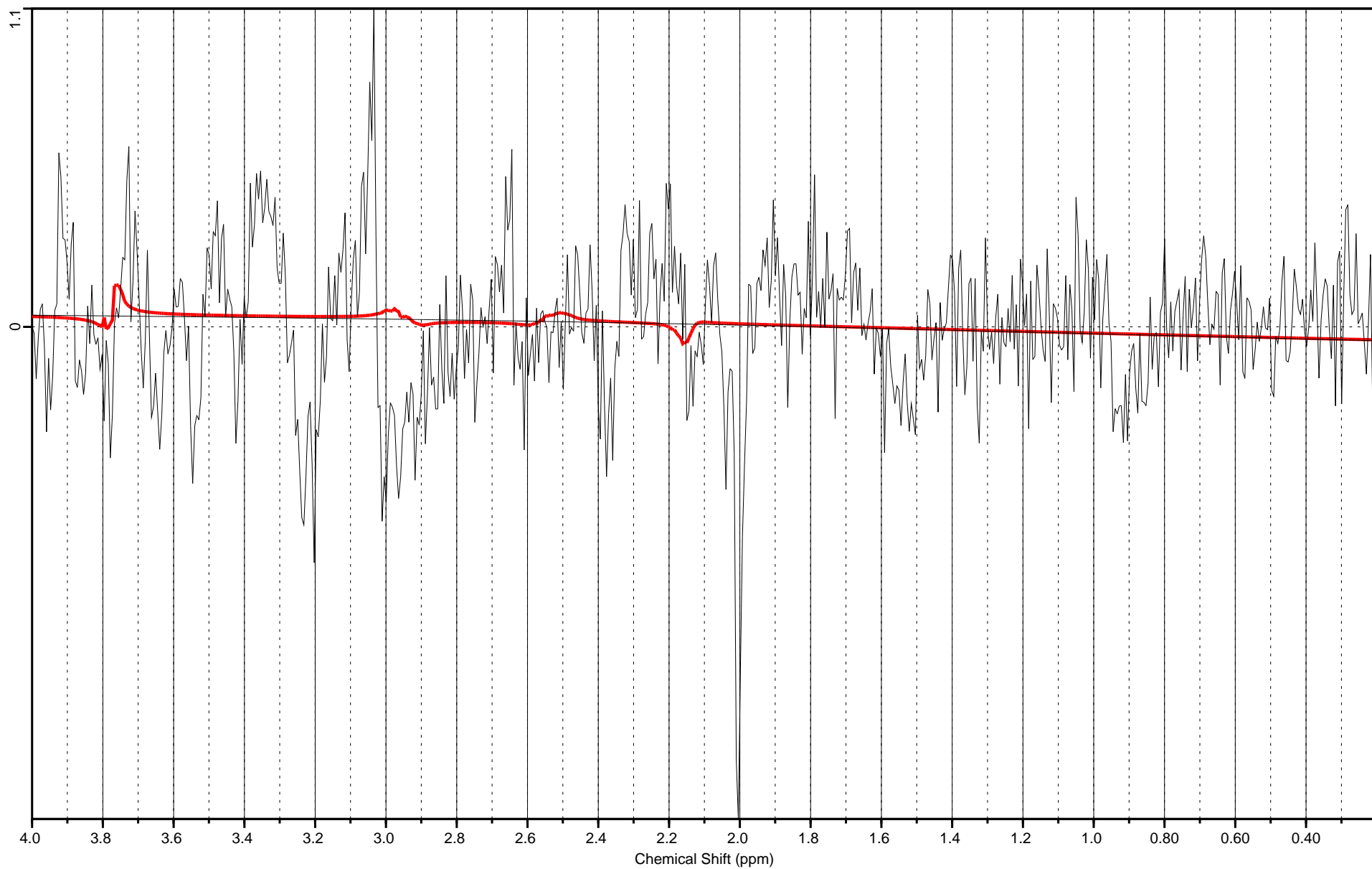
GSH Conc. = 8.28E-03

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Lac Conc. = 7.71E-04

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