

Figure S6: HPLC-MS analyses of the fraction C-3071-I-A06 from the *C. chinensis* extract enriched in astragalin. (A) Total ion chromatogram (TIC) in positive ion mode; (B) TIC in negative ion mode; (C) Chromatogram of the evaporative light scattering detector (ELSD); (D) Chromatogram of the photodiode array detector (PDA); (E-F) Mass spectra of the identified compound in positive and negative ion mode.

Current Data Parameters
NAME C3071IS.005
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters

Date_ 20191101
Time 11.06
INSTRUM spect
PROBHD 5 mm QNP 1H/1
PULPROG zg30
TD 32768
SOLVENT MeOH
NS 32
DS 2
SWH 8278.146 Hz
FIDRES 0.252629 Hz
AQ 1.9791873 sec
RG 128
DW 60.400 usec
DE 6.00 usec
TE 300.0 K
D1 1.00000000 sec

===== CHANNEL f1 =====

NUC1 1H
P1 11.60 usec
PL1 5.00 dB
SFO1 400.1324710 MHz

F2 - Processing parameters

SI 16384
SF 400.1300913 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00

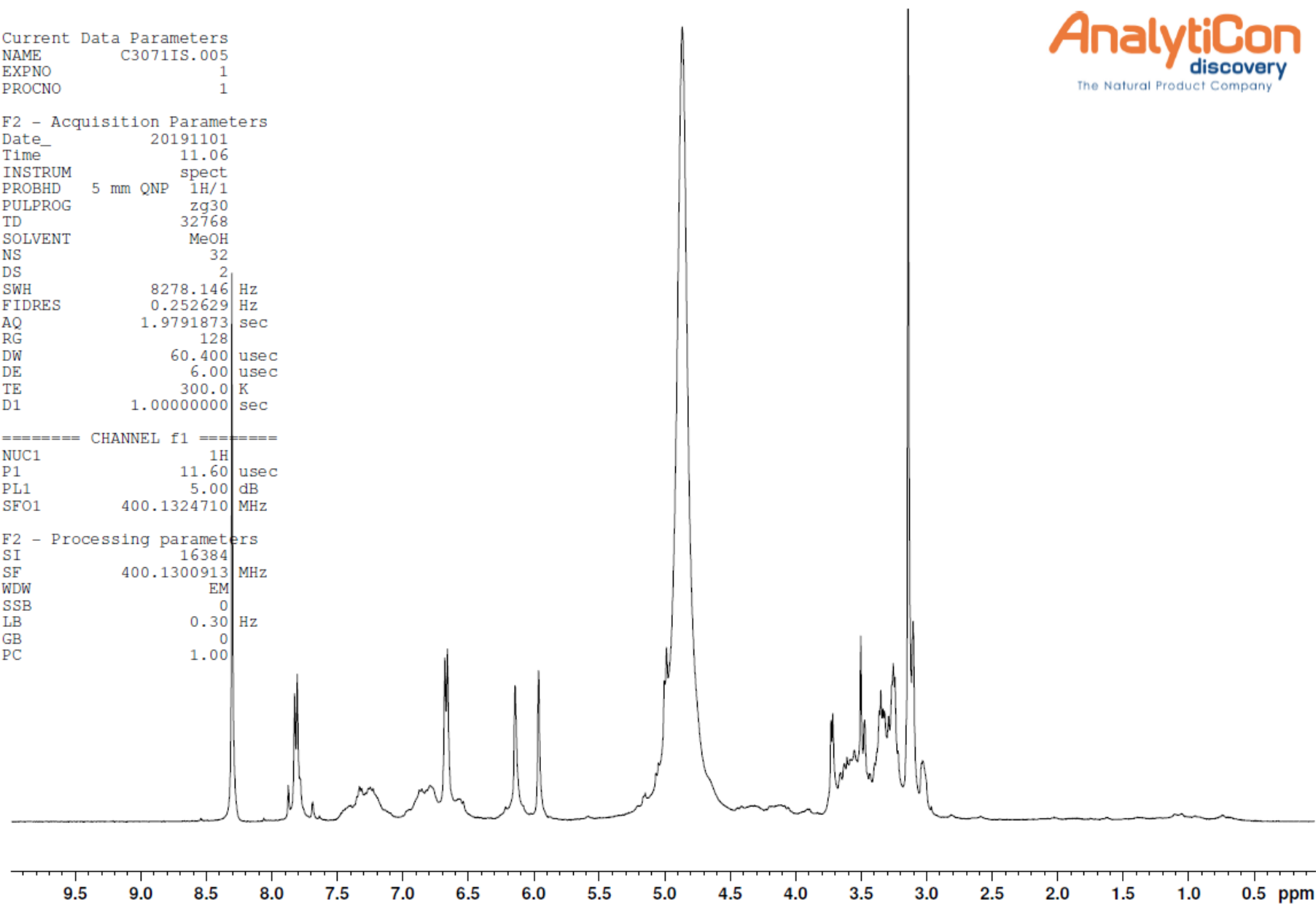


Figure S7: NMR spectrum of astragalin.

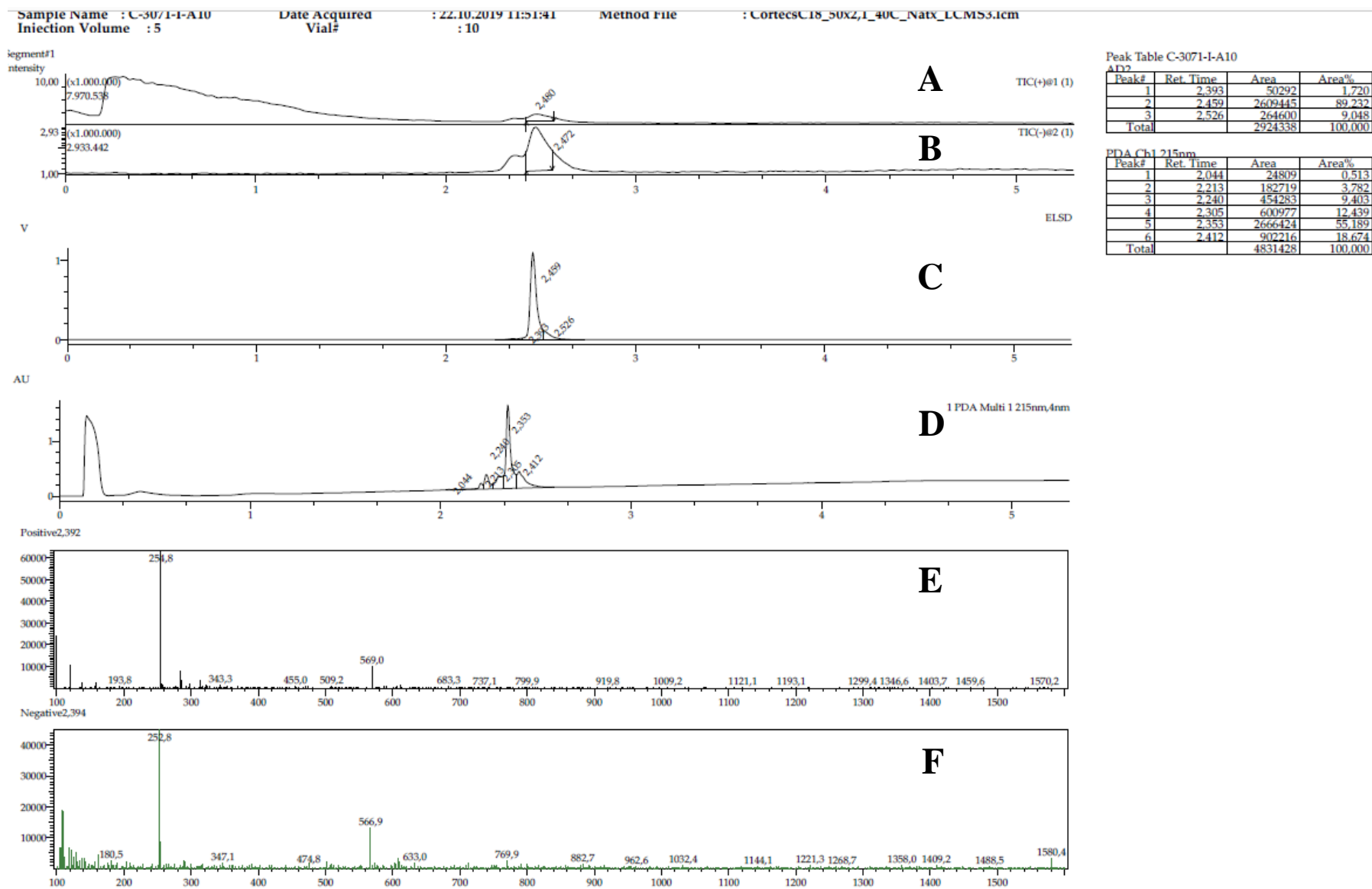


Figure S8: HPLC-MS analyses of the fraction C-3071-I-A10 from the *C. chinensis* extract enriched in pinoresinol. (A) Total ion chromatogram (TIC) in positive ion mode; (B) TIC in negative ion mode; (C) Chromatogram of the evaporative light scattering detector (ELSD); (D) Chromatogram of the photodiode array detector (PDA); (E-F) Mass spectra of the identified compound in positive and negative ion mode.

Current Data Parameters
NAME C3071IS.008
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters

Date_ 20191101
Time 11.34
INSTRUM spect
PROBHD 5 mm QNP 1H/1
PULPROG zg30
TD 32768
SOLVENT MeOH
NS 32
DS 2
SWH 8278.146 Hz
FIDRES 0.252629 Hz
AQ 1.9791873 sec
RG 228.1
DW 60.400 usec
DE 6.00 usec
TE 300.0 K
D1 1.00000000 sec

===== CHANNEL f1 =====

NUC1 1H
P1 11.60 usec
PL1 5.00 dB
SFO1 400.1324710 MHz

F2 - Processing parameters

SI 16384
SF 400.1300913 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00

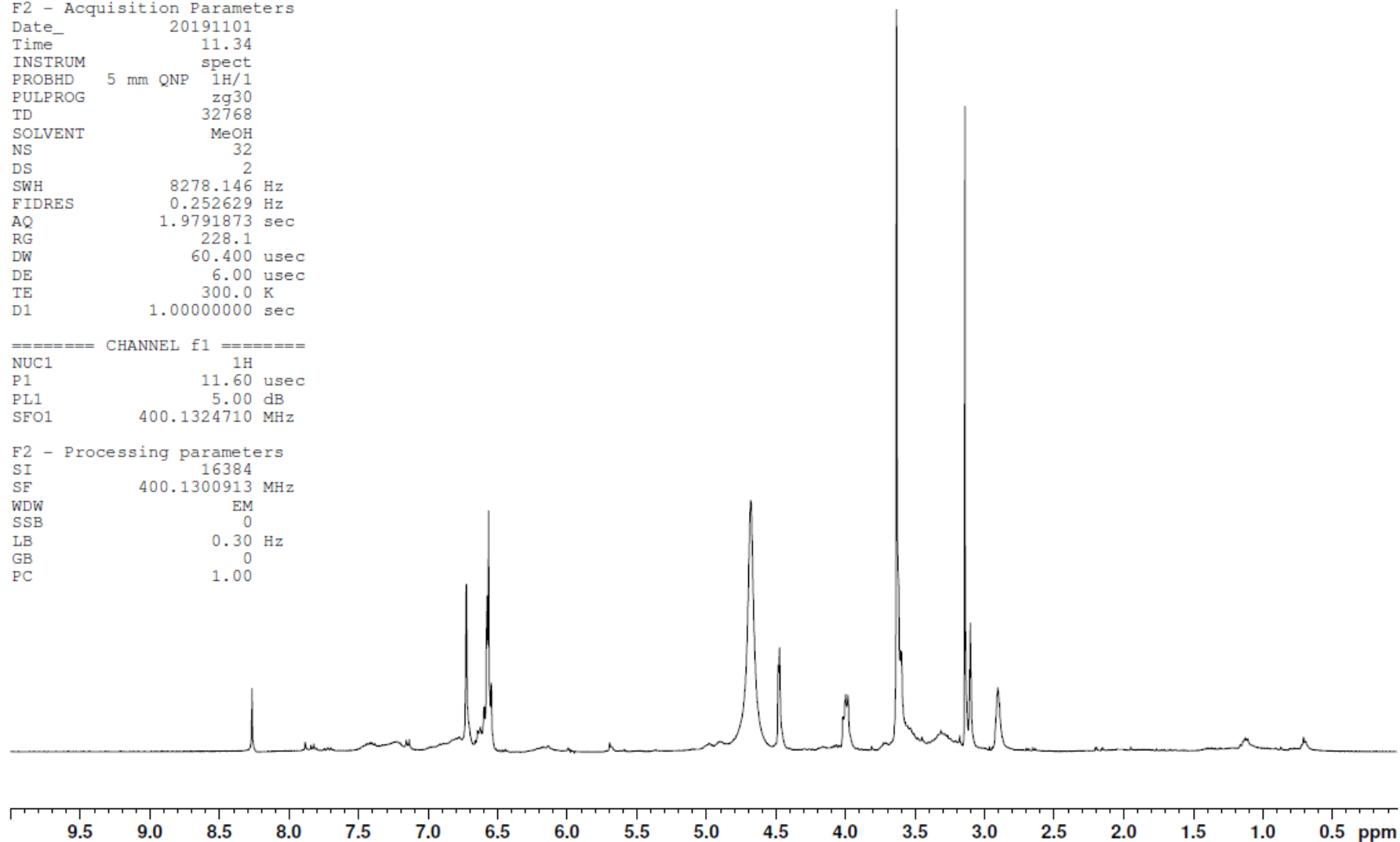


Figure S9: NMR spectrum of pinoresinol.

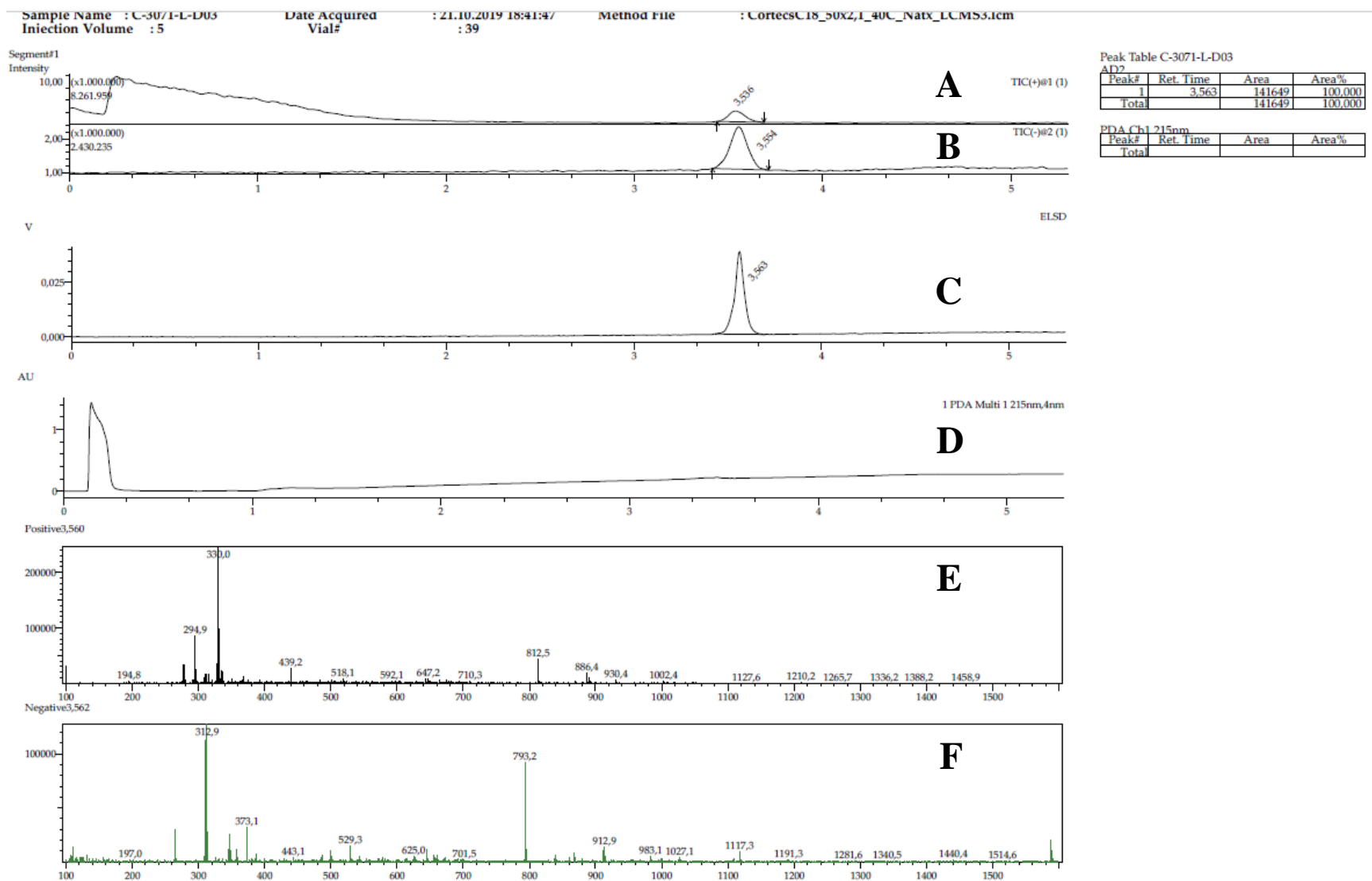


Figure S10: HPLC-MS analyses of the fraction C-3071-L-D03 from the *C. chinensis* extract enriched in zingibroside R1. (A) Total ion chromatogram (TIC) in positive ion mode; (B) TIC in negative ion mode; (C) Chromatogram of the evaporative light scattering detector (ELSD); (D) Chromatogram of the photodiode array detector (PDA); (E-F) Mass spectra of the identified compound in positive and negative ion mode.

Current Data Parameters
NAME C3071LS.013
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20191029
Time 11.16
INSTRUM spect
PROBHD 5 mm SEI 1H-13
PULPROG zg30
TD 32768
SOLVENT MeOD
NS 32
DS 2
SWH 10000.000 Hz
FIDRES 0.305176 Hz
AQ 1.6384000 sec
RG 181
DW 50.000 usec
DE 6.00 usec
TE 300.0 K
D1 2.00000000 sec

===== CHANNEL f1 =====
NUC1 1H
P1 10.50 usec
PL1 5.00 dB
SFO1 500.1240010 MHz

F2 - Processing parameters
SI 32768
SF 500.1200000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00

AnalytiCon
discovery
The Natural Product Company

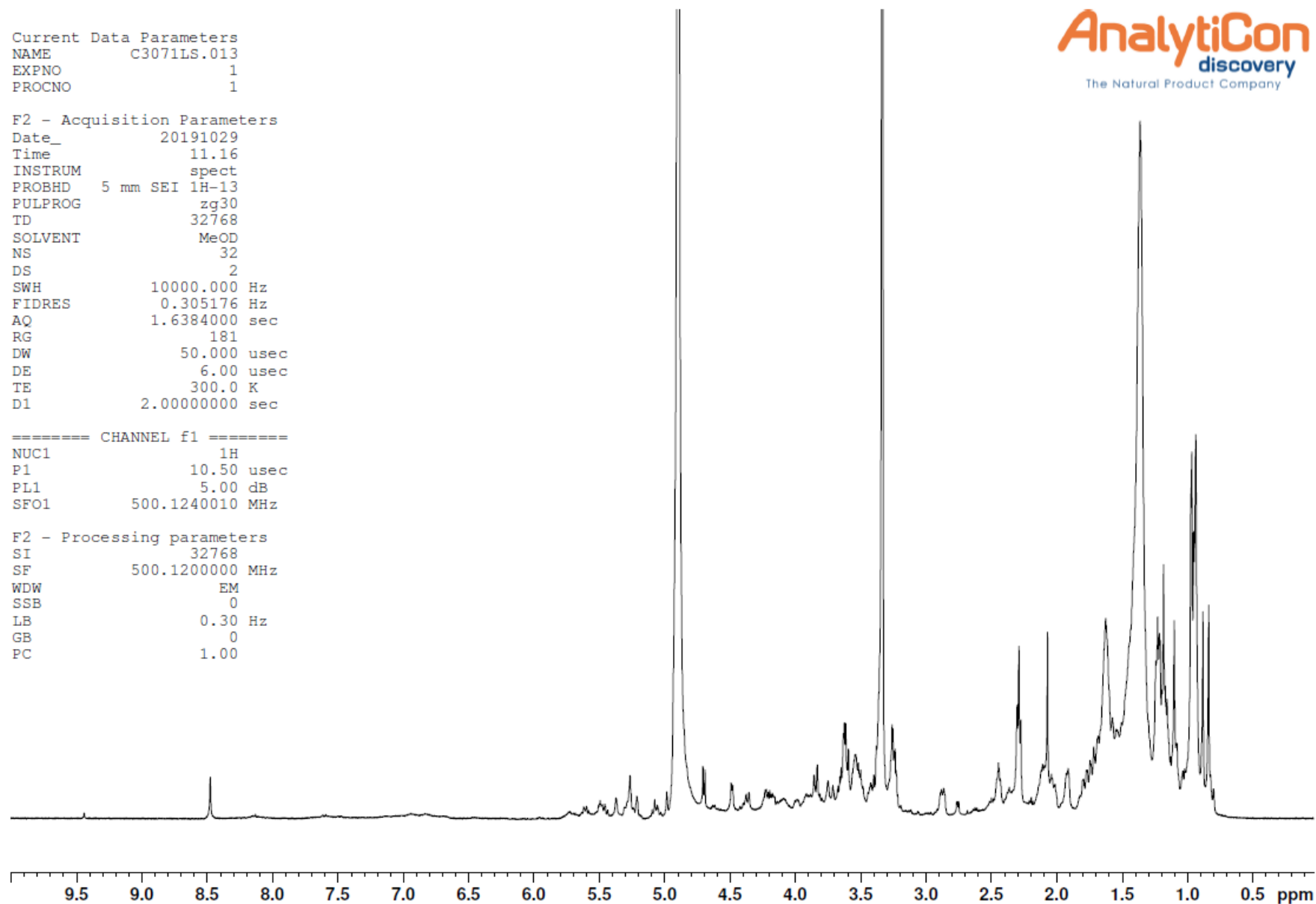


Figure S11: NMR spectrum of zingibroside R1.