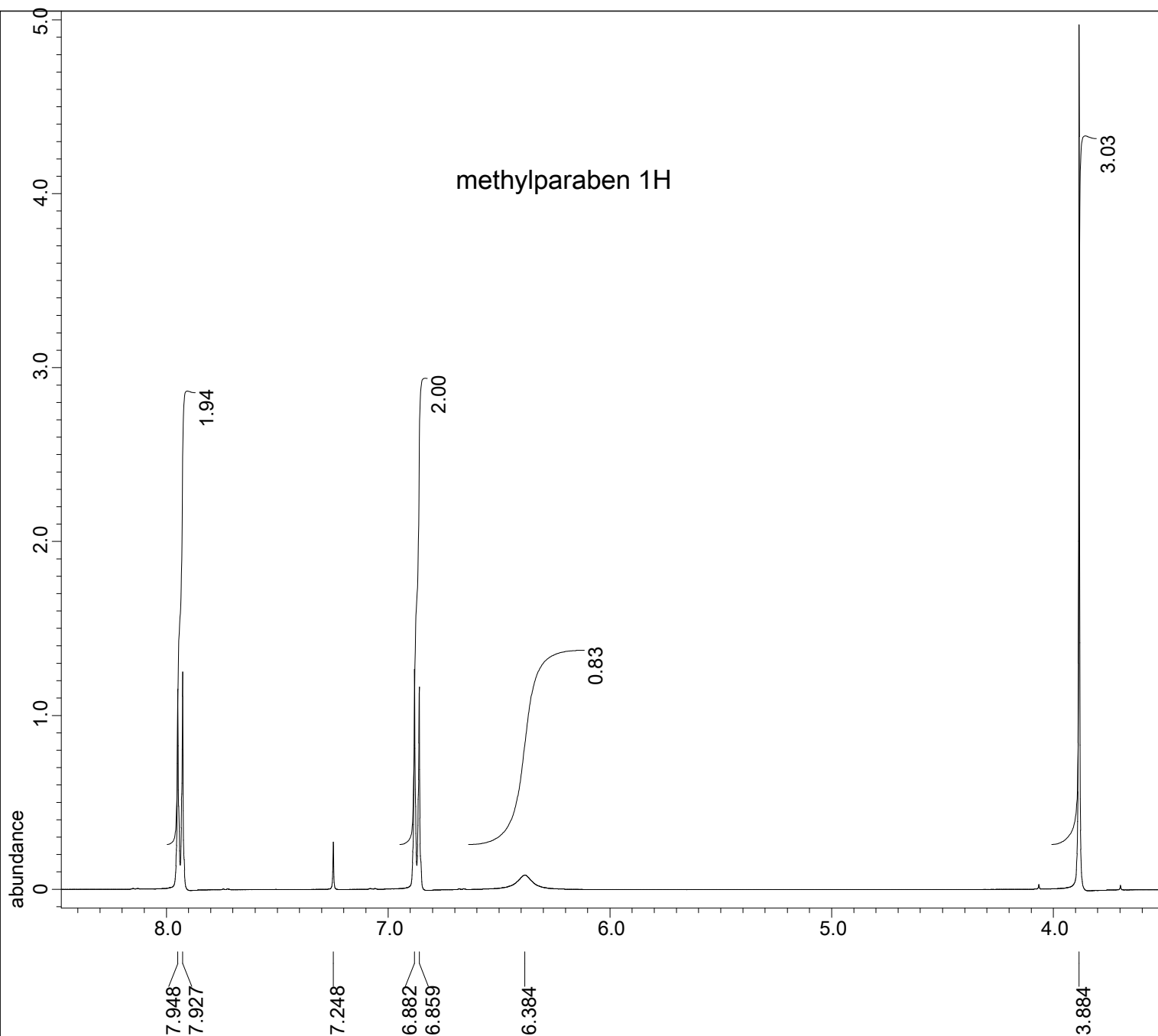




methylparaben 1H



X : parts per Million : Proton

Author = Farsa
Experiment = proton.jxp

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