

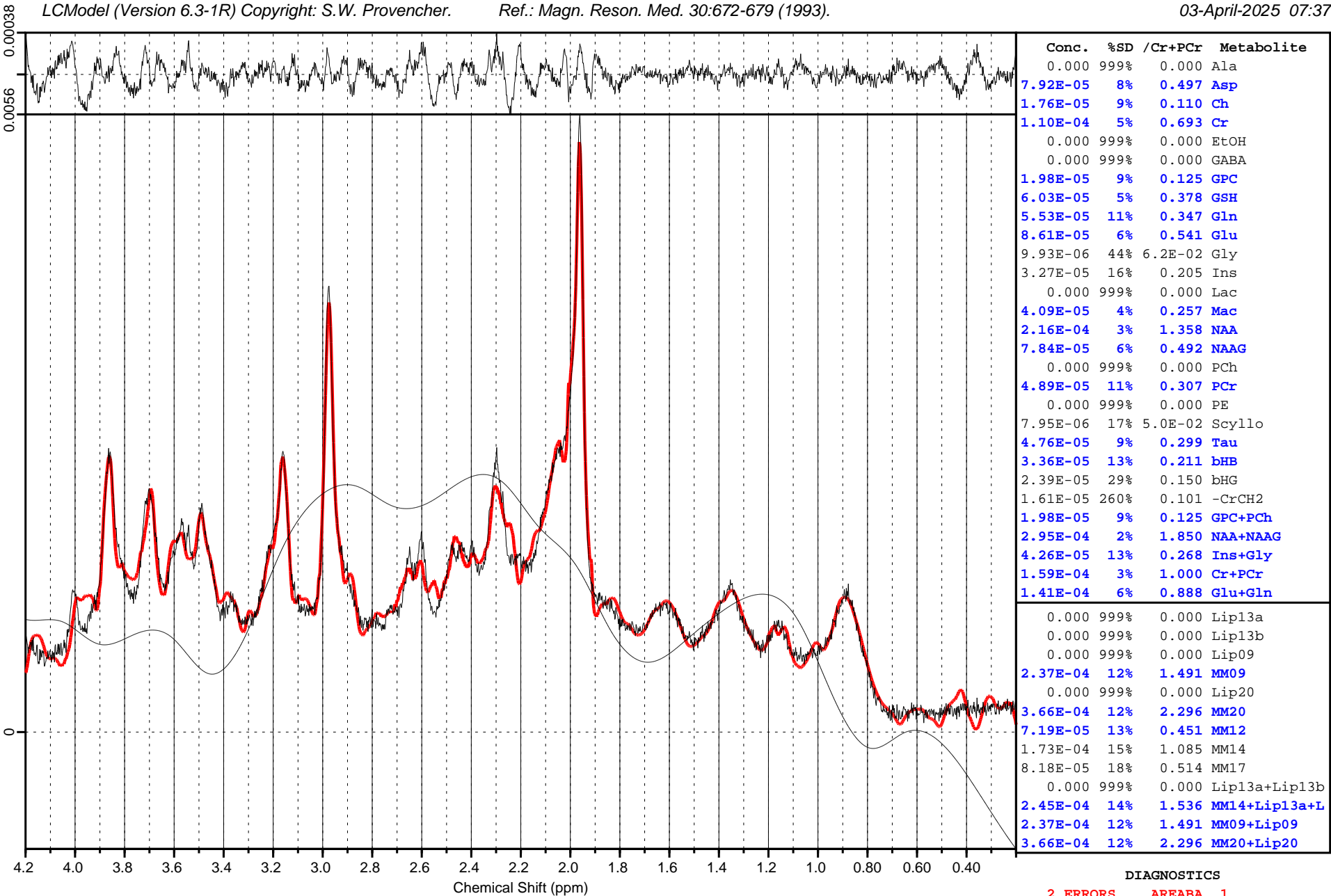
Fasting_Sub01_20241007_1150_PCC_Met

Data of: Institut fuer Diagnostische und Interventionelle Radiologie, Universitaetsklinikum Jena

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DIAGNOSTICS

2 ERRORS AREABA 1
1 info STARTV 24
1 info PNNLS 1

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Conc.	%SD	/Cr+PCr	Metabolite	2 ERRORS	AREABA	1	NDROWS=1
0.000	999%	0.000	Ala	1 info	STARTV	24	NDSLIC=1
7.92E-05	8%	0.497	Asp	1 info	PNNLS	1	FILH2O=' /data/projects/spectro-sequence-developm
1.76E-05	9%	0.110	Ch	1 info	RFALSI	12	ent/data/2024_Project_7T_BrainMrs_Jena/test_LC
1.10E-04	5%	0.693	Cr	MISCELLANEOUS OUTPUT			M/20250403_PrepapreRequest_MRShub/Raw/Fasting_
0.000	999%	0.000	EtOH	FWHM = 0.094 ppm S/N = 17			Sub01_20241007_1150_PCC_Met.RAW'
0.000	999%	0.000	GABA	Data shift =-0.033 ppm			FILRAW=' /data/projects/spectro-sequence-developm
1.98E-05	9%	0.125	GPC	Ph: -55 deg -21.1 deg/ppm			ent/data/2024_Project_7T_BrainMrs_Jena/test_LC
6.03E-05	5%	0.378	GSH	INPUT CHANGES			M/20250403_PrepapreRequest_MRShub/Raw/Fasting_
5.53E-05	11%	0.347	Gln	nameac(20)='Tau'			Sub01_20241007_1150_PCC_Met.RAW'
8.61E-05	6%	0.541	Glu	nameac(19)='NAAG'			FILBAS=' /data/projects/spectro-sequence-developm
9.93E-06	44%	6.2E-02	Gly	nameac(18)='NAA'			ent/data/2024_Project_7T_BrainMrs_Jena/test_LC
3.27E-05	16%	0.205	Ins	nameac(17)='PE'			M/20250403_PrepapreRequest_MRShub/steam_7T_TE6
0.000	999%	0.000	Lac	nameac(16)='PCho'			.BASIS'
4.09E-05	4%	0.257	Mac	nameac(15)='PCr'			filtab=' /data/projects/spectro-sequence-developm
2.16E-04	3%	1.358	NAA	nameac(14)='Lac'			ent/data/2024_Project_7T_BrainMrs_Jena/test_LC
7.84E-05	6%	0.492	NAAG	nameac(13)='sIns'			M/20250403_PrepapreRequest_MRShub/Results/Fast
0.000	999%	0.000	PCh	nameac(12)='Ins'			ing_Sub01_20241007_1150_PCC_Met.table'
4.89E-05	11%	0.307	PCr	nameac(11)='GSH'			ltable=7
0.000	999%	0.000	PE	nameac(10)='GPC'			FILCOO=' /data/projects/spectro-sequence-developm
7.95E-06	17%	5.0E-02	Scyllo	nameac(9)='Glu'			ent/data/2024_Project_7T_BrainMrs_Jena/test_LC
4.76E-05	9%	0.299	Tau	nameac(8)='Gln'			M/20250403_PrepapreRequest_MRShub/Results/Fast
3.36E-05	13%	0.211	bHB	nameac(7)='Glc'			ing_Sub01_20241007_1150_PCC_Met.coord'
2.39E-05	29%	0.150	bHG	nameac(6)='GABA'			LCOORD=9
1.61E-05	260%	0.101	-CrCH2	nameac(5)='Cr'			FILCSV=' /data/projects/spectro-sequence-developm
1.98E-05	9%	0.125	GPC+PCh	nameac(4)='Asc'			ent/data/2024_Project_7T_BrainMrs_Jena/test_LC
2.95E-04	2%	1.850	NAA+NAAG	nameac(3)='Asp'			M/20250403_PrepapreRequest_MRShub/Results/Fast
4.26E-05	13%	0.268	Ins+Gly	nameac(2)='Ala'			ing_Sub01_20241007_1150_PCC_Met.csv'
1.59E-04	3%	1.000	Cr+PCr	nameac(1)='Mac'			LCSV=11
1.41E-04	6%	0.888	Glu+Gln	neach=20			FILPS=' /data/projects/spectro-sequence-developme
0.000	999%	0.000	Lip13a	nunfil=8224			nt/data/2024_Project_7T_BrainMrs_Jena/test_LCM
0.000	999%	0.000	Lip13b	wconc=1			/20250403_PrepapreRequest_MRShub/Results/Fasti
0.000	999%	0.000	Lip09	atth2o=1			ng_Sub01_20241007_1150_PCC_Met.ps'
2.37E-04	12%	1.491	MM09	PPMST=4.2			LPS=8
0.000	999%	0.000	Lip20	PPMEND=0.2			ECHOT=5.52
3.66E-04	12%	2.296	MM20	doecc=F			HZPPPM=2.9717e+02
7.19E-05	13%	0.451	MM12	dows=F			DELTAT=0.0001
1.73E-04	15%	1.085	MM14	attmet=1			NUNFIL=8224
8.18E-05	18%	0.514	MM17	ICOLEN=1			KEY(1)=210387309
0.000	999%	0.000	Lip13a+Lip13b	ICOLST=1			
2.45E-04	14%	1.536	MM14+Lip13a+Lip13b+MM12	IROWEN=1			
2.37E-04	12%	1.491	MM09+Lip09	IROWST=1			
3.66E-04	12%	2.296	MM20+Lip20	ISLICE=1			
DIAGNOSTICS				NDCOLS=1			

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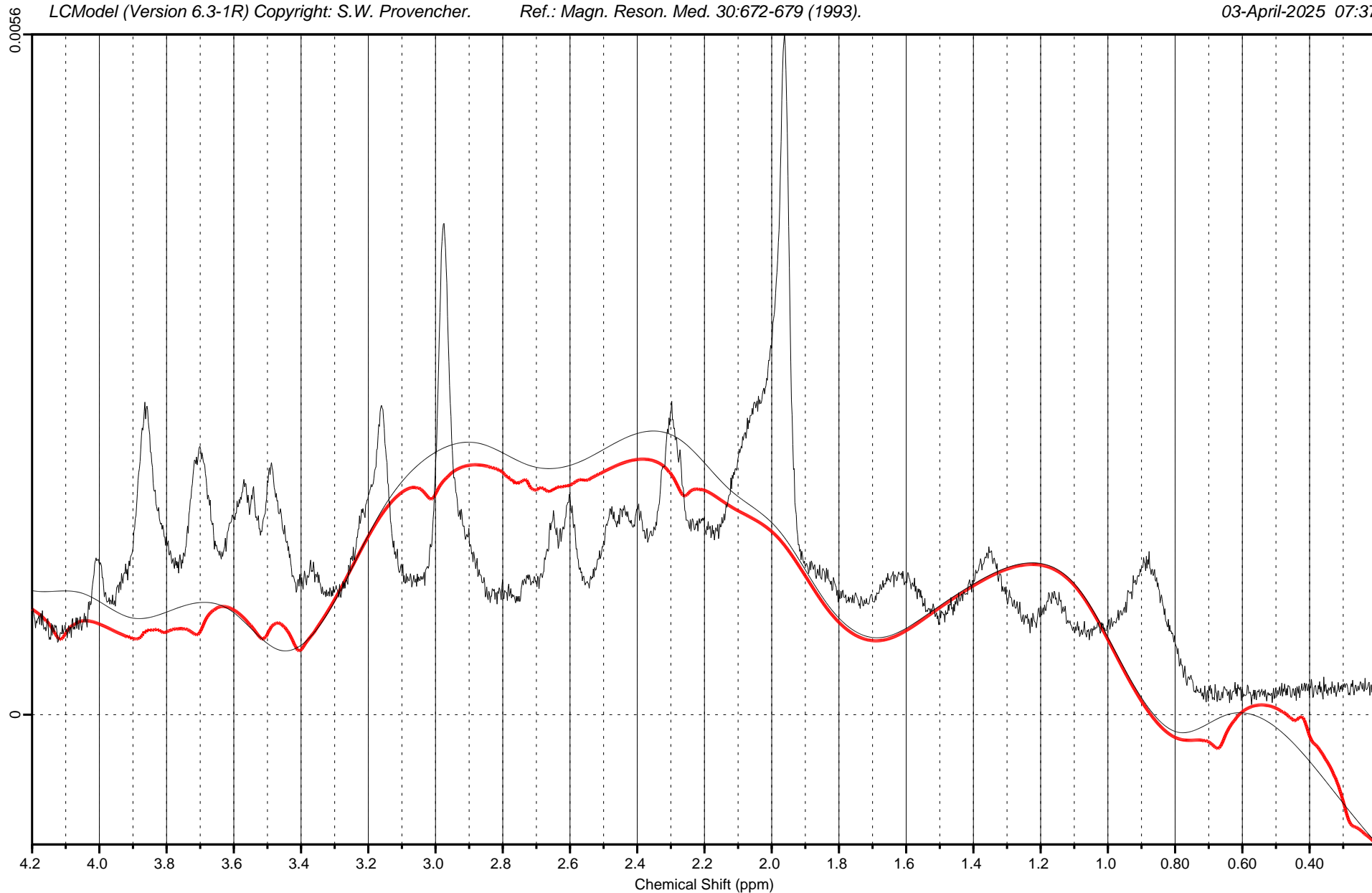
Asp Conc. = 7.92E-05

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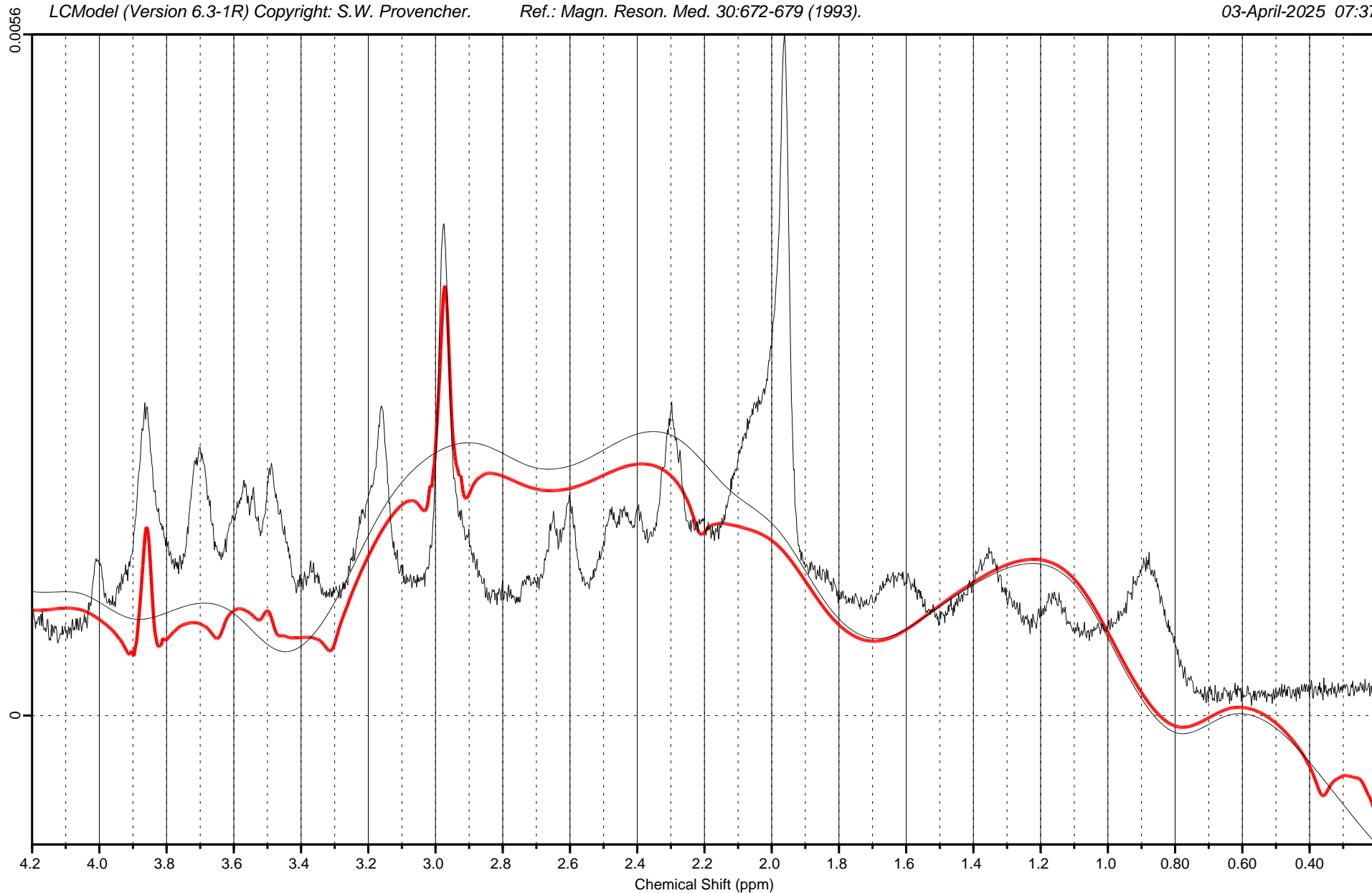
Cr Conc. = 1.10E-04

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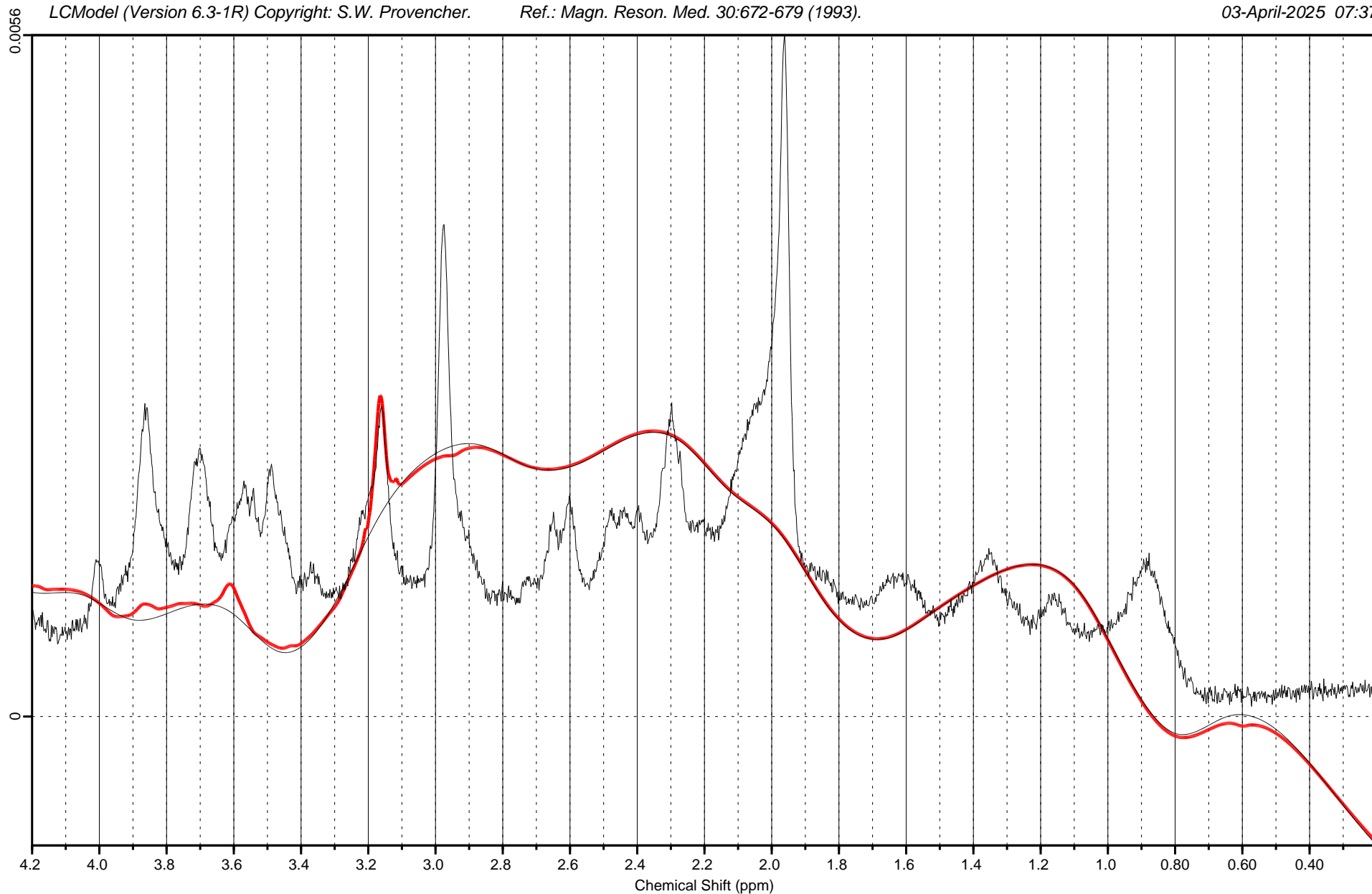
GPC Conc. = 1.98E-05

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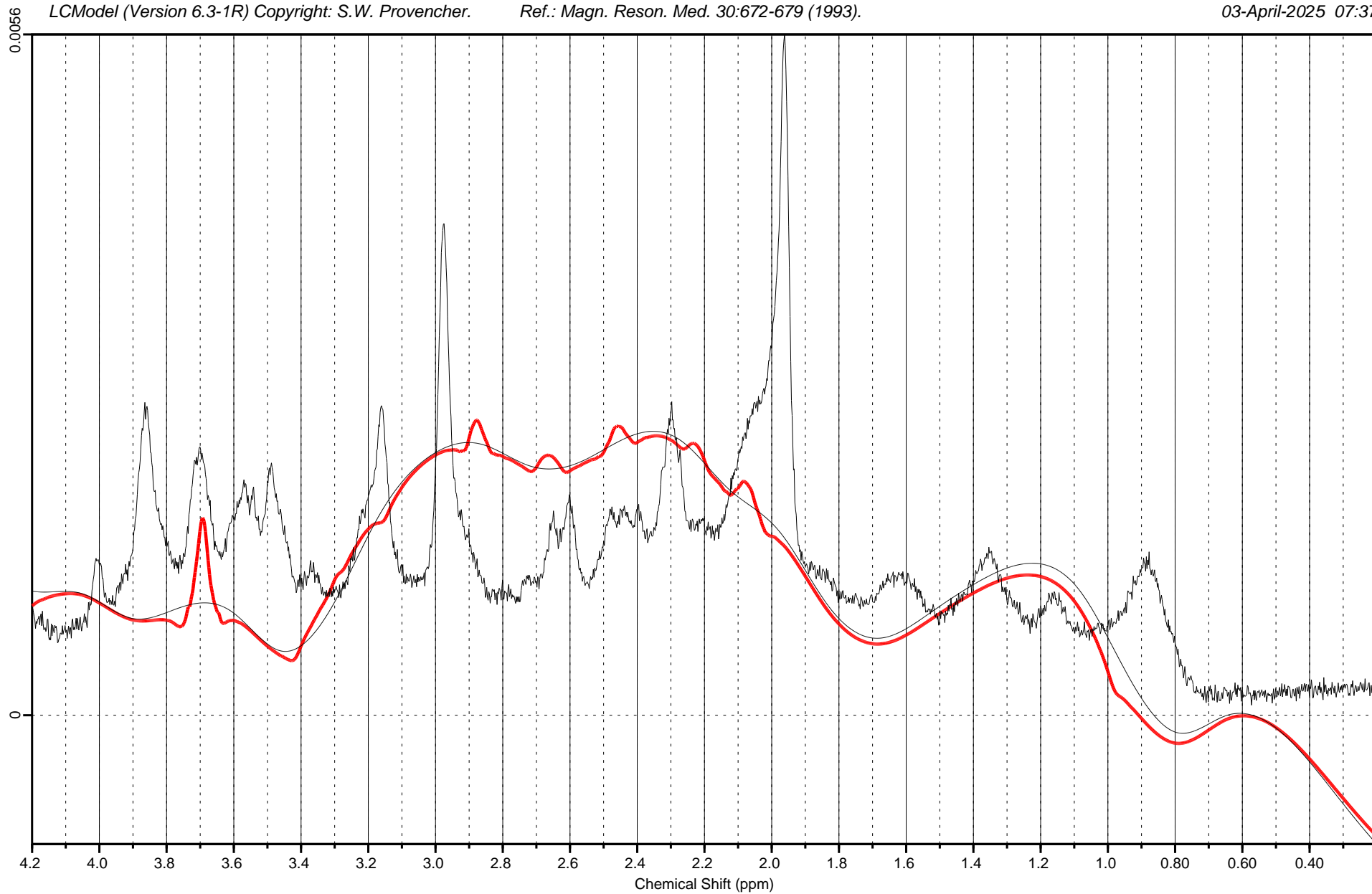
GSH Conc. = 6.03E-05

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Fasting_Sub01_20241007_1150_PCC_Met

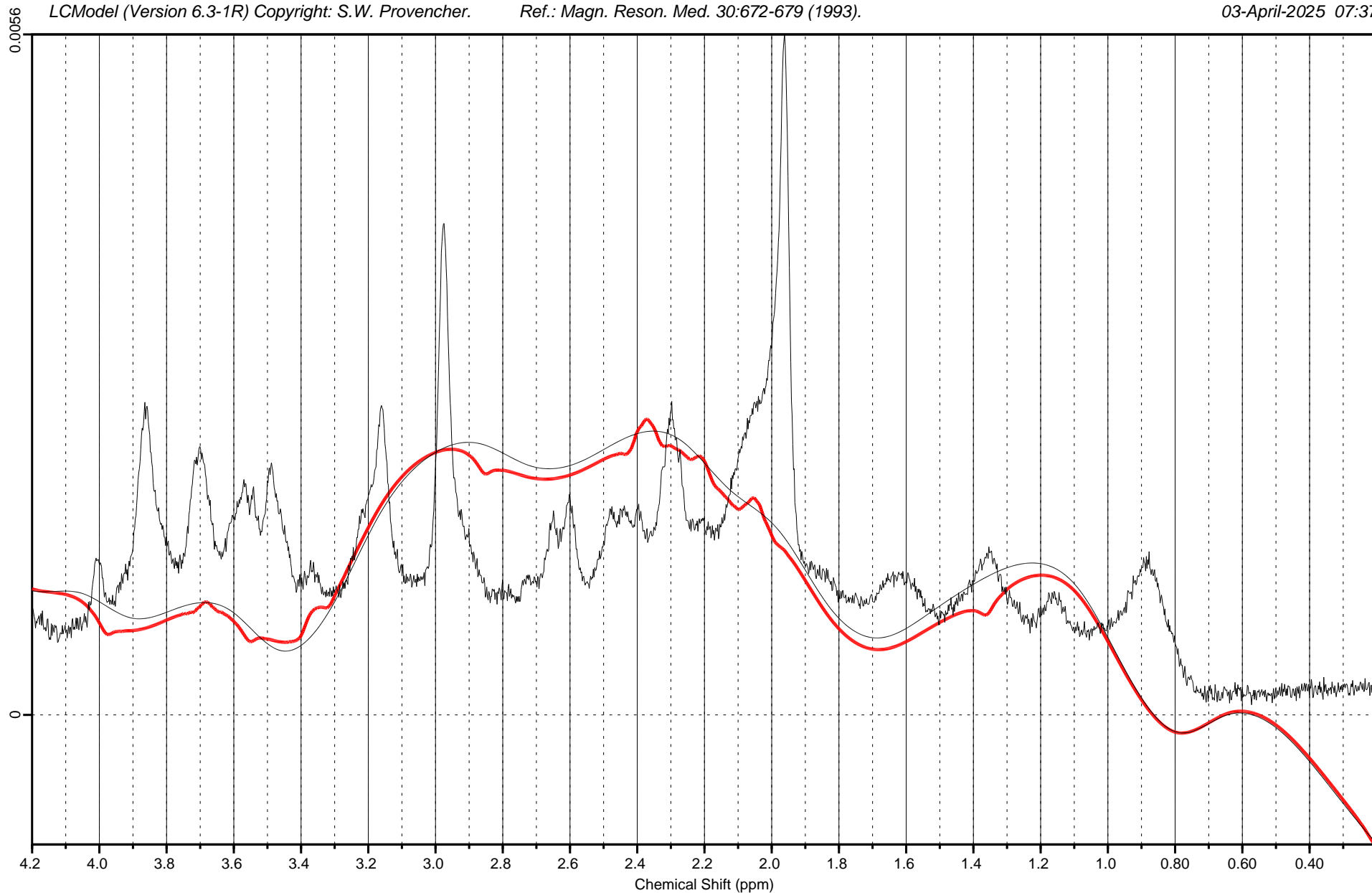
Gln Conc. = 5.53E-05

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Fasting_Sub01_20241007_1150_PCC_Met

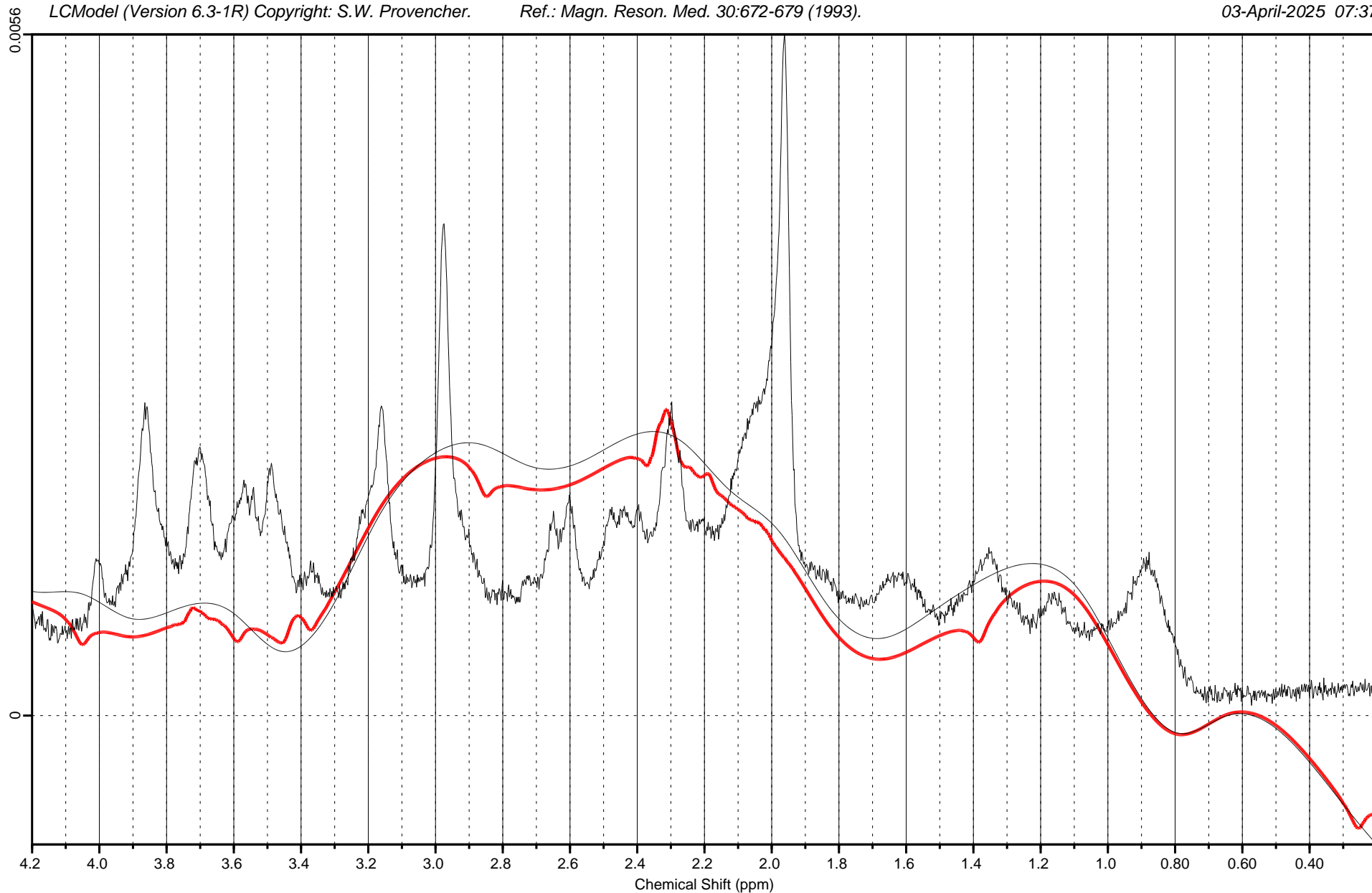
Glu Conc. = 8.61E-05

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Fasting_Sub01_20241007_1150_PCC_Met

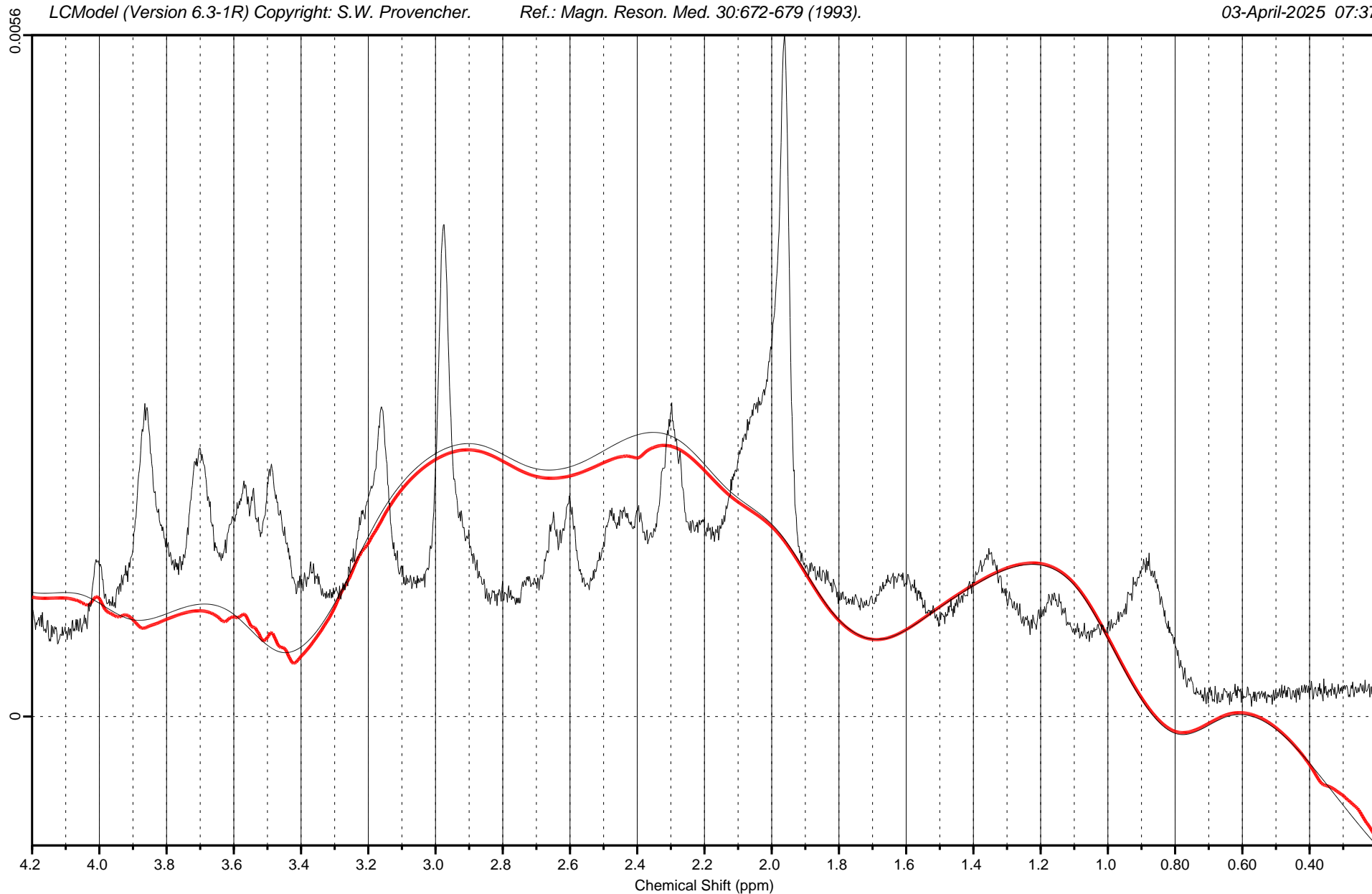
Ins Conc. = 3.27E-05

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Fasting_Sub01_20241007_1150_PCC_Met

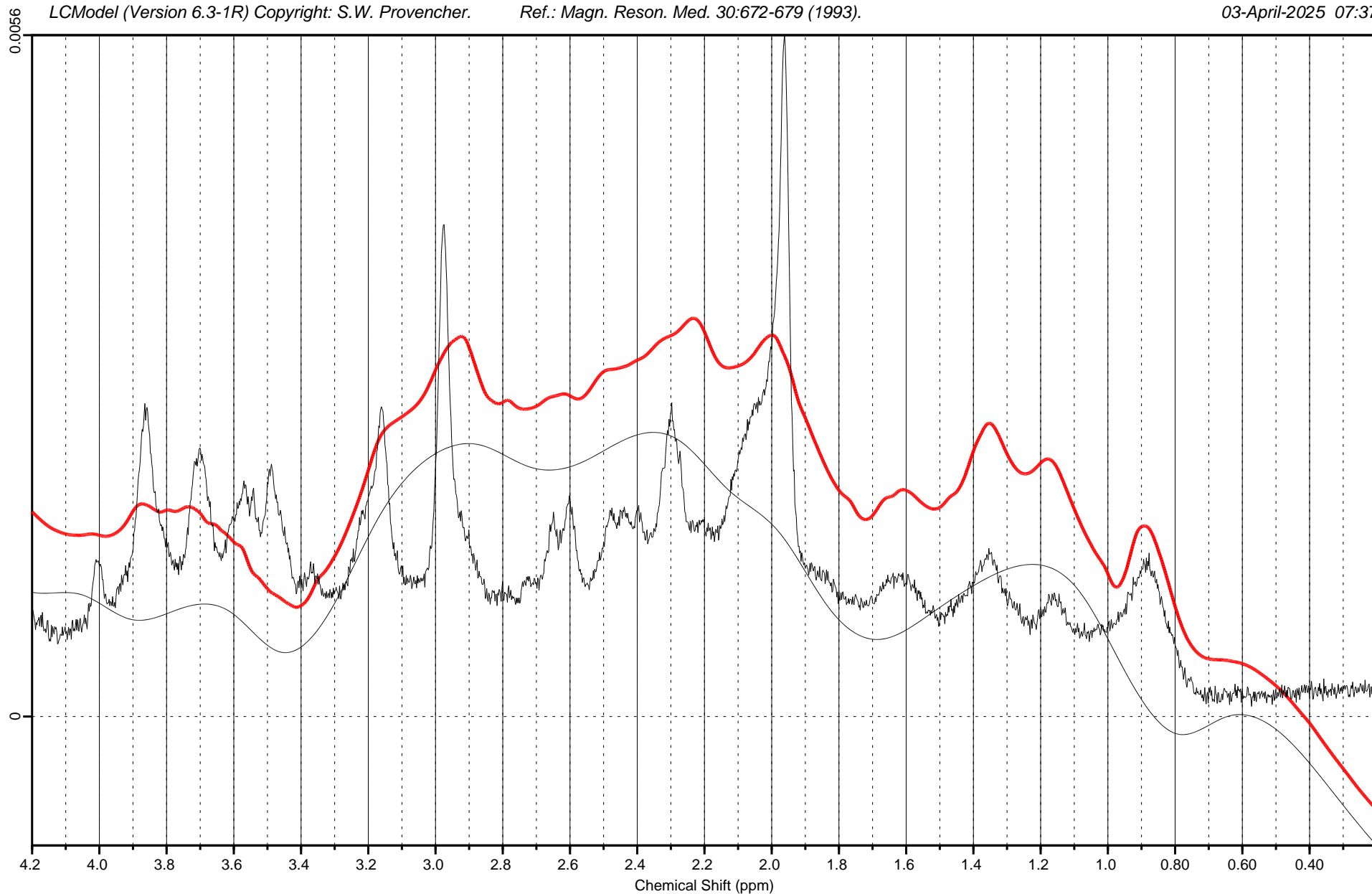
Mac Conc. = 4.09E-05

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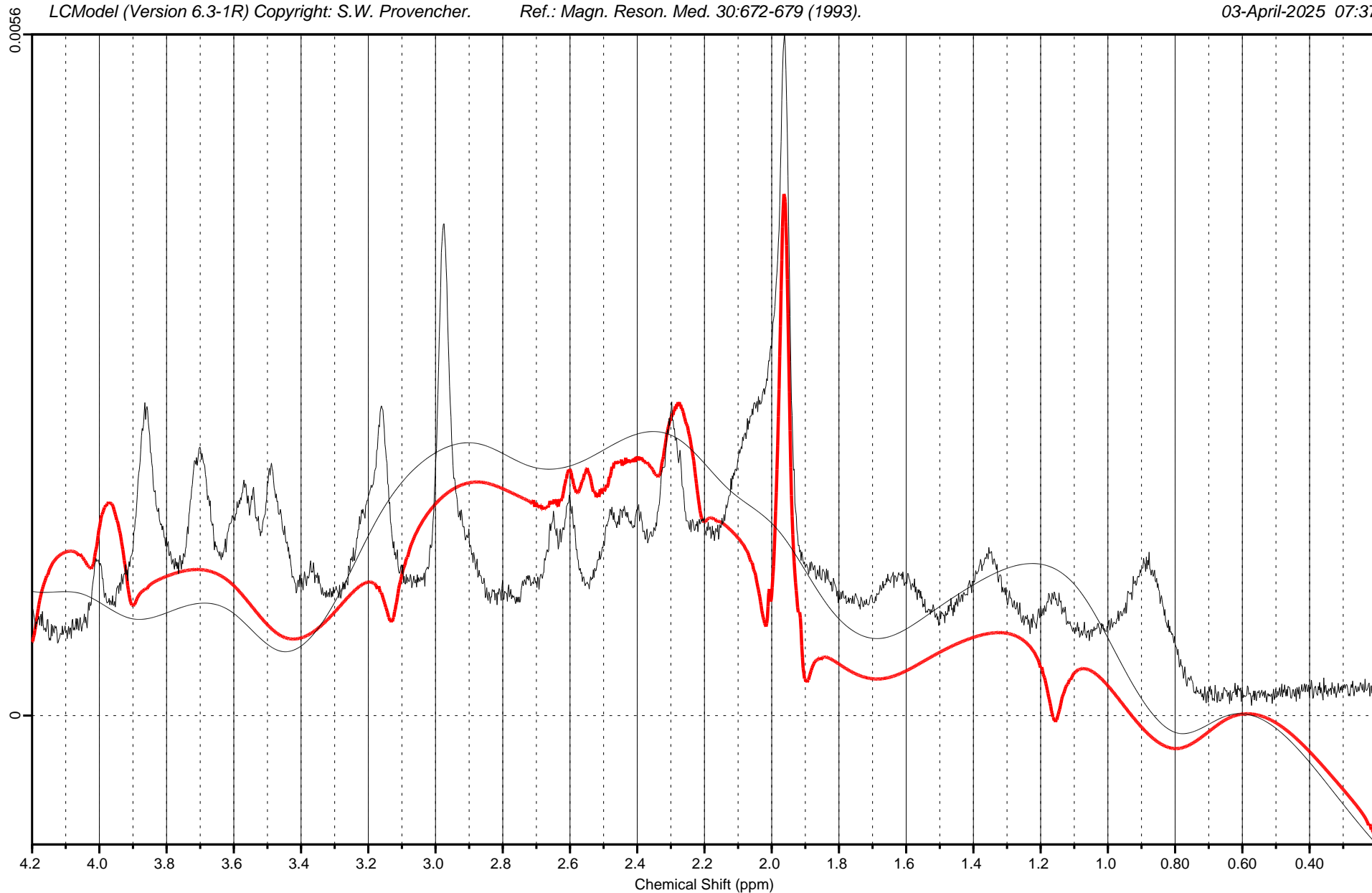
NAA Conc. = 2.16E-04

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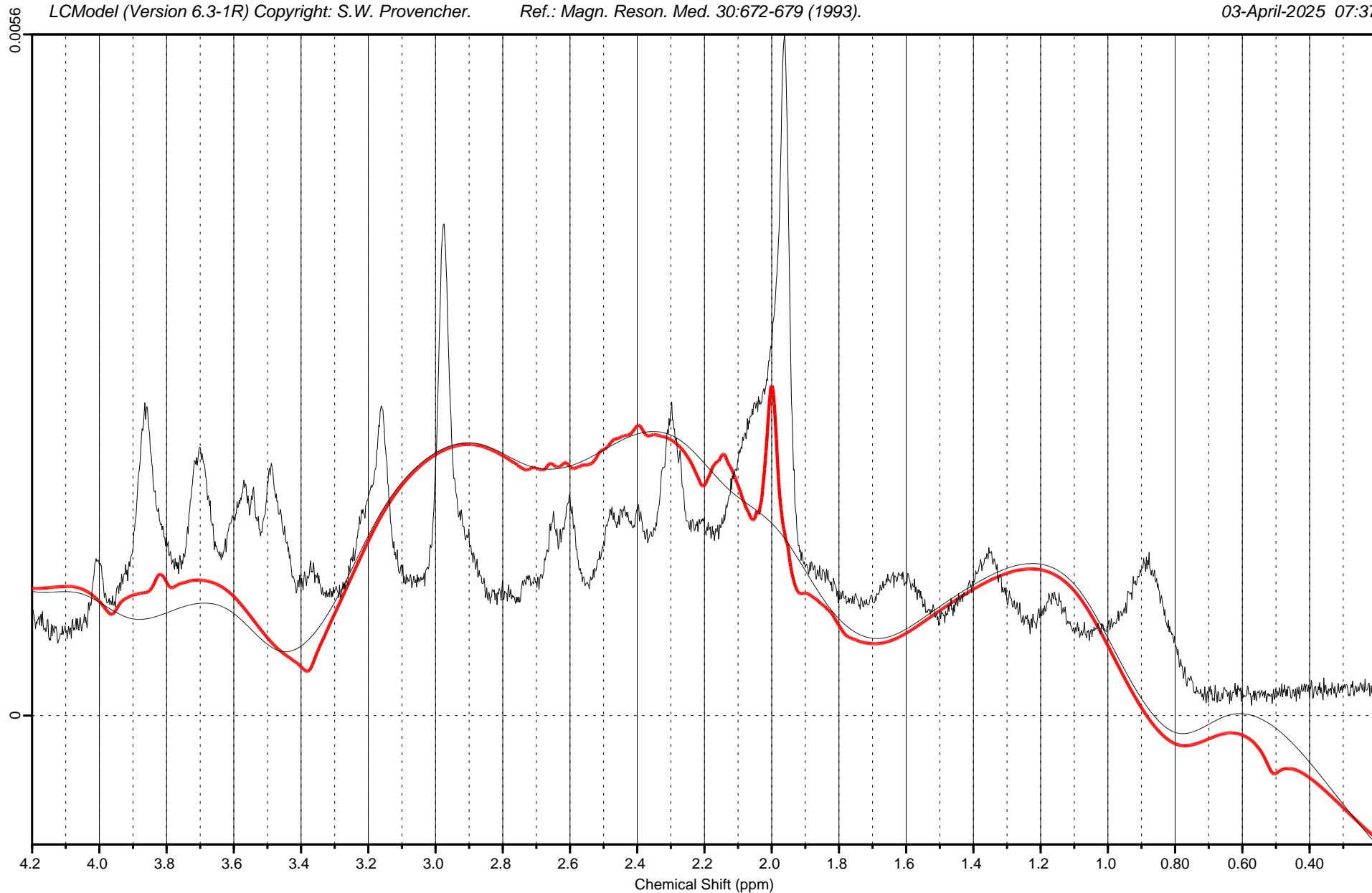
NAAG Conc. = 7.84E-05

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Fasting_Sub01_20241007_1150_PCC_Met

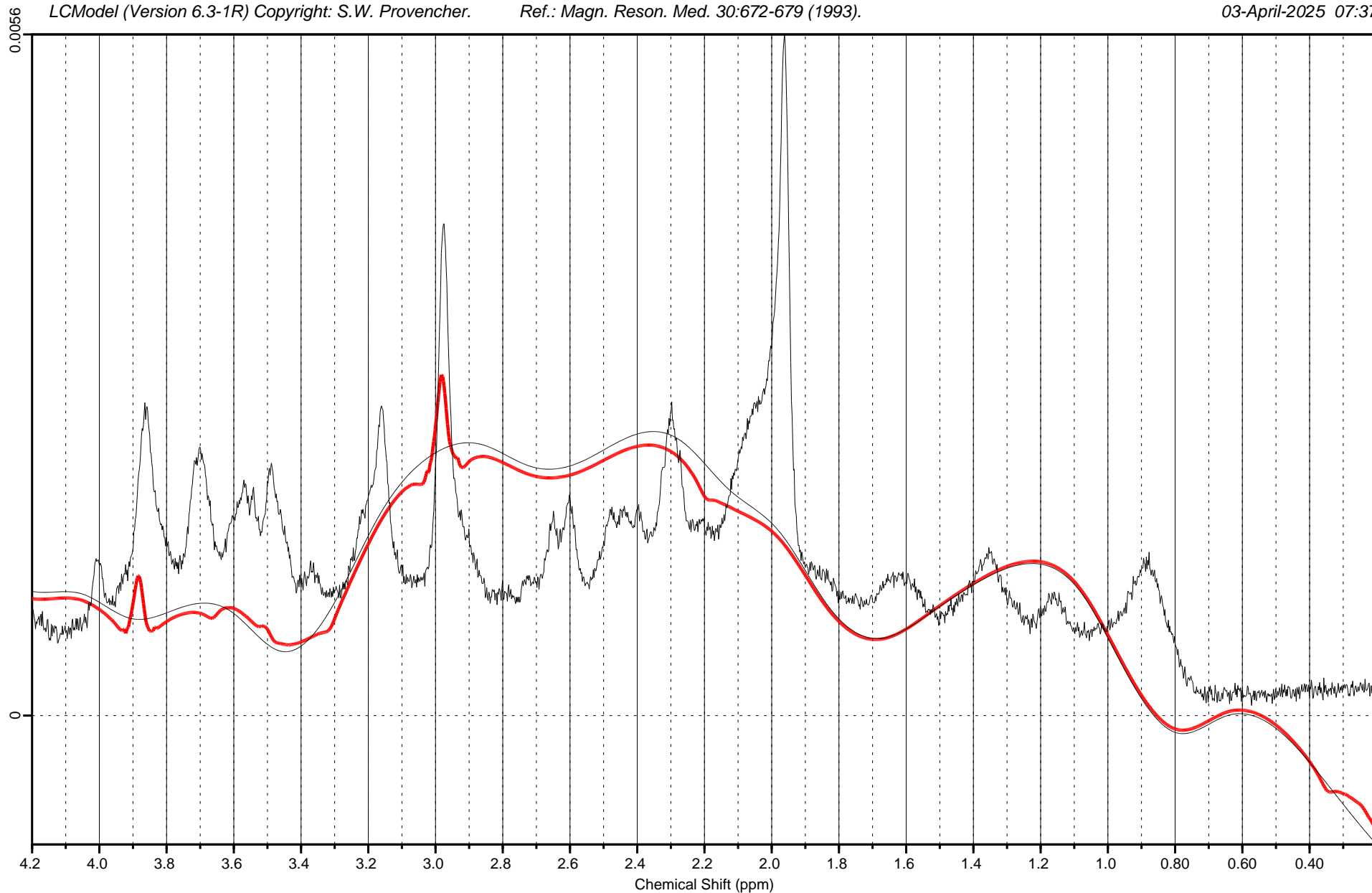
PCr Conc. = 4.89E-05

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Fasting_Sub01_20241007_1150_PCC_Met

Tau Conc. = 4.76E-05

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