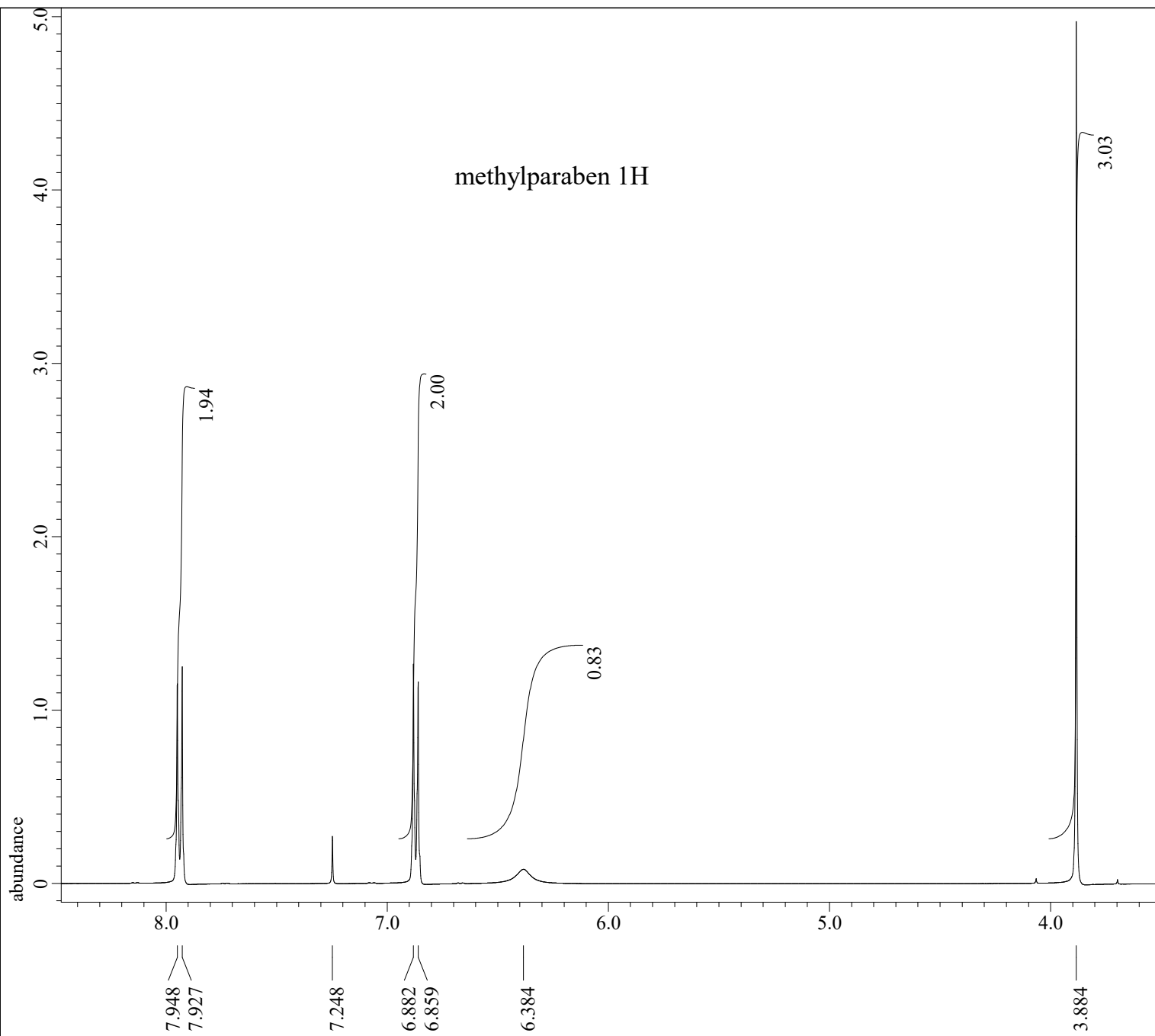




methylparaben 1H



X : parts per Million : Proton

Filename	= MetPar_Beres_Sindler_
Author	= Farsa
Experiment	= proton.jxp
Sample_Id	= MetPar_Beres_Sindler
Solvent	= CHLOROFORM-D
Actual_Start_Time	= 4-OCT-2019 11:59:58
End_Time	= 4-OCT-2019 12:02:03
Revision_Time	= 11-DEC-2019 11:49:03
Comment	= single_pulse
Data_Format	= 1D COMPLEX
Dim_Size	= 13107
Dim_Title	= Proton
Dim_Units	= [ppm]
Dimensions	= X
Spectrometer	= JNM-ECZ400R/S1
Field_Strength	= 9.389766[T] (400[MHz])
X_Acq_Duration	= 2.18628096[s]
X_Domain	= 1H
X_Freq	= 399.78219838[MHz]
X_Offset	= 5[ppm]
X_Points	= 16384
X_Prescans	= 1
X_Resolution	= 0.45739775[Hz]
X_Sweep	= 7.4940048[kHz]
X_Sweep_Clippped	= 5.99520384[kHz]
Irr_Domain	= Proton
Irr_Freq	= 399.78219838[MHz]
Irr_Offset	= 5[ppm]
Tri_Domain	= Proton
Tri_Freq	= 399.78219838[MHz]
Tri_Offset	= 5[ppm]
Blanking	= 2[us]
Clipped	= FALSE
Scans	= 16
Total_Scans	= 16
Relaxation_Delay	= 5[s]
Recvr_Gain	= 56
Temp_Get	= 21.3[dC]
X_90_Width	= 5.8[us]
X_Acq_Time	= 2.18628096[s]
X_Angle	= 45[deg]
X_Atn	= 4.7[dB]
X_Pulse	= 2.9[us]
Irr_Mode	= Off
Tri_Mode	= Off
Comment_1	= *** Pulse ***
Comment_111	= *** presat_time ***
Comment_201	= *** obs_dante_presatu
Comment_202	= *** irr_preaturation
Comment_203	= *** tri_preaturation
Comment_7	= *** Pulse Delay ***
Dante_Loop	= 500
Dante_Presat	= FALSE
Decimation_Rate	= 0
Get_90	= pulse_service::get_90
Get_Atn	= pulse_service::get_at
Get_Freq	= pulse_service::get_fr
Get_Gamma	= pulse_service::get_ga