

Discovery of novel coumarin-Schiff base hybrids as potential acetylcholinesterase inhibitors: Design, synthesis, enzyme inhibition, and computational studies

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EXPERIMENTAL

In vitro inhibition study on AChE

The assay for AChE inhibitory activity was done based on a modified protocol [1-5]. Acetylcholinesterase (Type VI-S, from *Electrophorus electricus*), 5,5'-dithiobis-2-nitrobenzoic acid (Ellman's reagent, DTNB) and acetylthiocholine chloride (ATCI) were purchased from Sigma Aldrich. Enzyme solutions were prepared to give 0.28 units/mL aliquots. Stock solutions were prepared by dissolving 1 mg of tested compounds in 1 mL of DMSO and diluted to a final concentration of 1000, 800, 600, 400, 200, 100, and 50 µg/mL. The assay buffer solution A (pH = 8.0) was prepared by taking 93.2 mL of solution (6 g of sodium dihydrogen phosphate was dissolved in 500 mL of deionized water) and mixed with 6.8 mL of solution (7 g of sodium hydrogen phosphate was dissolved in 500 mL of deionized water) and adjusted. 57.7 mL of sodium dihydrogen phosphate and 42.3 mL of sodium hydrogen phosphate were mixed and adjusted to prepare the buffer B (pH = 7.0). Furthermore, fresh solutions of 0.01 M DTNB (0.396 g of DTNB and 15 mg NaHCO₃ were dissolved in 100 mL of the buffer B) in a dark place, and 0.075 M ATCI (0.2048 g of ATCI was dissolved in 10 mL of deionized water), were prepared. In 96-well plates, 20 µL of the test compounds, 10 µL of DTNB, 15 µL of the enzyme (AChE) and 140 µL of the buffer solution A, were added in dark conditions and incubated for 15 min, followed by the addition of ATCI (10 µL) and incubated again for the same period. The activity was measured by reading the absorbance of the solution at 412 nm. Blanks containing all the components except the enzyme were carried out. IC₅₀ values were calculated as the concentration of a compound that produces 50% enzyme activity inhibition using the GraphPad Prism 8 programme package. Results are expressed as the mean ± SD of at least three different experiments performed in triplicate.

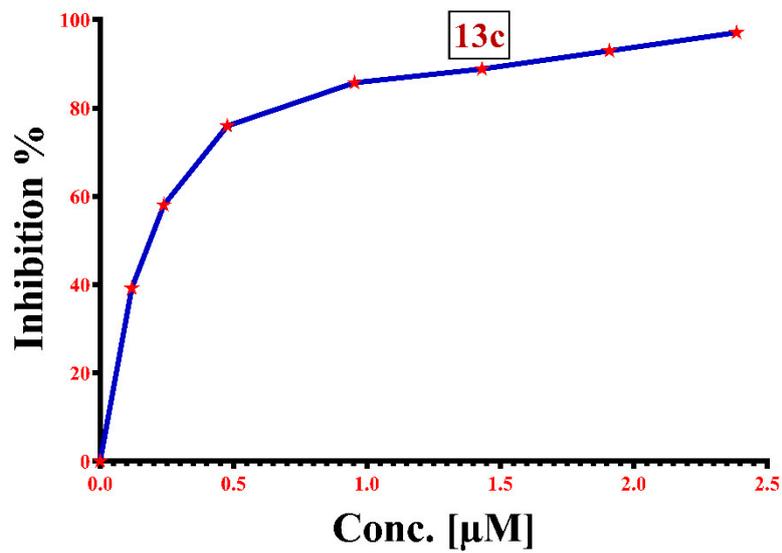
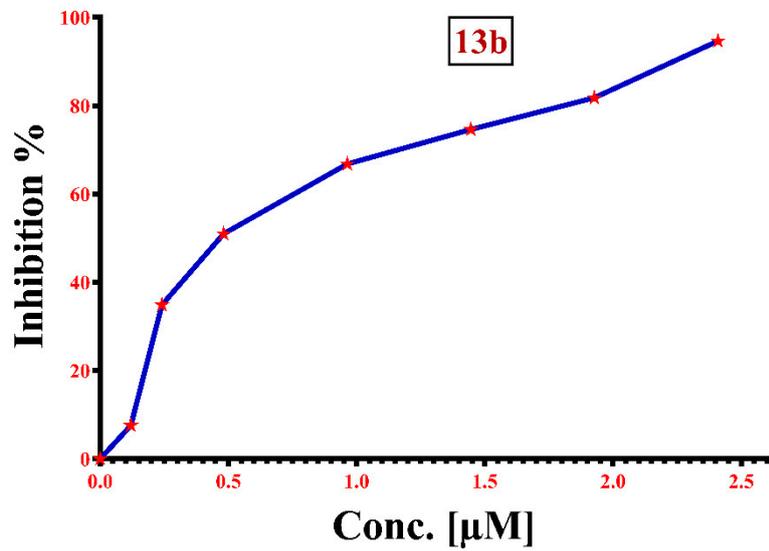
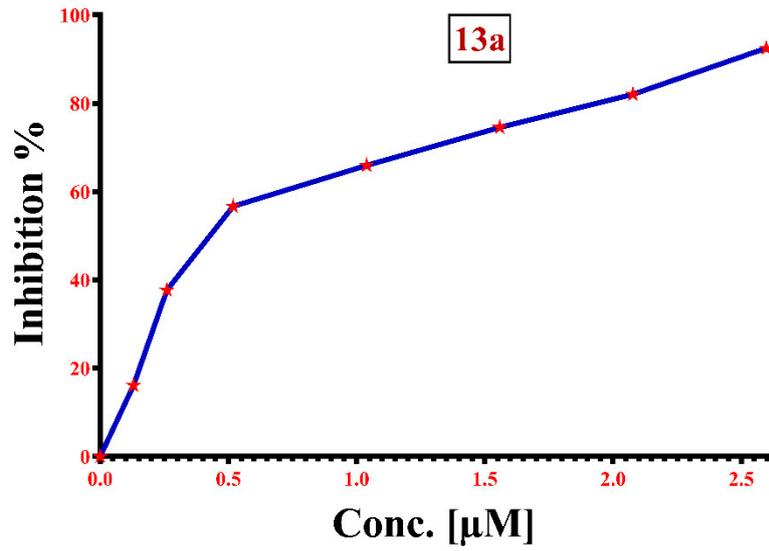
IC₅₀ Calculations

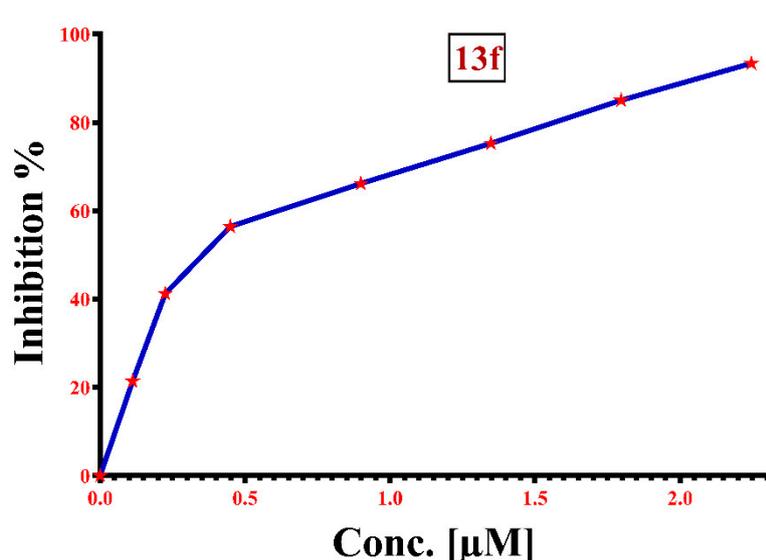
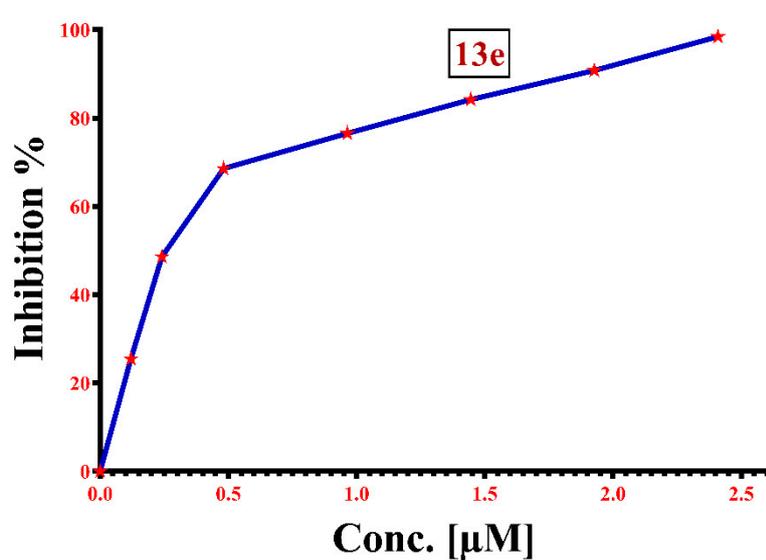
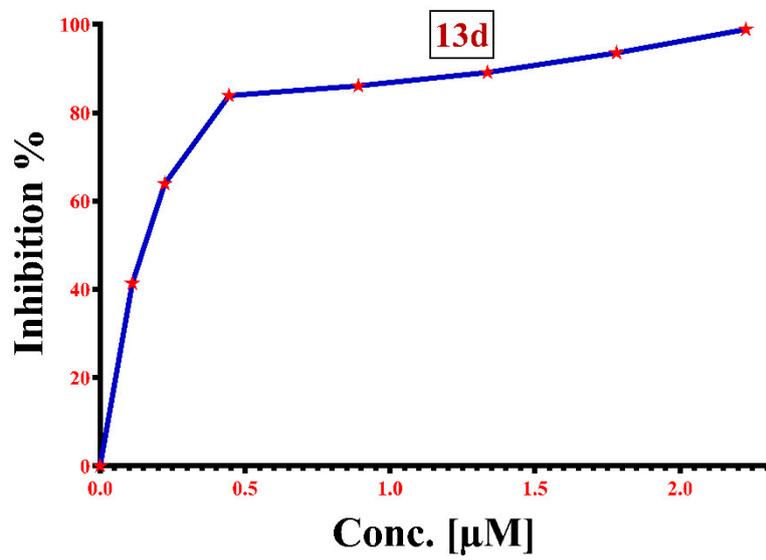
Percentage inhibition was calculated by this formula:

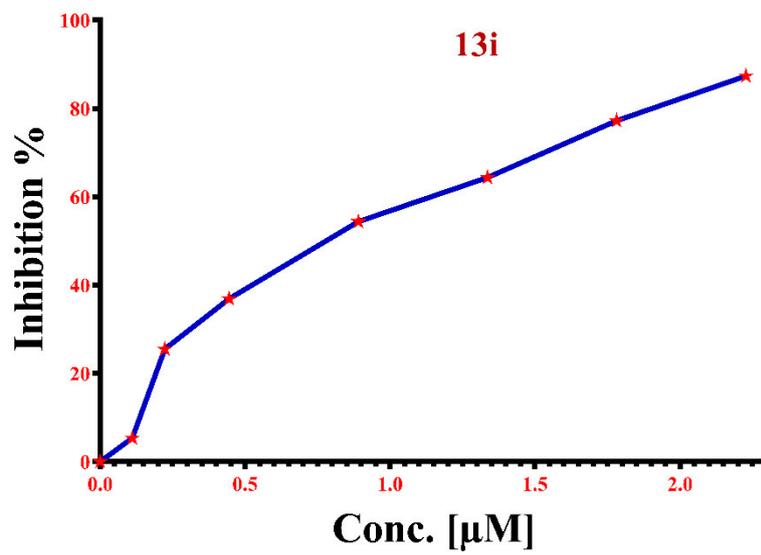
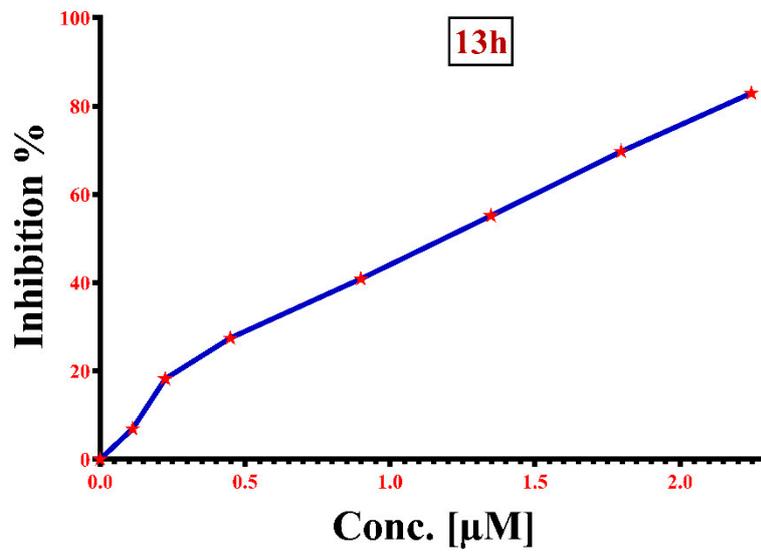
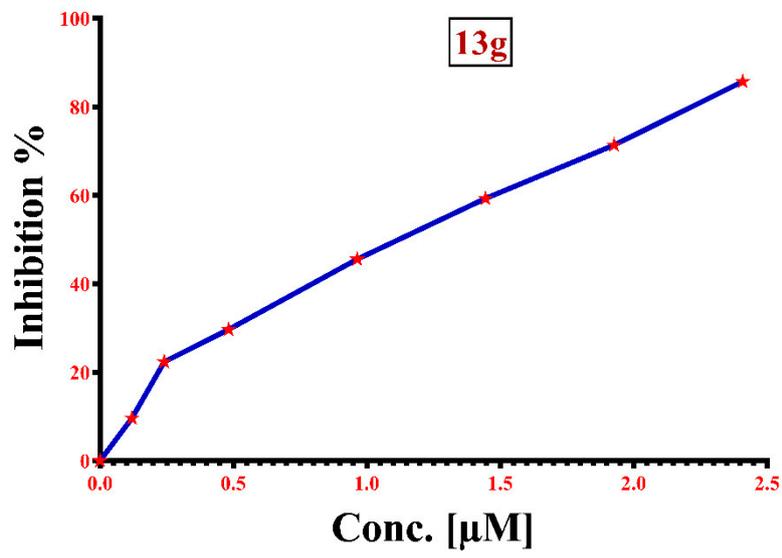
$$\% \text{ (Inhibition)} = \frac{B - A}{B} \times 100$$

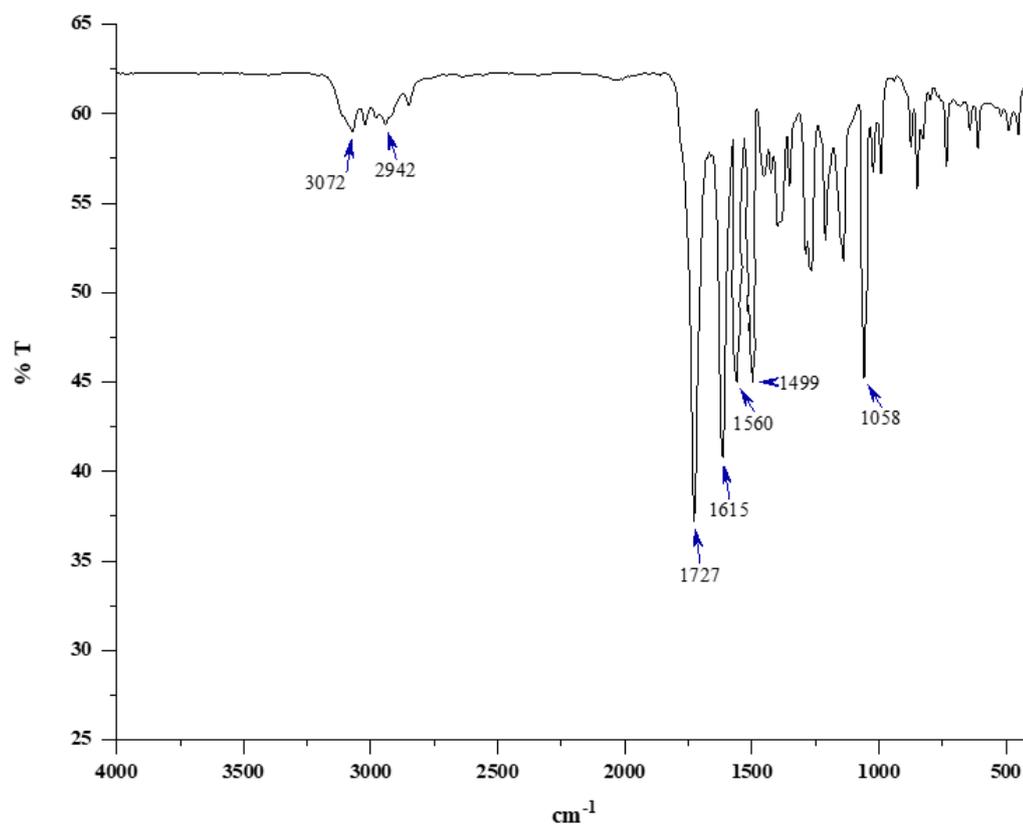
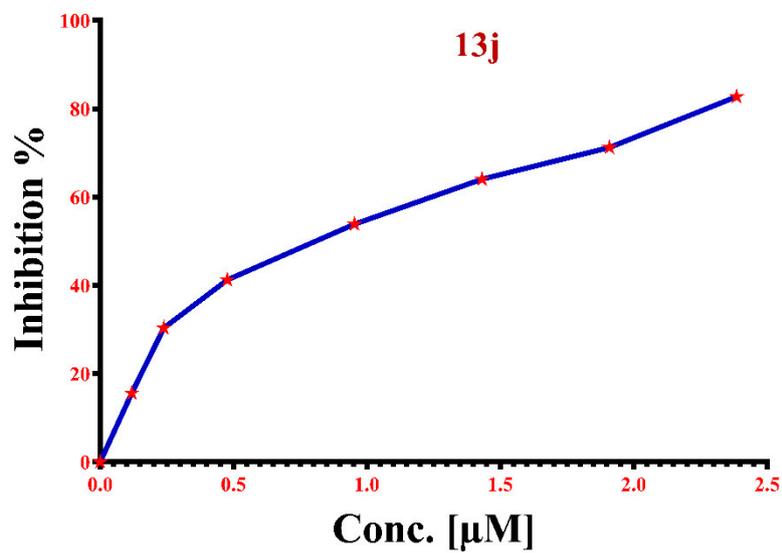
- A= absorbance of the enzyme with the test sample;
- B = absorbance of the enzyme without test sample.
- Each experiment was repeated thrice.

- Concentrations were converted from $\mu\text{g/mL}$ to μM .
- IC_{50} value was calculated by GraphPad Prism.
- Galantamine was used as a reference compound.

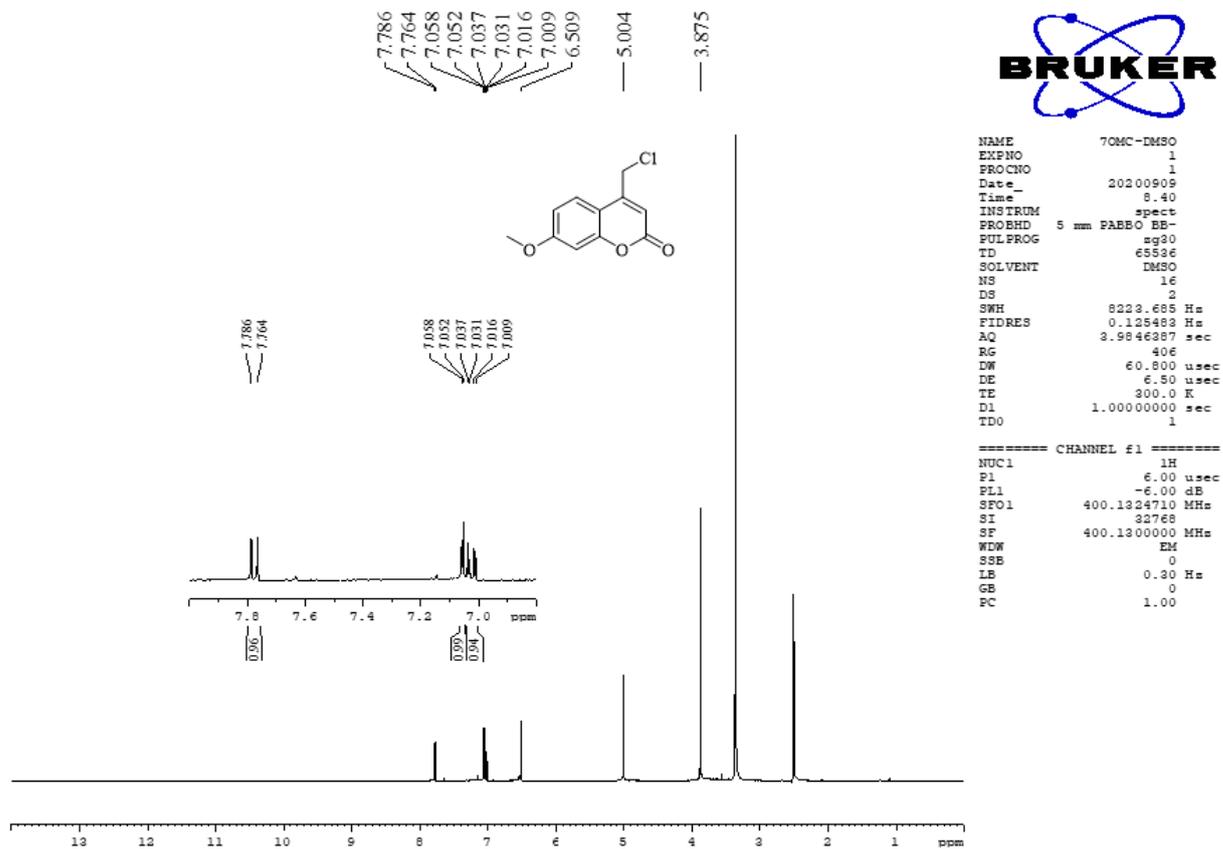




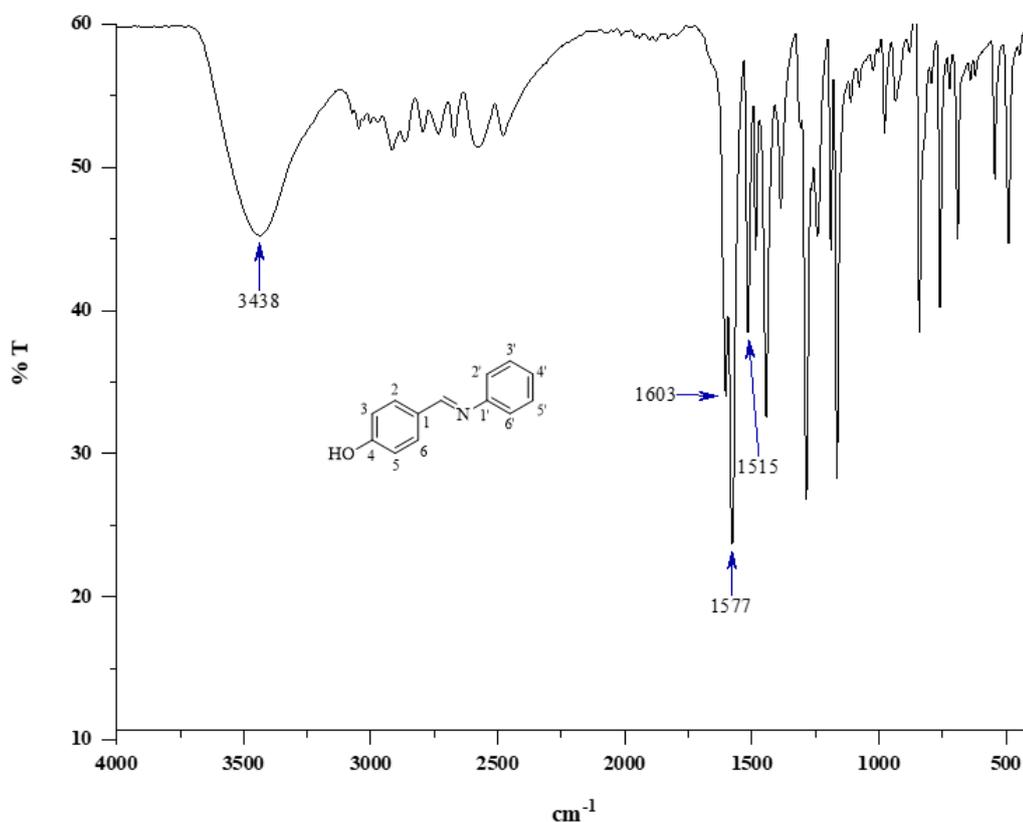




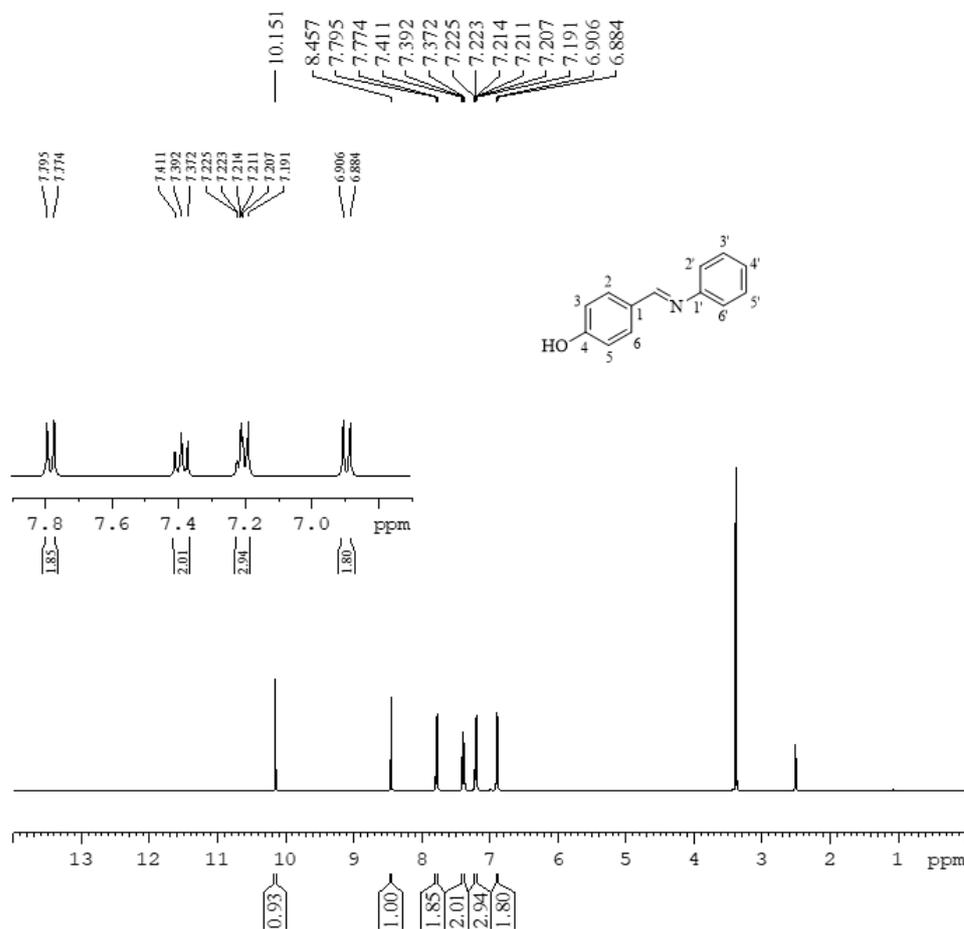
IR spectrum of 4-(chloromethyl)-7-methoxy-2H-chromen-2-one (9)



¹H NMR spectrum of 4-(chloromethyl)-7-methoxy-2H-chromen-2-one (9)



IR spectrum of (*E*)-4-((phenylimino)methyl)phenol (12a)

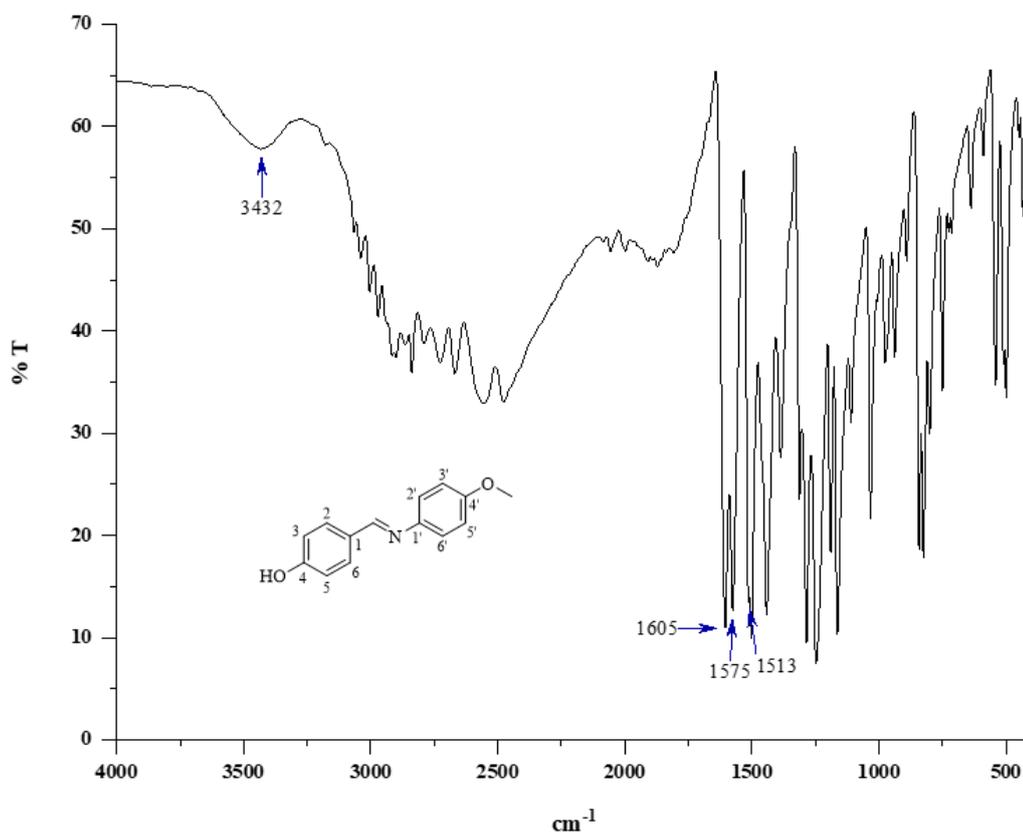


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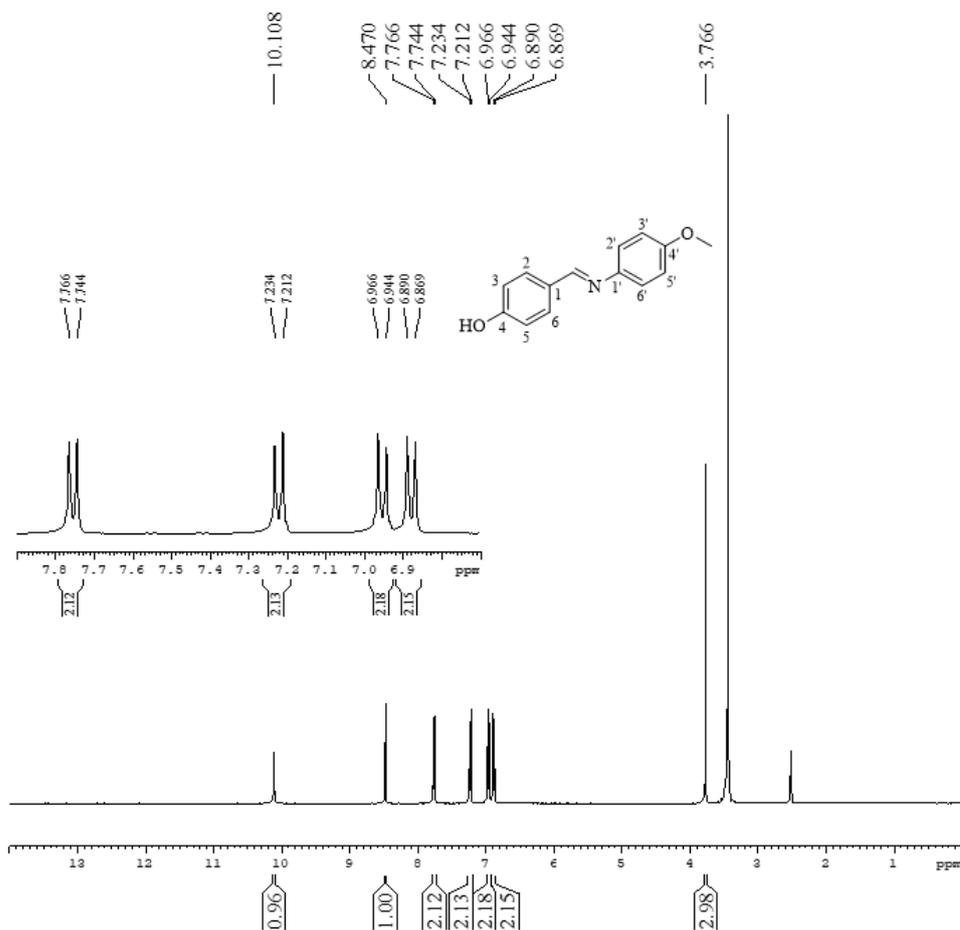
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PULPROG       zg30
TD            65536
SOLVENT       DMSO
NS            16
DS            2
SWH           8223.685 Hz
FIDRES        0.125483 Hz
AQ            3.9846387 sec
RG            203
DW            60.800 usec
DE            6.50 usec
TE            300.0 K
D1            1.00000000 sec
TD0           1

===== CHANNEL f1 =====
NUC1          1H
P1            6.00 usec
PL1           -6.00 dB
SFO1          400.1324710 MHz
SI            32768
SF            400.1300000 MHz
WDW           EM
SSB           0
LB            0.30 Hz
GB            0
PC            1.00
  
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¹H NMR spectrum of 4-((phenylimino)methyl)phenol (12a)



IR spectrum of (E)-4-(((4-methoxyphenyl)imino)methyl)phenol (12b)



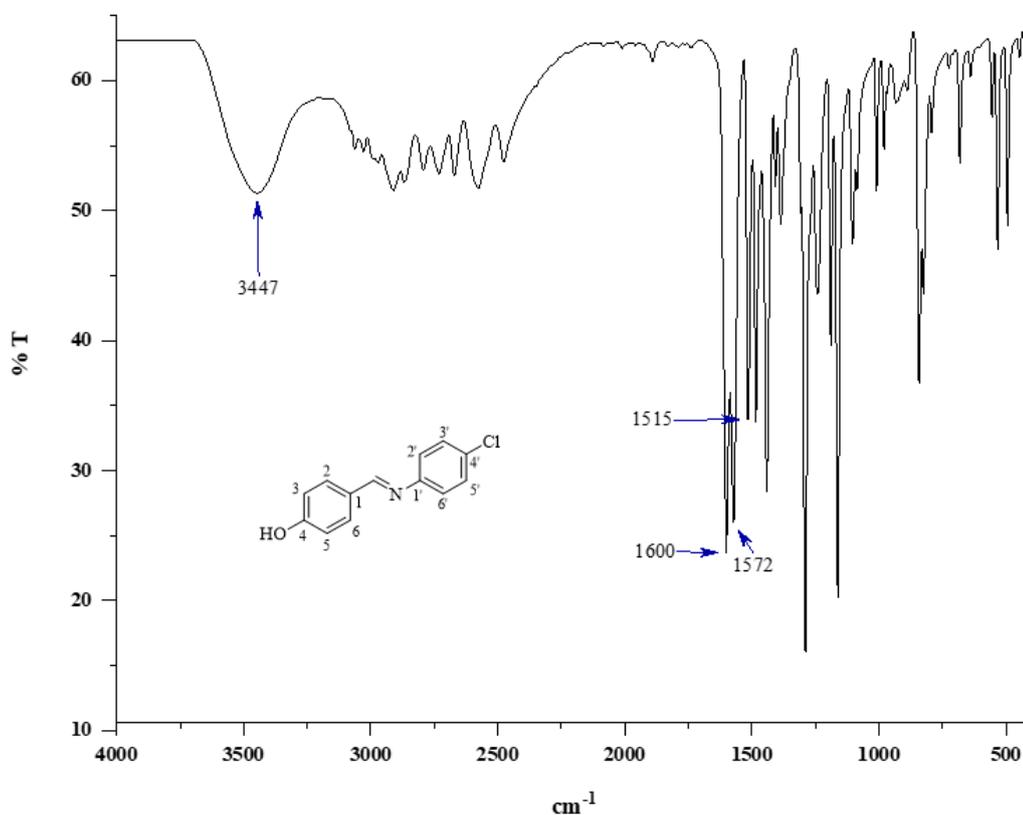
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PROCNO    1
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PULPROG   zg30
TD         65526
SOLVENT   DMSO
NS         16
DS         2
SWH        8223.685 Hz
FIDRES     0.125483 Hz
AQ         3.9846287 sec
RG         144
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DE         6.50 usec
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TD0        1
  
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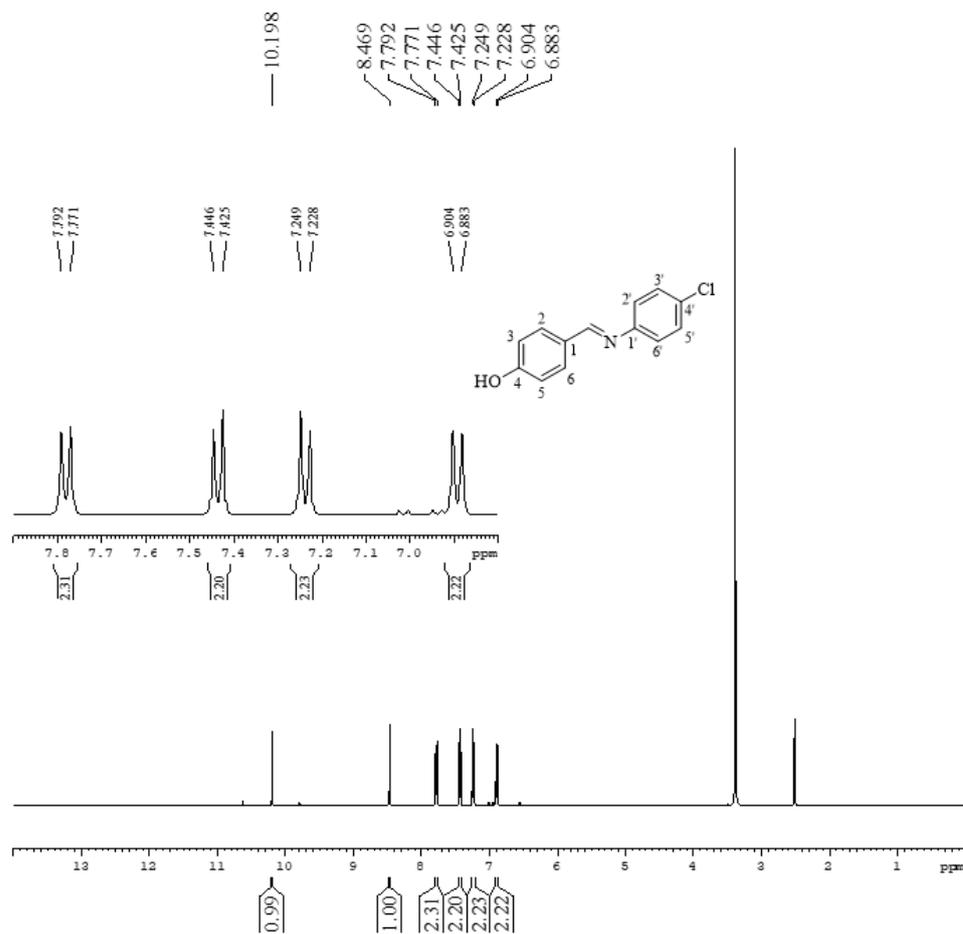
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SI        32768
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WDW       EM
SSB       0
LB        0.30 Hz
GB        0
PC        1.00
  
```

¹H NMR spectrum of (*E*)-4-(((4-methoxyphenyl)imino)methyl)phenol (12b)



IR spectrum of (*E*)-4-(((4-chlorophenyl)imino)methyl)phenol (12c)



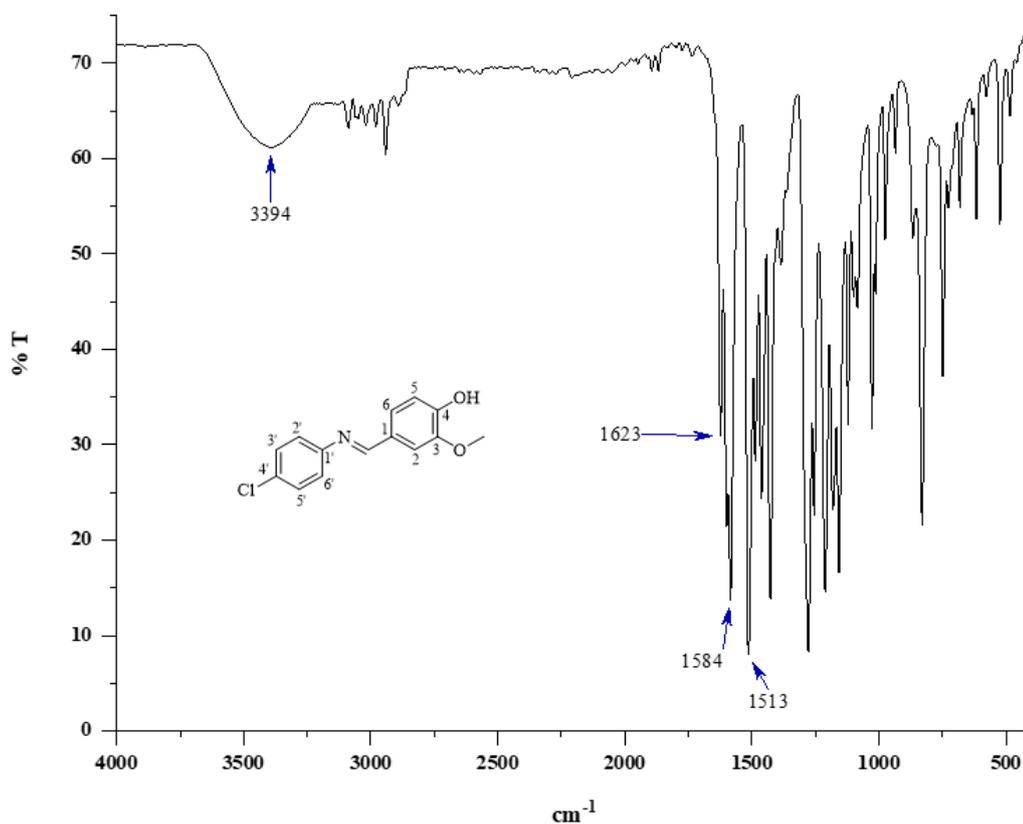
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PROCNO        1
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TD            65536
SOLVENT       DMSO
NS            16
DS            2
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FIDRES        0.125483 Hz
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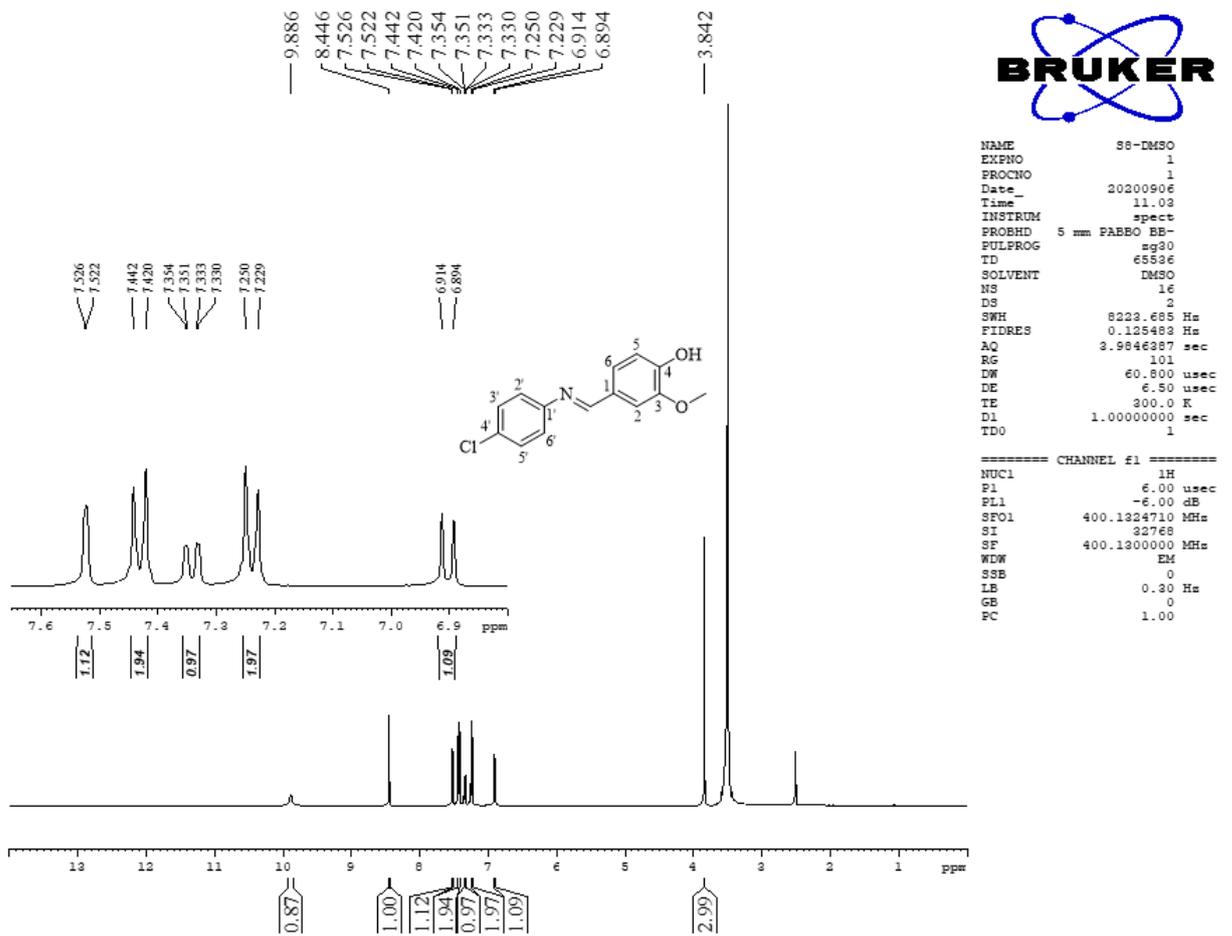
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SF           400.1300000 MHz
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SSB           0
LB            0.30 Hz
GB            0
PC            1.00
  
```

¹H NMR spectrum of *(E)*-4-(((4-chlorophenyl)imino)methyl)phenol (**12c**)



IR spectrum of *(E)*-4-(((4-chlorophenyl)imino)methyl)-2-methoxyphenol (**12d**)



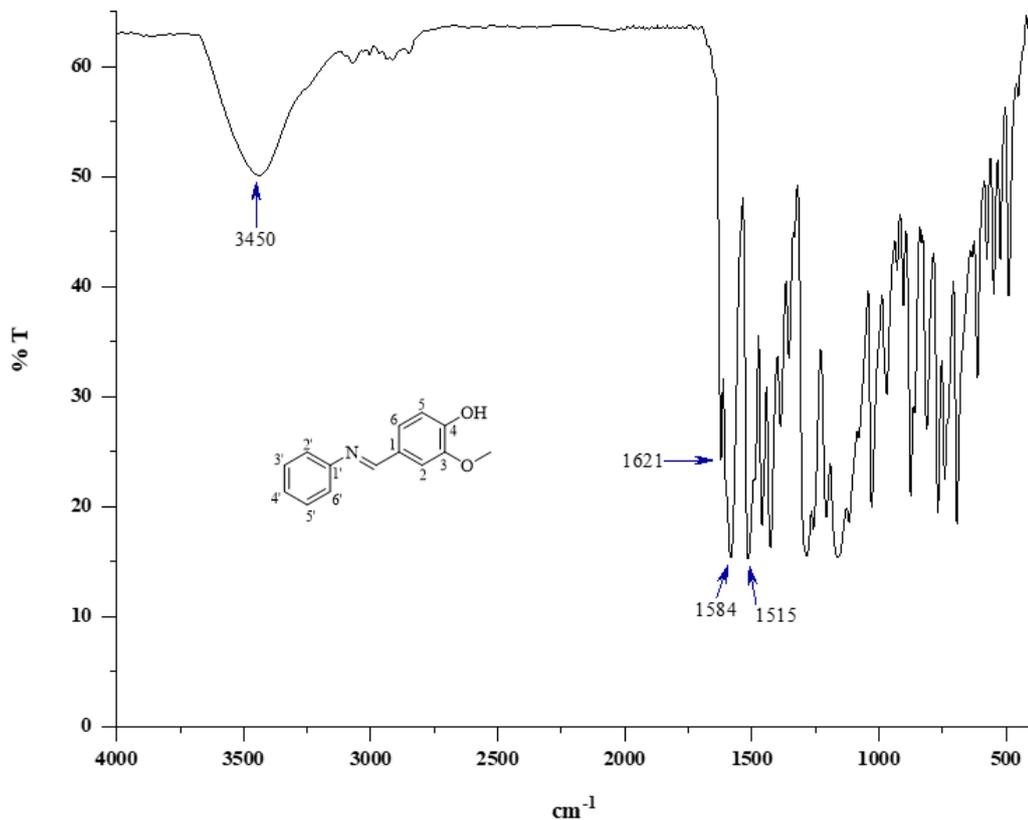
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PROCNO        1
Date_         20200906
Time          11.03
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PULPROG       zg30
TD            65536
SOLVENT       DMSO
NS            16
DS            2
SWH           8222.685 Hz
FIDRES        0.125483 Hz
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RG            401
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DE            6.50 usec
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D1            1.00000000 sec
TD0           1
  
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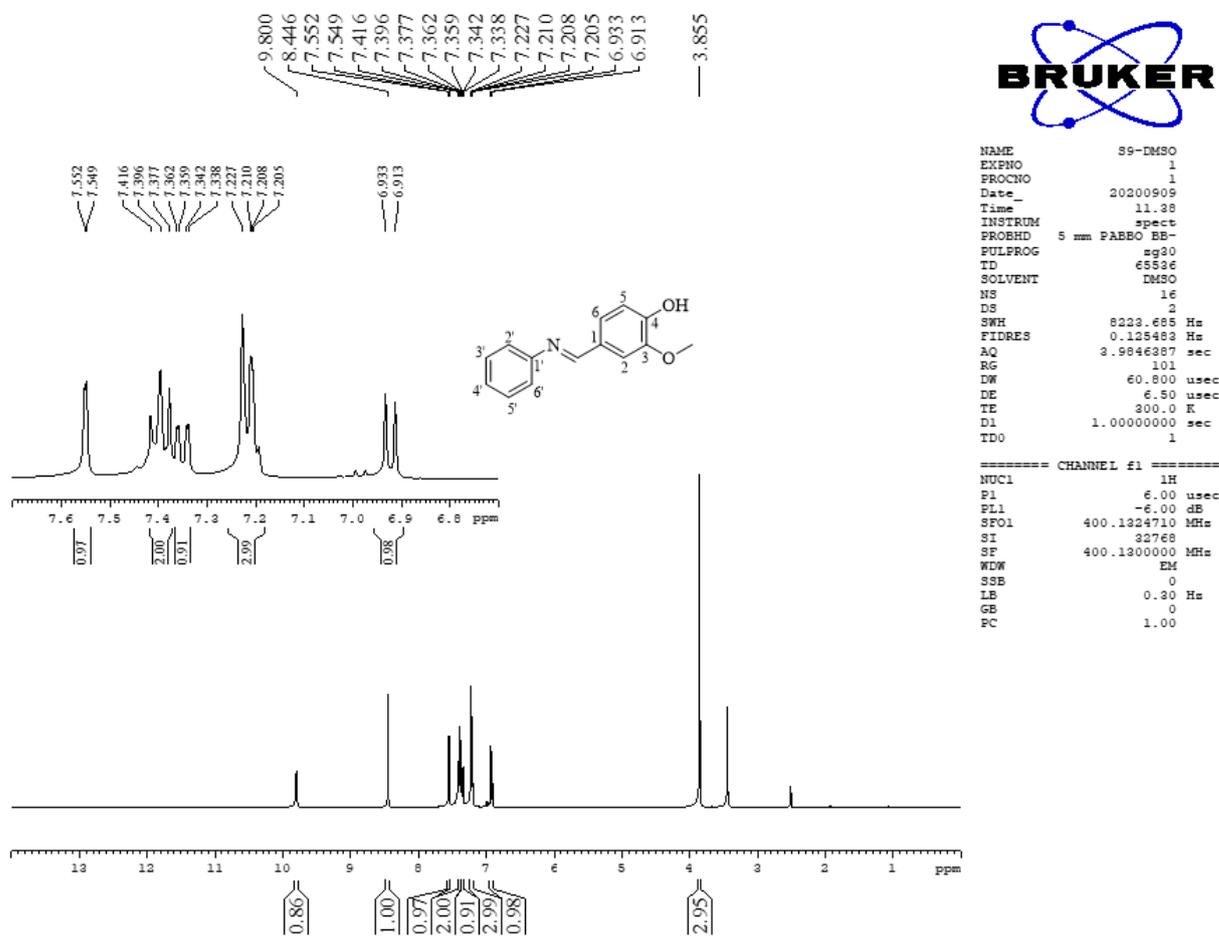
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PL1           -6.00 dB
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SSB           0
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PC            1.00
  
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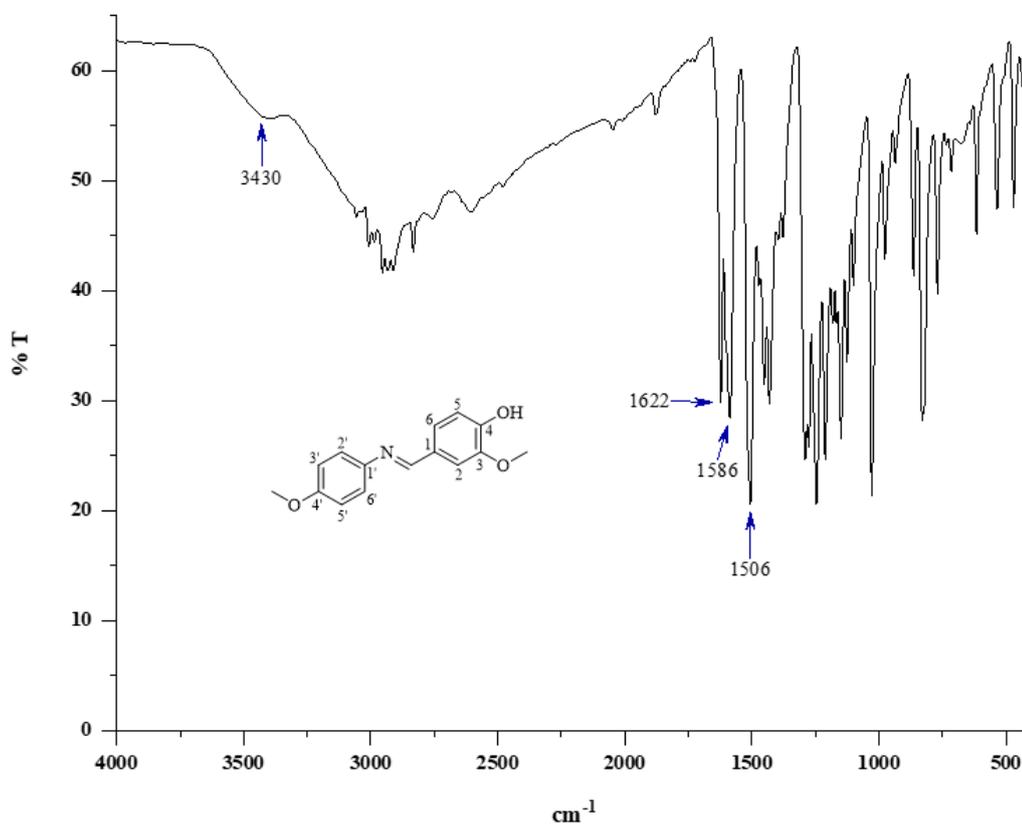
¹H NMR spectrum of (E)-4-(((4-chlorophenyl)imino)methyl)-2-methoxyphenol (12d)



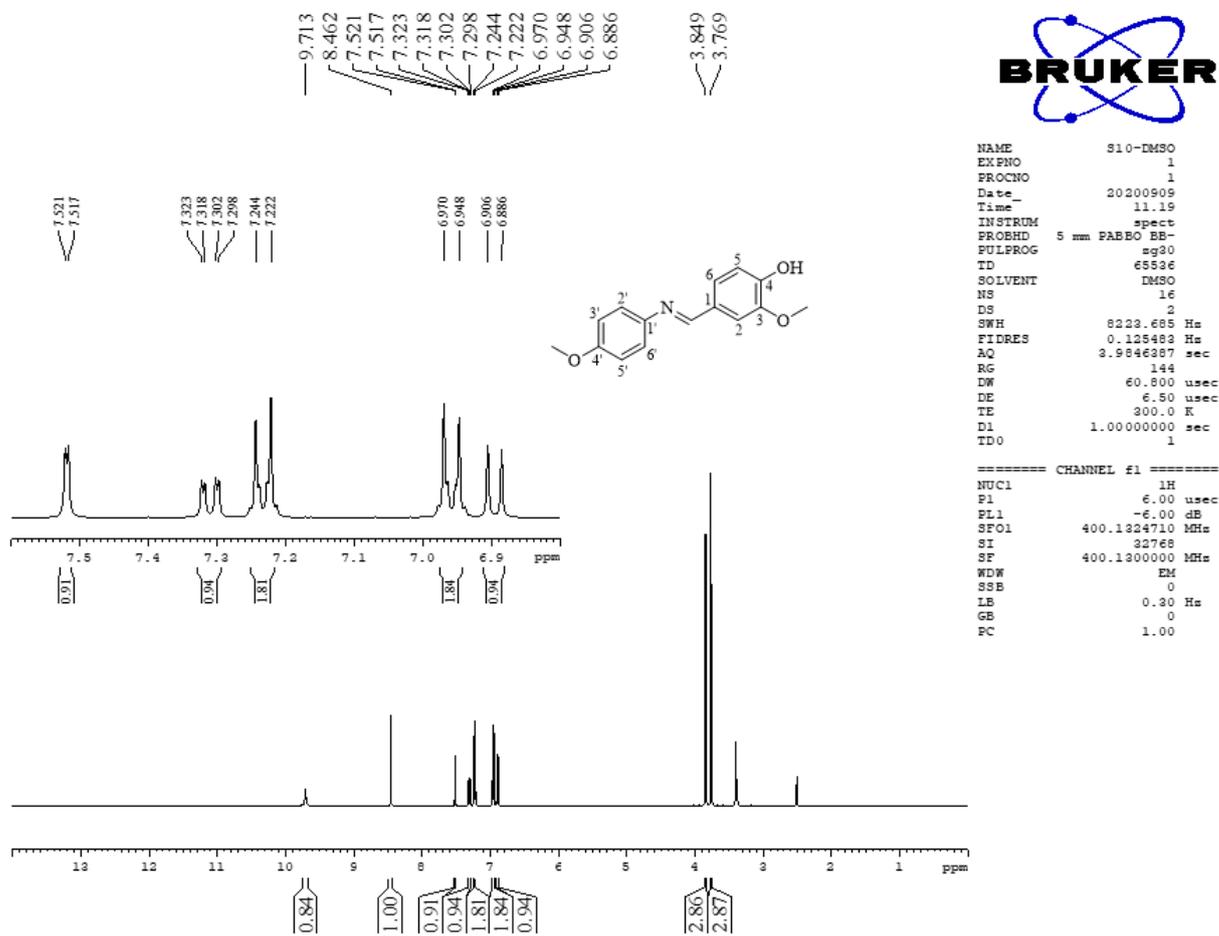
IR spectrum of (E)-2-methoxy-4-((phenylimino)methyl)phenol (12e)



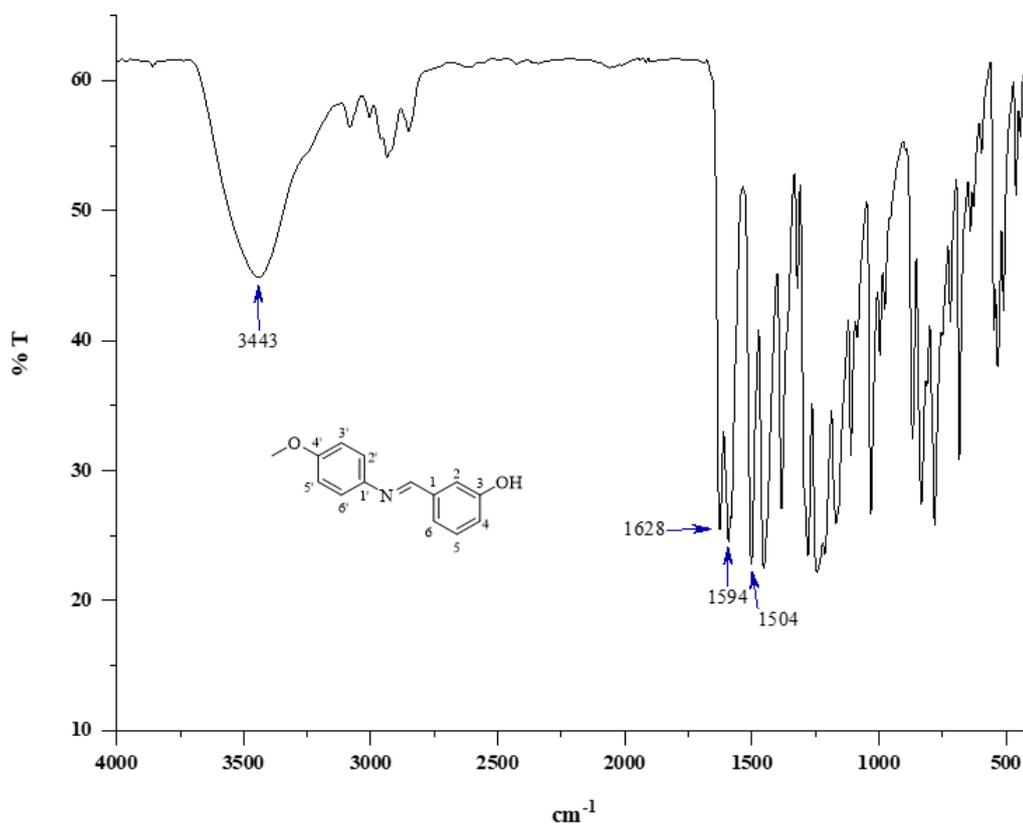
¹H NMR spectrum of (*E*)-2-methoxy-4-((phenylimino)methyl)phenol (**12e**)



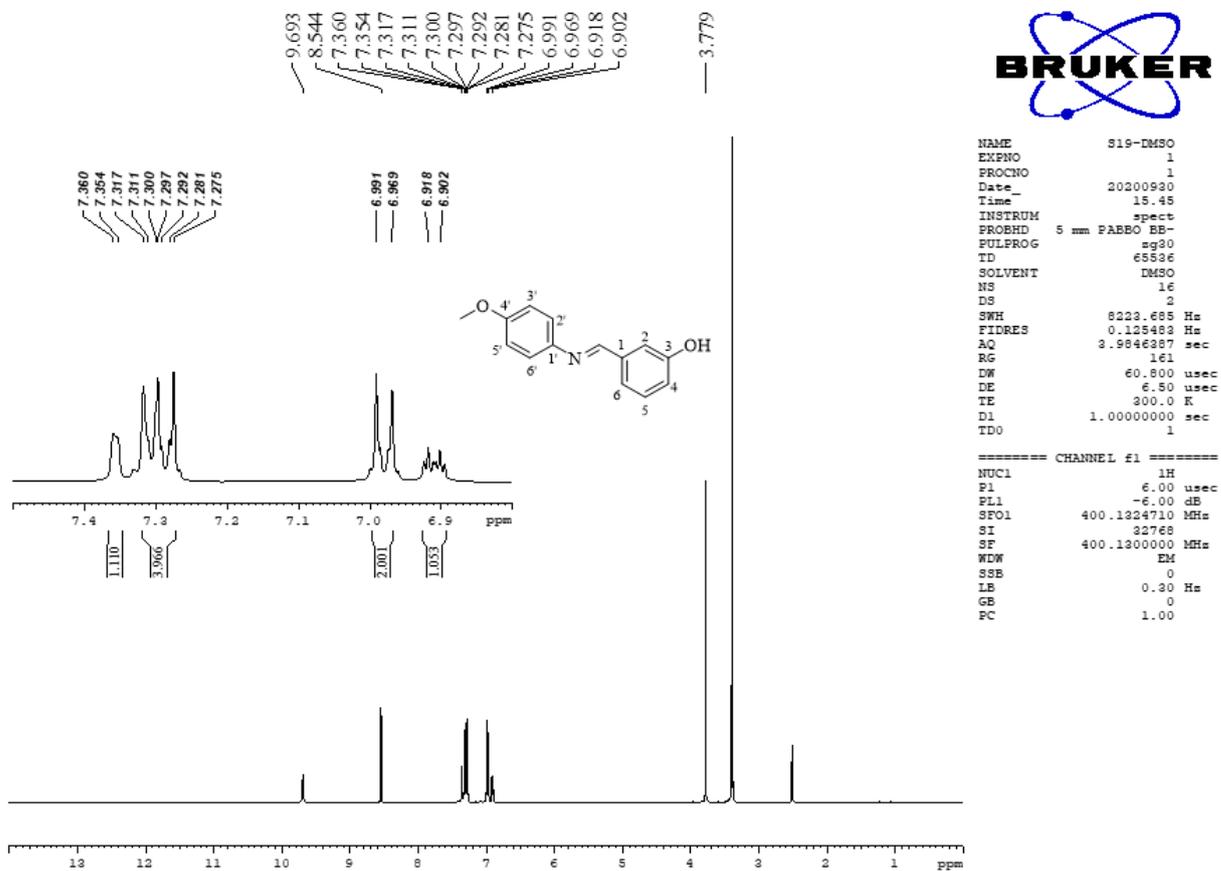
IR spectrum of (*E*)-2-methoxy-4-(((4-methoxyphenyl)imino)methyl)phenol (**12f**)



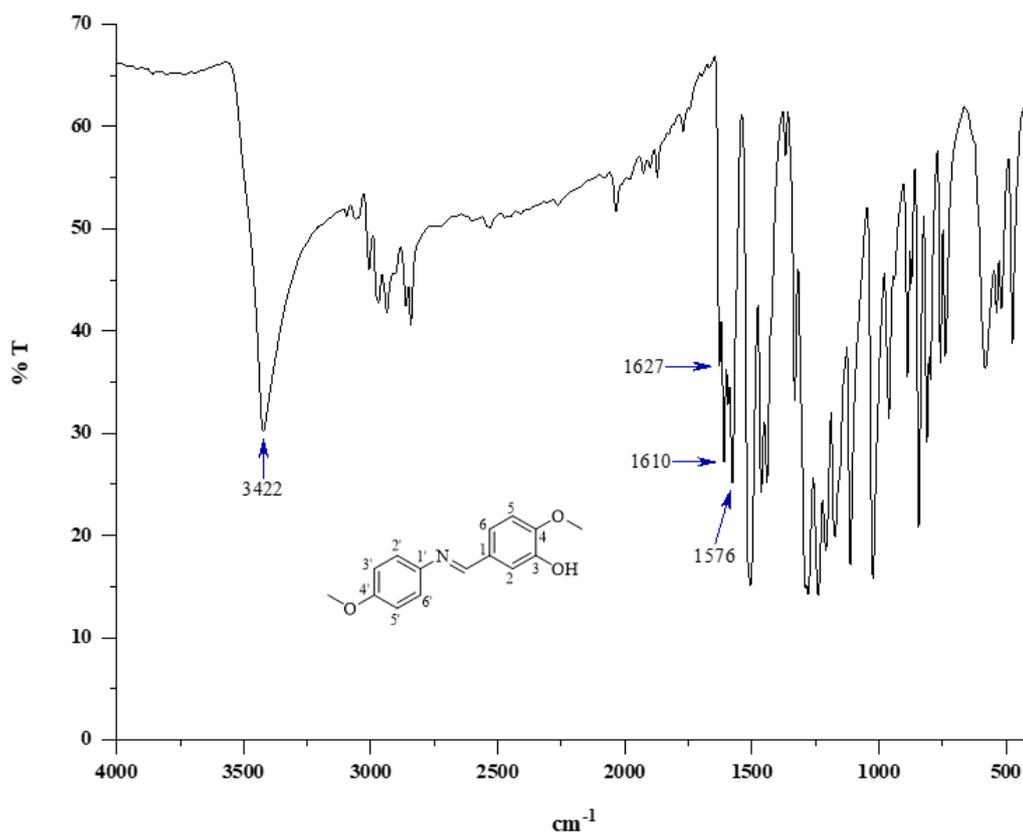
^1H NMR spectrum of (*E*)-2-methoxy-4-(((4-methoxyphenyl)imino)methyl)phenol (**12f**)



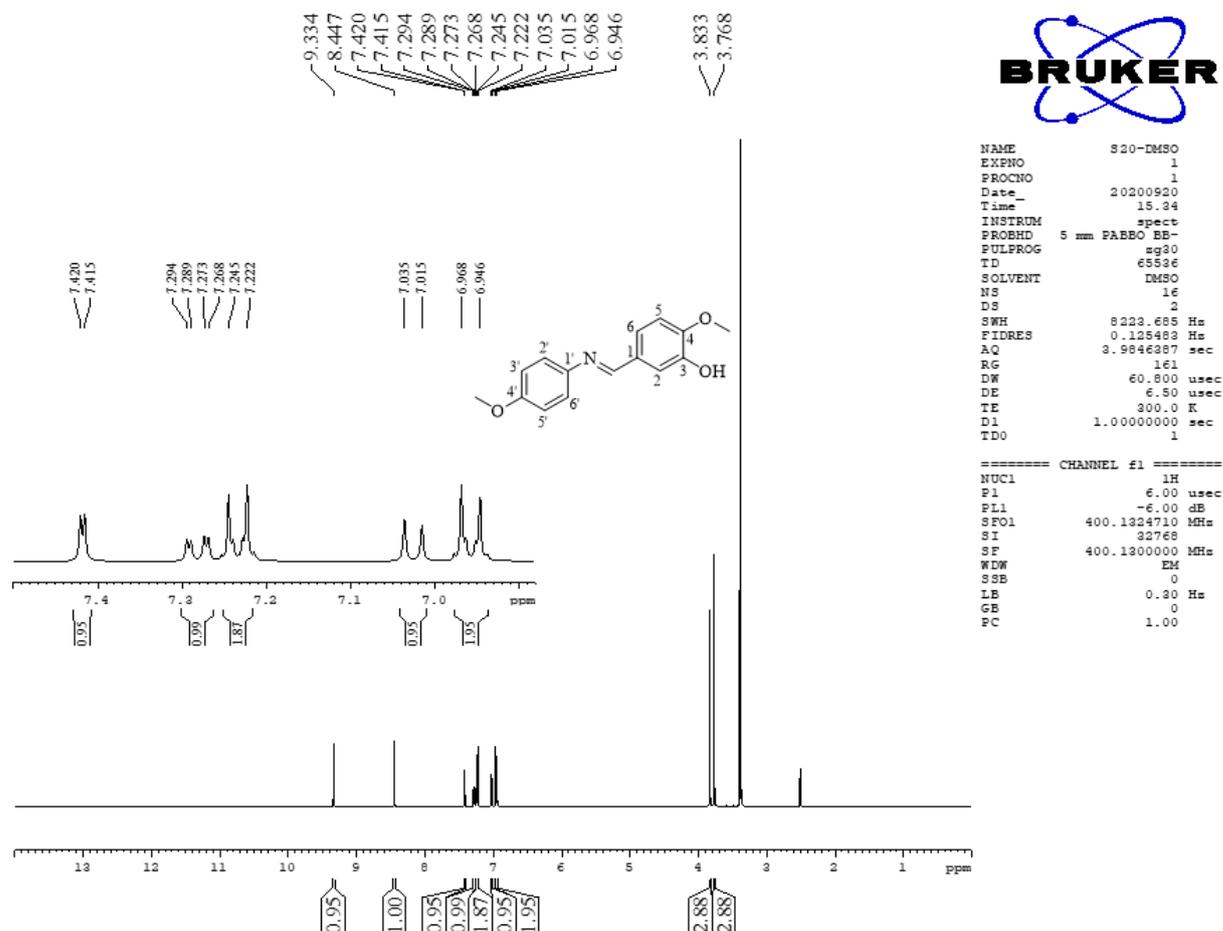
IR spectrum of (*E*)-3-(((4-methoxyphenyl)imino)methyl)phenol (**12g**)



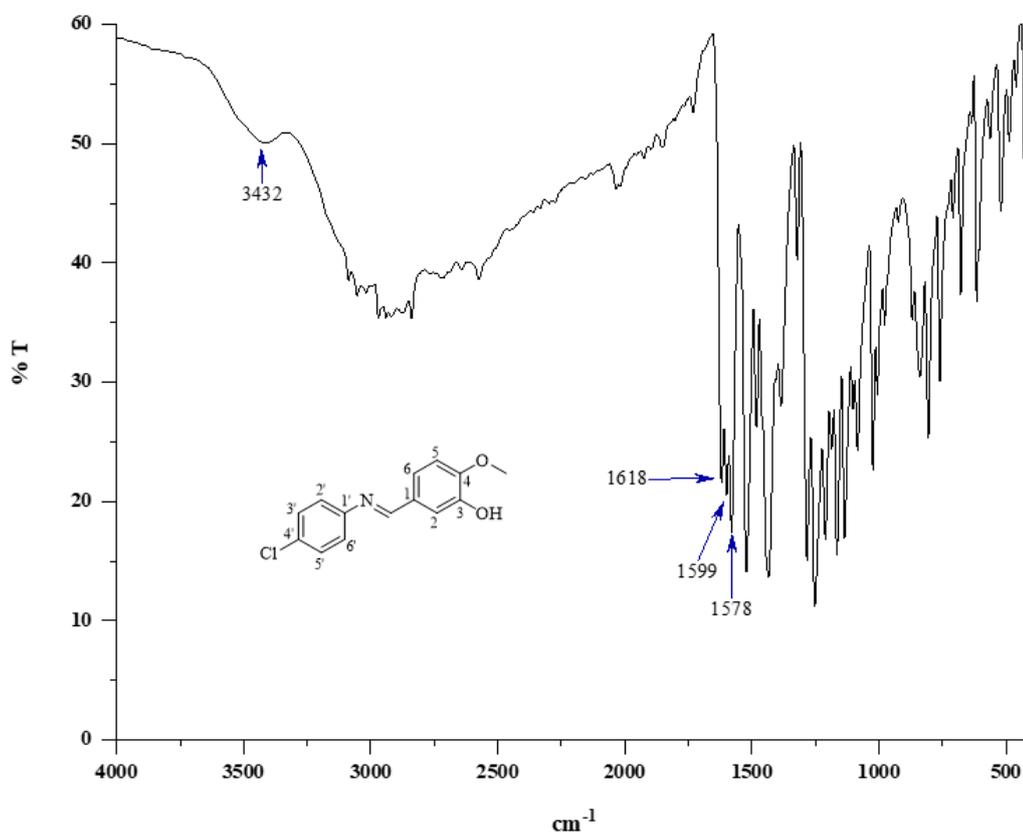
¹H NMR spectrum of (E)-3-(((4-methoxyphenyl)imino)methyl)phenol (12g)



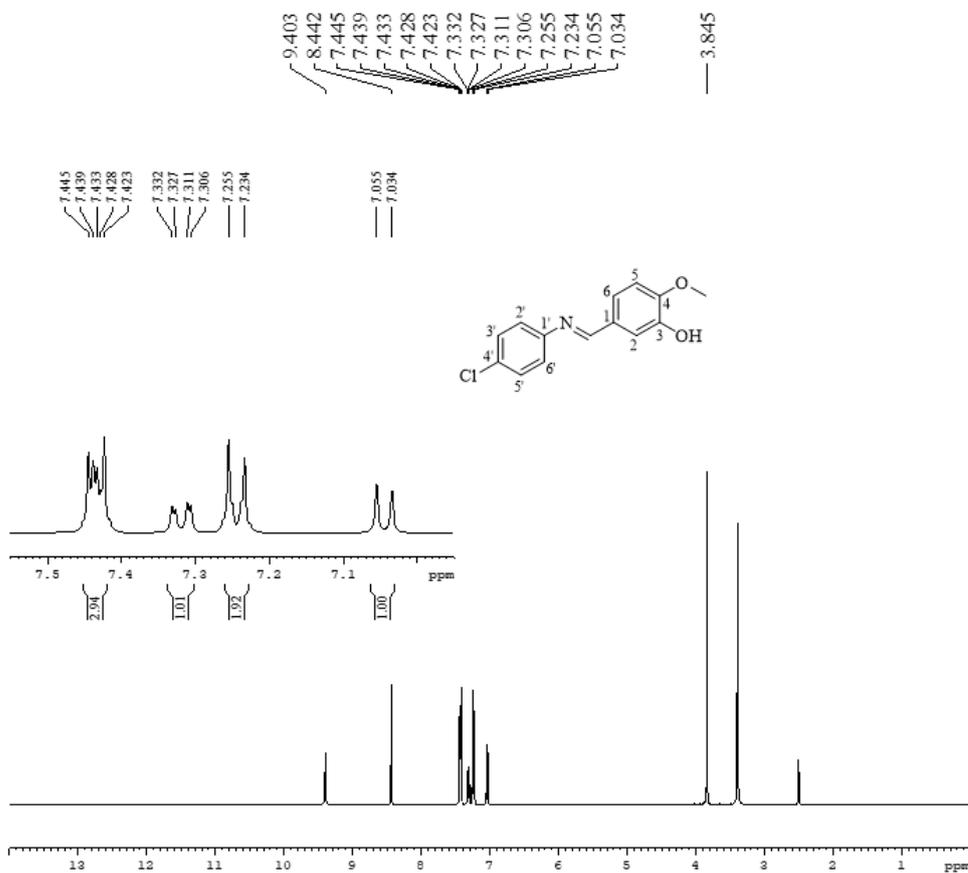
IR spectrum of (E)-2-methoxy-5-(((4-methoxyphenyl)imino)methyl)phenol (12h)



¹H NMR spectrum of (*E*)-2-methoxy-5-(((4-methoxyphenyl)imino)methyl) phenol (**12b**)



IR spectrum of (*E*)-5-(((4-chlorophenyl)imino)methyl)-2-methoxyphenol (**12i**)



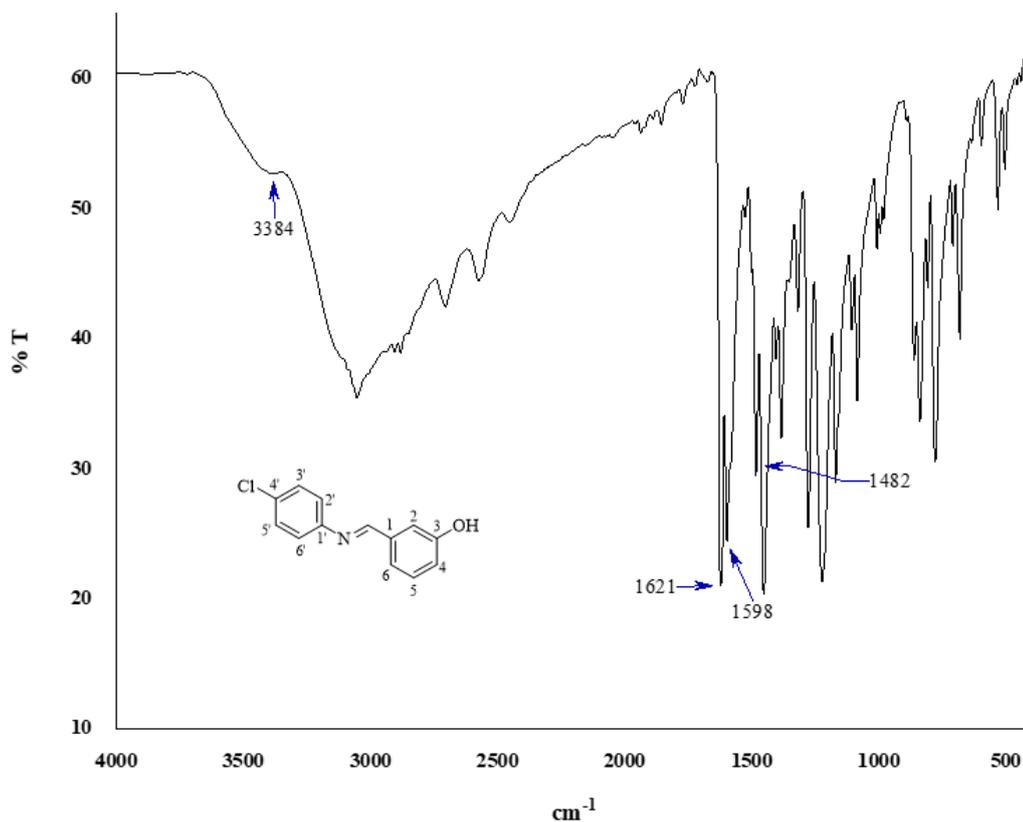
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PROCNO   1
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PULPROG  zg30
TD       65536
SOLVENT  DMSO
NS       16
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FIDRES   0.125483 Hz
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TE       300.0 K
D1       1.00000000 sec
TD0      1
  
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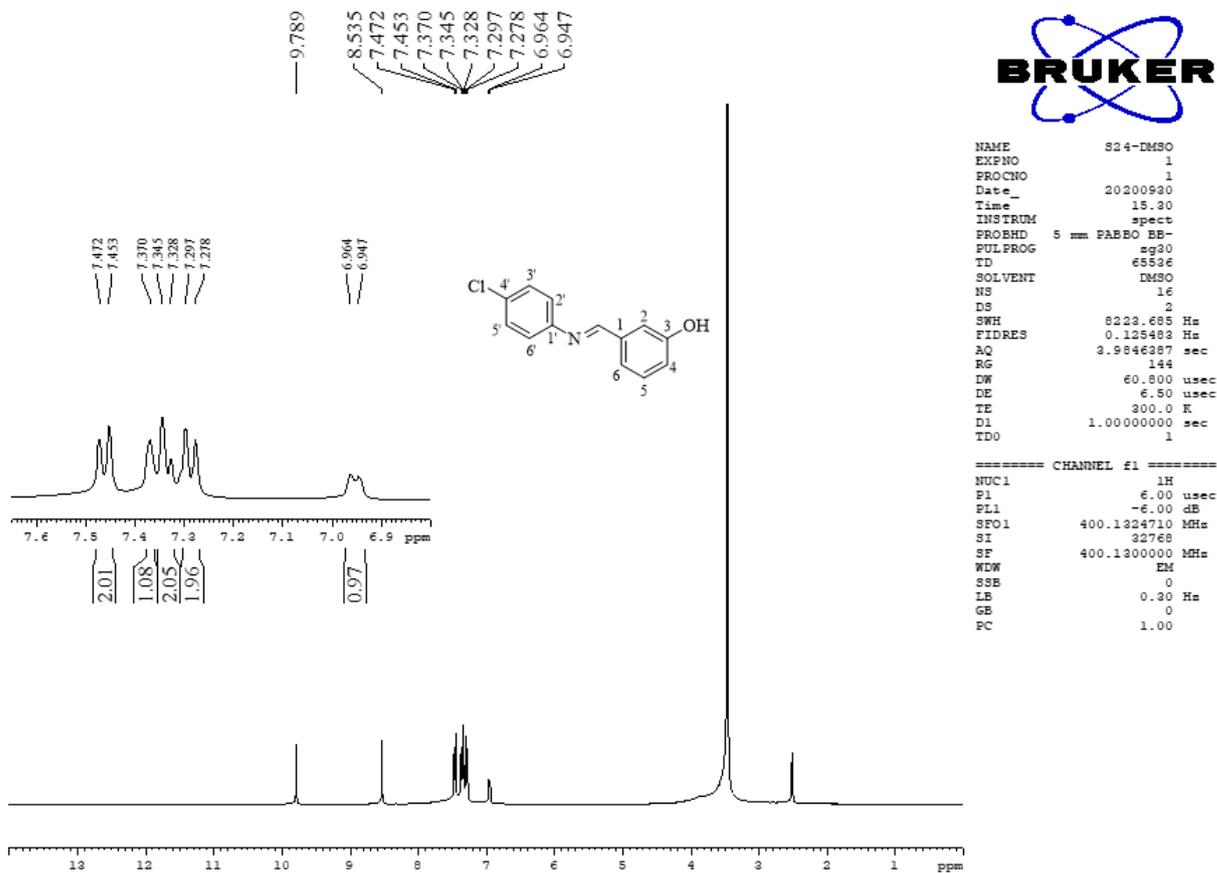
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SSB      0
LB       0.30 Hz
GB       0
PC       1.00
  
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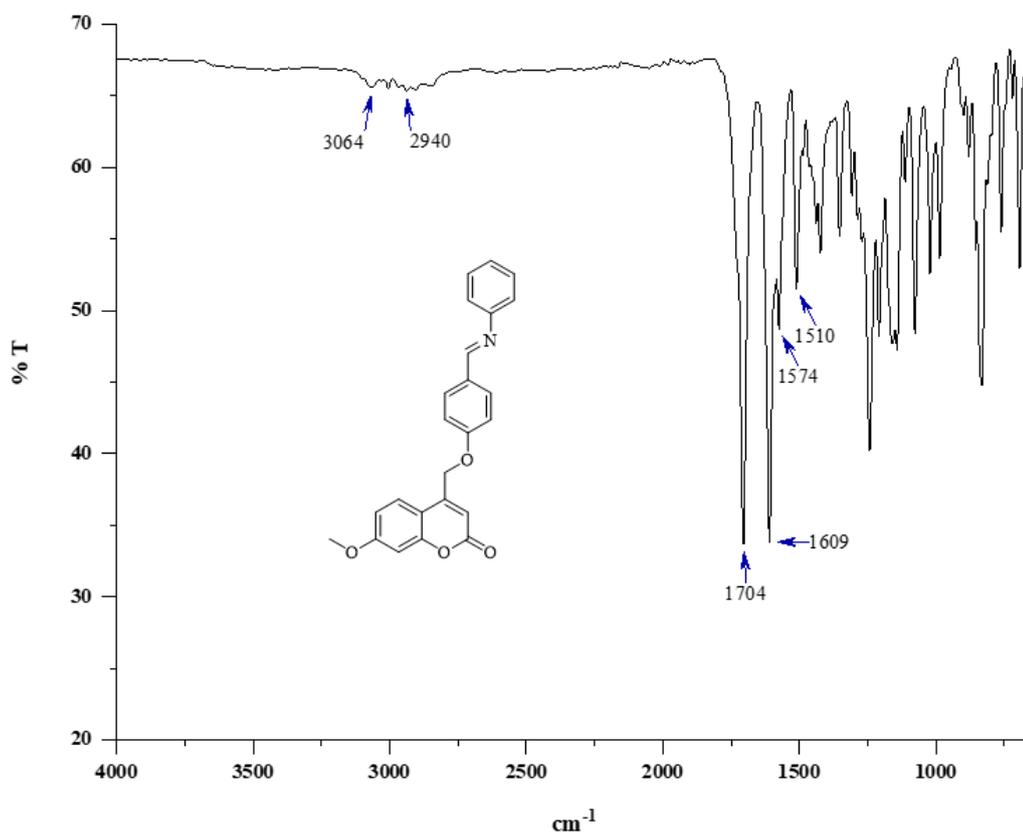
¹H NMR spectrum of (E)-5-(((4-chlorophenyl)imino)methyl)-2-methoxyphenol (12i)



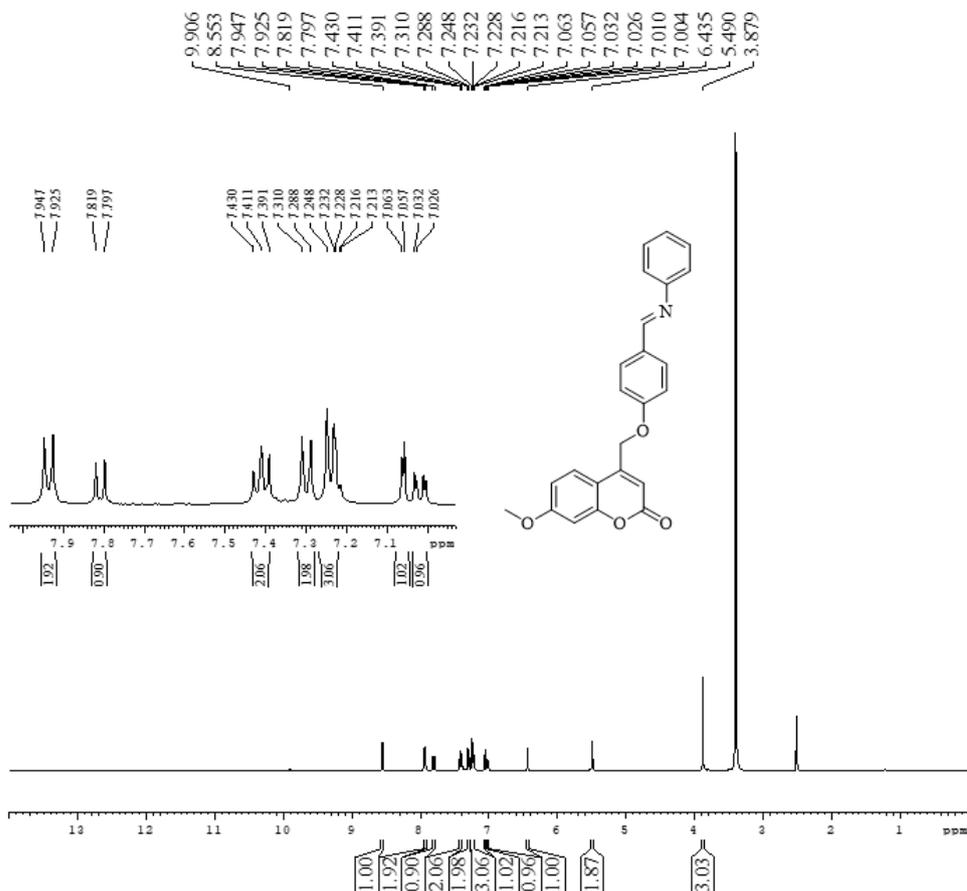
IR spectrum of (E)-3-(((4-chlorophenyl)imino)methyl)phenol (12j)



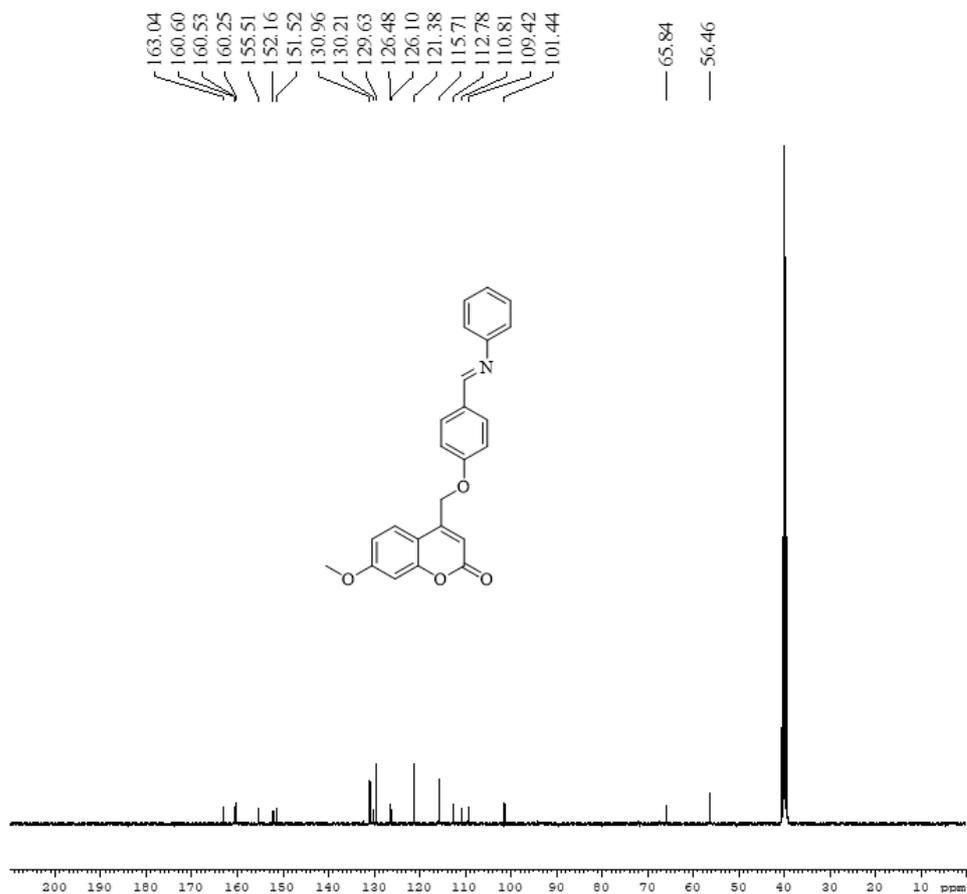
¹H NMR spectrum of *(E)*-3-(((4-chlorophenyl)imino)methyl)phenol (**12j**)



IR spectrum of *(E)*-7-methoxy-4-((4-((phenylimino)methyl)phenoxy)methyl)-2H-chromen-2-one (**13a**)



¹H NMR spectrum of (E)-7-methoxy-4-((4-((phenylimino)methyl)phenoxy)methyl)-2H-chromen-2-one (13a)



¹³C NMR spectrum of (E)-7-methoxy-4-((4-((phenylimino)methyl)phenoxy)methyl)-2H-chromen-2-one (13a)



```

NAME          S170MC-DMSO
EXPNO         1
PROCNO        1
Date_         20200909
Time         11.05
INSTRUM       spect
PROBHD        5 mm FAPBO BB-
PULPROG       zg30
TD            65536
SOLVENT       DMSO
NS            16
DS            2
SWH           8223.685 Hz
FIDRES        0.125483 Hz
AQ            3.9846387 sec
RG            181
DW            60.800 usec
DE            6.50 usec
TE            300.0 K
D1            1.00000000 sec
TD0           1

===== CHANNEL f1 =====
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P1            6.00 usec
PL1           -6.00 dB
SFO1          400.1324710 MHz
SI            32768
SF            400.1300000 MHz
WDW           EM
SSB           0
LB            0.30 Hz
GB            0
PC            1.00
  
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NAME          S170MC-DMSO
EXPNO         3
PROCNO        1
Date_         20200909
Time         11.54
INSTRUM       spect
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PULPROG       zgpg30
TD            65536
SOLVENT       DMSO
NS            2048
DS            4
SWH           24038.461 Hz
FIDRES        0.366798 Hz
AQ            1.3631988 sec
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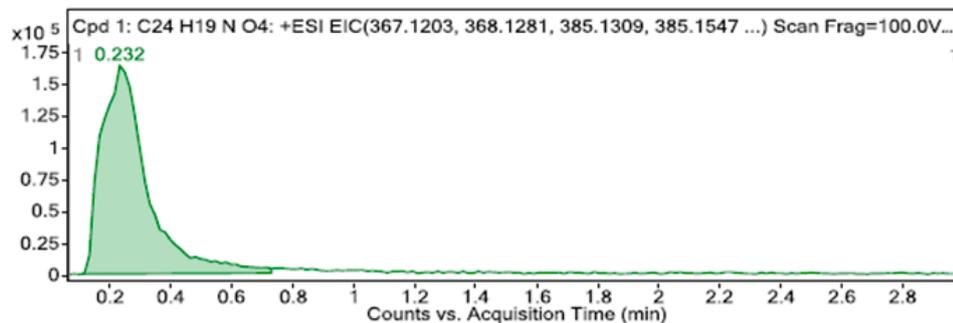
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NUC2          1H
PCPD2         80.00 usec
PL2           -6.00 dB
PL12          19.50 dB
PL13          19.50 dB
SFO2          400.1316005 MHz
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SF            100.6127690 MHz
WDW           EM
SSB           0
LB            1.00 Hz
GB            0
PC            1.40
  
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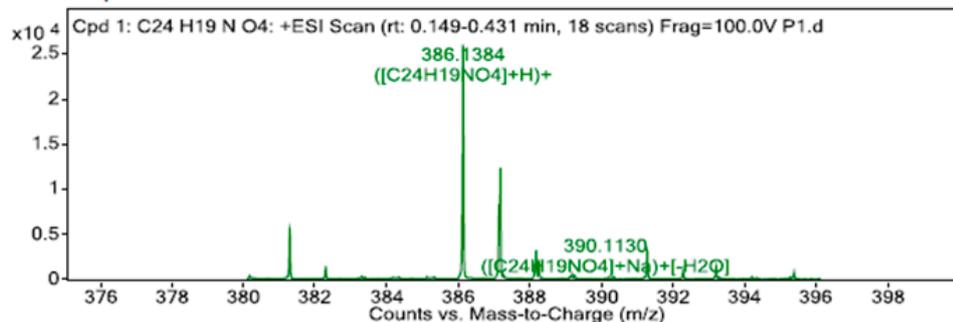
Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
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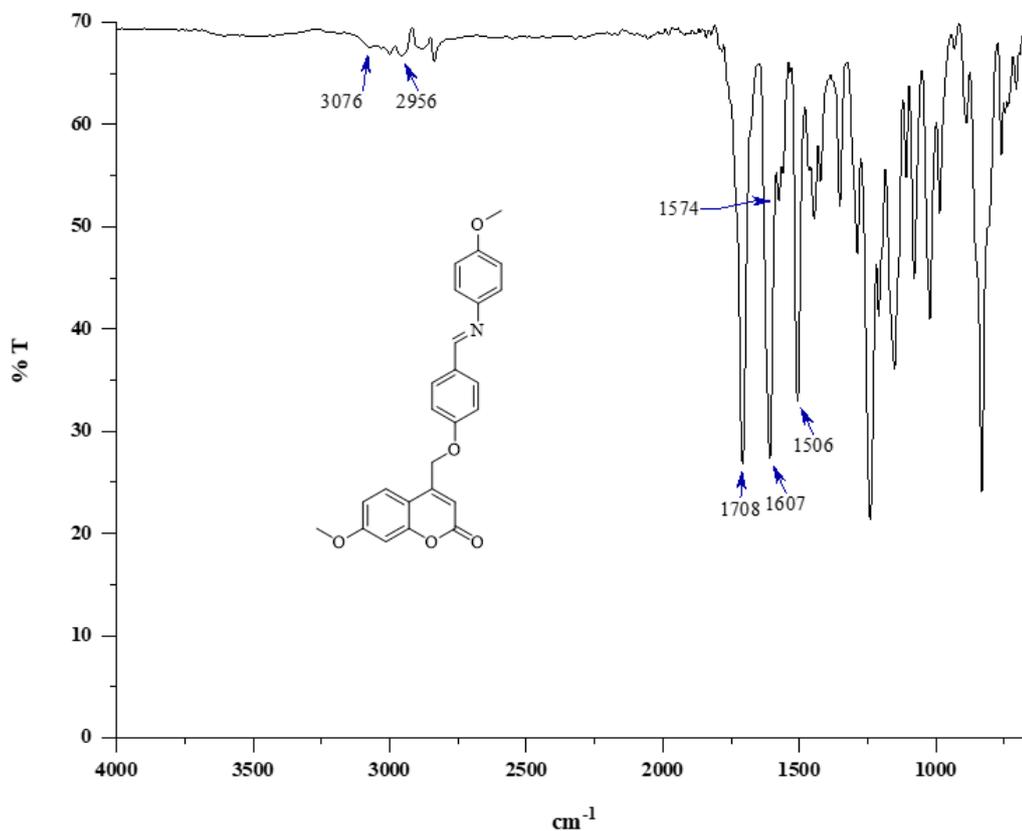
Compound Label	m/z	RT	Algorithm	Mass
Cpd 1: C ₂₄ H ₁₉ N O ₄	390.113	0.232	Find By Formula	385.1312

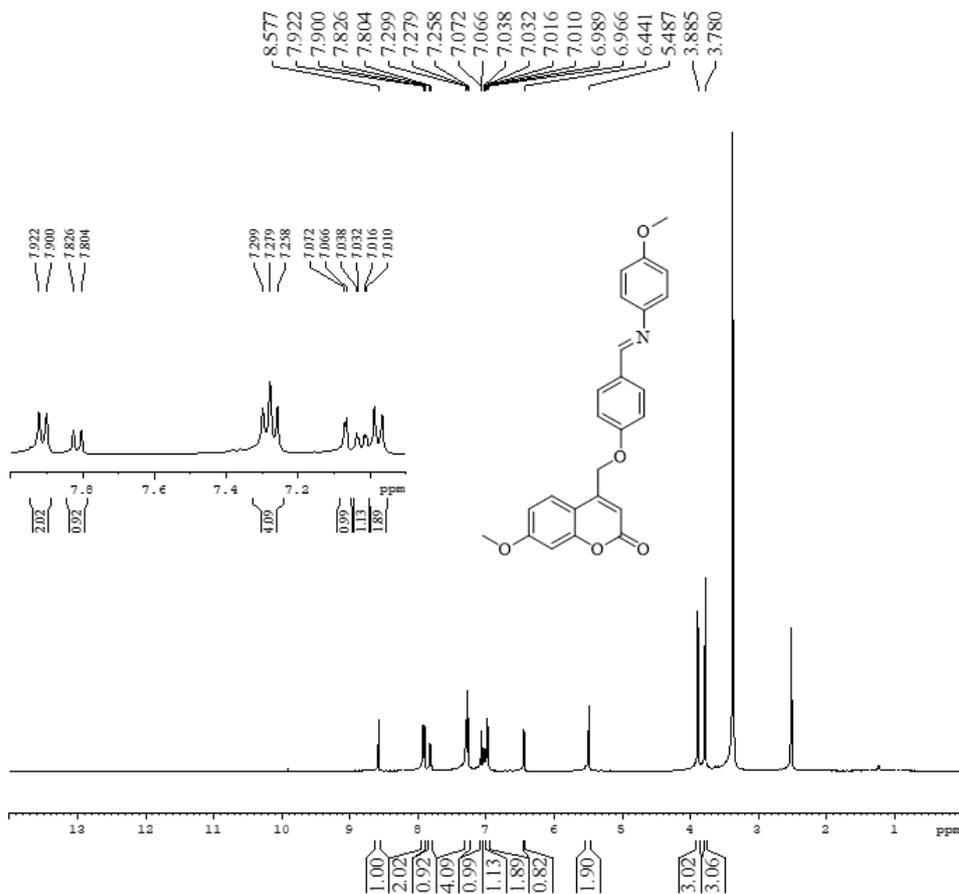


MS Zoomed Spectrum



HRMS of (*E*)-7-methoxy-4-(((4-((phenylimino)methyl)phenoxy)methyl)-2*H*-chromen-2-one (13a)

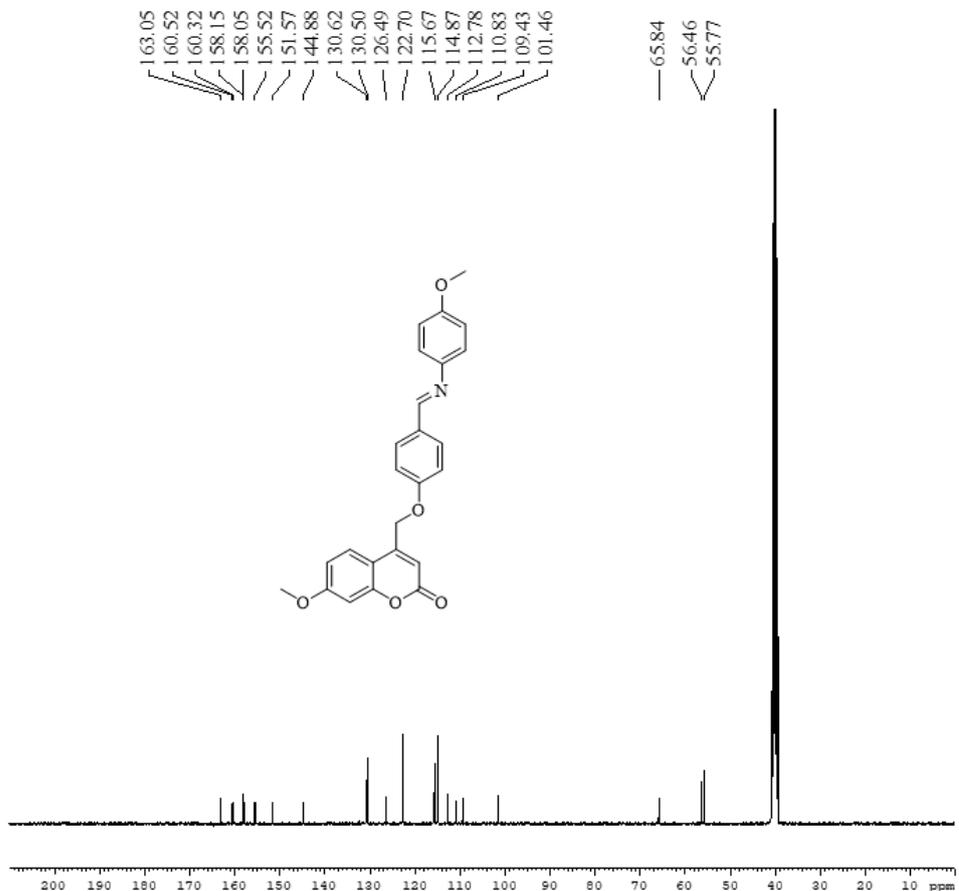




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PROCNO 1
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PULPROG sg30
TD 65536
SOLVENT DMSO
NS 16
DS 2
SWH 8223.685 Hz
FIDRES 0.125488 Hz
AQ 3.9846287 sec
RG 228
DW 60.800 usec
DE 6.50 usec
TE 300.0 K
D1 1.00000000 sec
TD0 1
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```
===== CHANNEL f1 =====
NUC1 1H
P1 6.00 usec
PL1 -6.00 dB
SF01 400.1324710 MHz
SI 32768
SF 400.13000000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00
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¹H NMR spectrum of *(E)*-7-methoxy-4-((4-((4-methoxyphenyl)imino)methyl) phenoxy)methyl)-2*H*-chromen-2-one (**13b**)



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NAME S270MC-DMSO
EXPNO 2
PROCNO 1
Date_ 20200917
Time 12.29
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG sggg30
TD 65536
SOLVENT DMSO
NS 15260
DS 4
SWH 24038.461 Hz
FIDRES 0.366798 Hz
AQ 1.3631988 sec
RG 203
DW 20.800 usec
DE 6.50 usec
TE 300.0 K
D1 2.00000000 sec
D11 0.03000000 sec
TD0 1
```

```
===== CHANNEL f1 =====
NUC1 13C
P1 12.00 usec
PL1 -1.00 dB
SF01 100.6228298 MHz
```

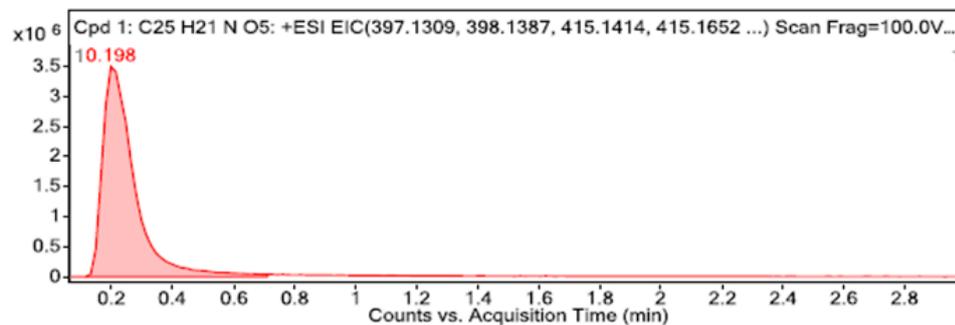
```
===== CHANNEL f2 =====
CPDPRG2 waltz65
NUC2 1H
PCPD2 80.00 usec
PL2 -6.00 dB
PL12 19.50 dB
PL13 19.50 dB
SF02 400.1316005 MHz
SI 32768
SF 100.6127690 MHz
WDW EM
SSB 0
LB 1.00 Hz
GB 0
PC 1.40
```

¹³C NMR spectrum of *(E)*-7-methoxy-4-((4-((4-methoxyphenyl)imino)methyl) phenoxy)methyl)-2*H*-chromen-2-one (**13b**)

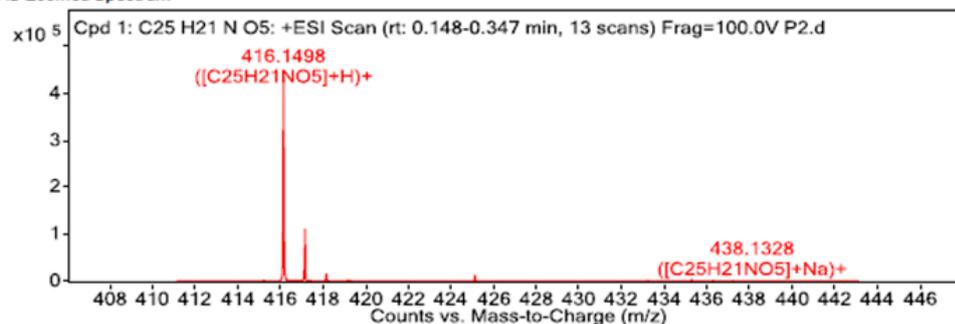
Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C ₂₅ H ₂₁ N O ₅	0.198	415.1424	432183	C ₂₅ H ₂₁ N O ₅	415.142	1.13

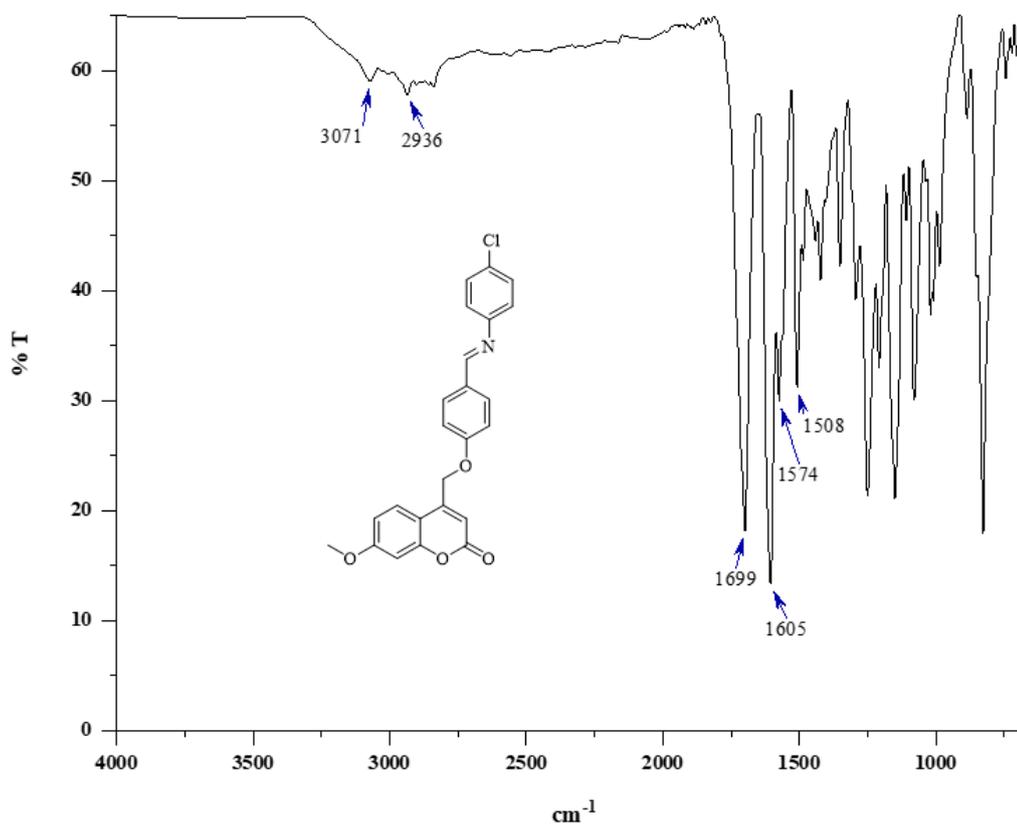
Compound Label	m/z	RT	Algorithm	Mass
Cpd 1: C ₂₅ H ₂₁ N O ₅	416.1498	0.198	Find By Formula	415.1424

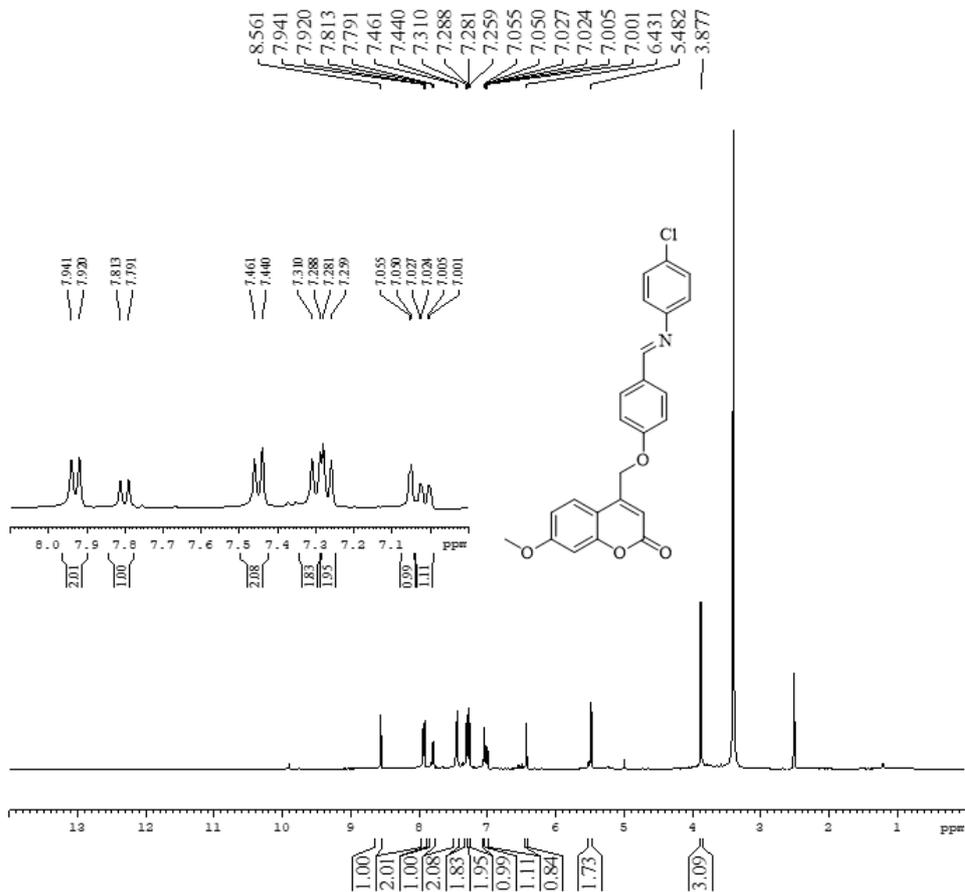


MS Zoomed Spectrum



HRMS of (*E*)-7-methoxy-4-((4-(((4-methoxyphenyl)imino)methyl) phenoxy)methyl)-2*H*-chromen-2-one (**13b**)





¹H NMR spectrum of *(E)*-4-((4-(((4-chlorophenyl)imino)methyl)phenoxy)methyl)-7-methoxy-2*H*-chromen-2-one (13c)

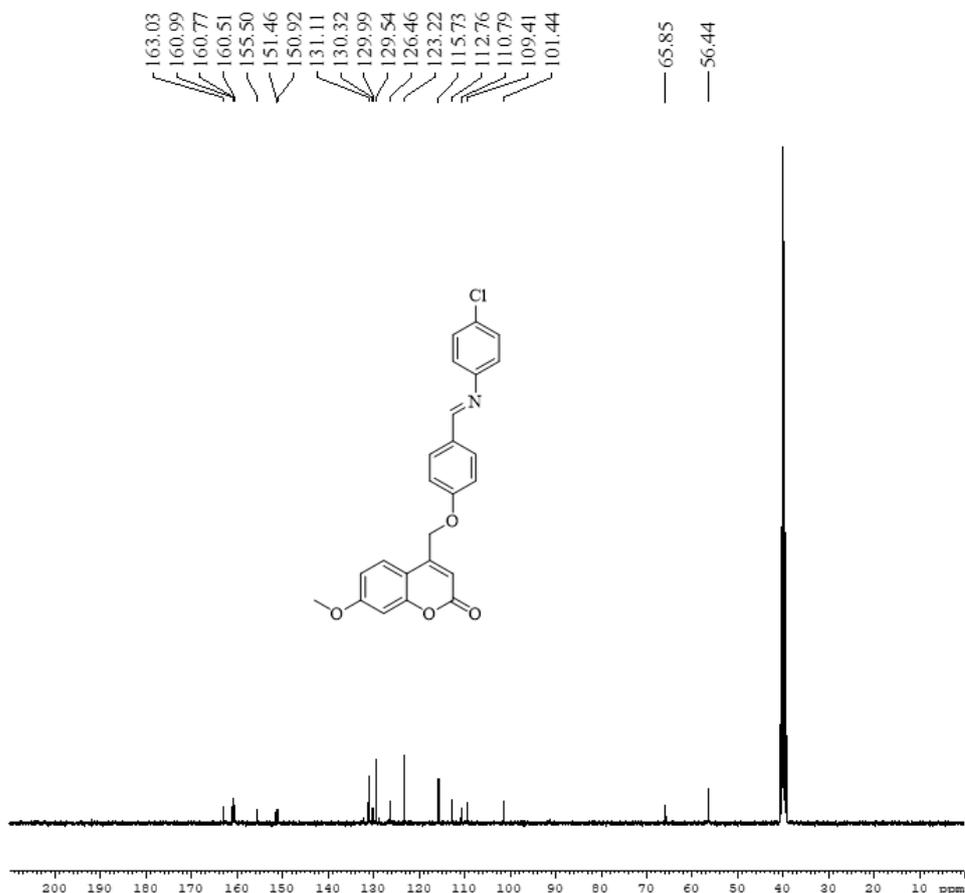


```

NAME      S770MC-DMSO
EXPNO    1
PROCNO   1
Date_    20200909
Time     8.58
INSTRUM spect
PROBHD   5 mm PABBO BB-
PULPROG  sg30
TD       65536
SOLVENT  DMSO
NS       16
DS       2
SWH      8222.685 Hz
FIDRES   0.125488 Hz
AQ       3.9846387 sec
RG       161
DW       60.800 usec
DE       6.50 usec
TE       300.0 K
D1       1.00000000 sec
TD0      1

===== CHANNEL f1 =====
NUC1     1H
P1       -6.00 usec
PL1      -6.00 dB
SFO1     400.1324710 MHz
SI       32768
SF       400.1300000 MHz
WDW      EM
SSB      0
LB       0.30 Hz
GB       0
PC       1.00

```



¹³C NMR spectrum of *(E)*-4-((4-(((4-chlorophenyl)imino)methyl)phenoxy)methyl)-7-methoxy-2*H*-chromen-2-one (13c)



```

NAME      S770MC-DMSO
EXPNO    2
PROCNO   1
Date_    20200909
Time     18.02
INSTRUM spect
PROBHD   5 mm PABBO BB-
PULPROG  sgpg30
TD       65536
SOLVENT  DMSO
NS       2048
DS       4
SWH      24028.461 Hz
FIDRES   0.366798 Hz
AQ       1.3631988 sec
RG       203
DW       20.800 usec
DE       6.50 usec
TE       300.0 K
D1       2.00000000 sec
D11      0.03000000 sec
TD0      1

===== CHANNEL f1 =====
NUC1     13C
P1       12.00 usec
PL1      -1.00 dB
SFO1     100.6228298 MHz

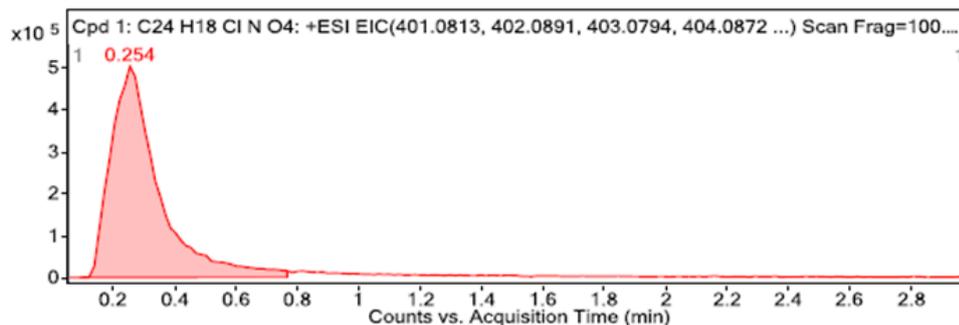
===== CHANNEL f2 =====
CFDPRG2  waltz65
NUC2     1H
PCPD2    80.00 usec
PL2      -6.00 dB
PL12     19.50 dB
PL13     19.50 dB
SFO2     400.1316005 MHz
SI       32768
SF       100.6127690 MHz
WDW      EM
SSB      0
LB       1.00 Hz
GB       0
PC       1.40

```

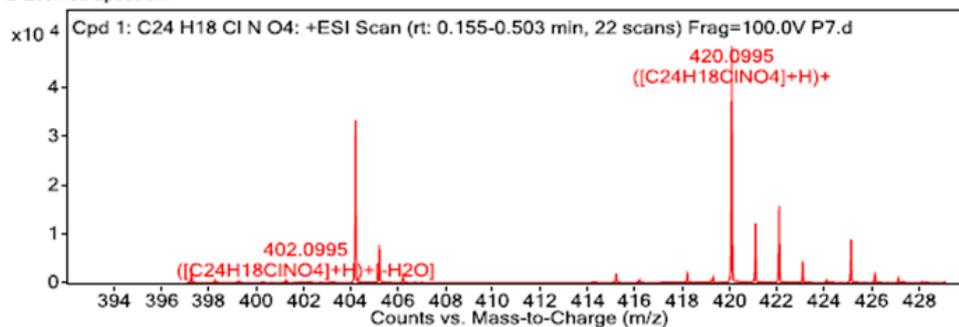
Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C ₂₄ H ₁₈ ClN ₄ O ₄	0.254	419.092	49039	C ₂₄ H ₁₈ ClN ₄ O ₄	419.0924	-1.06

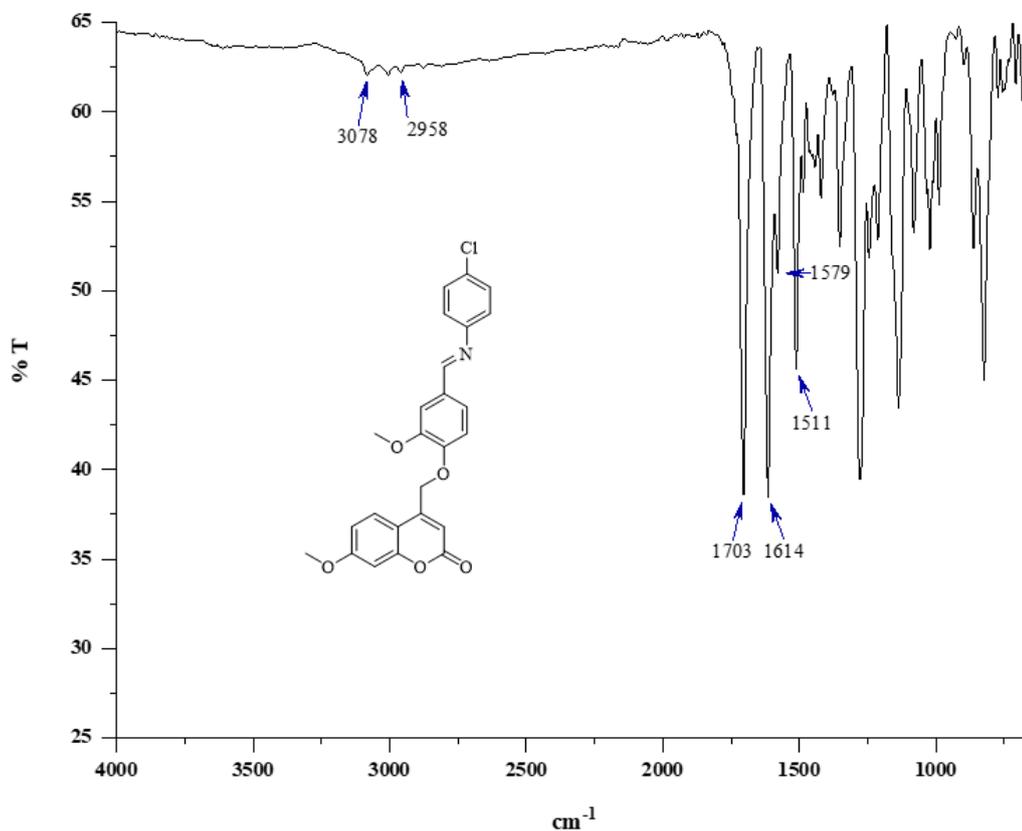
Compound Label	m/z	RT	Algorithm	Mass
Cpd 1: C ₂₄ H ₁₈ ClN ₄ O ₄	420.0995	0.254	Find By Formula	419.092

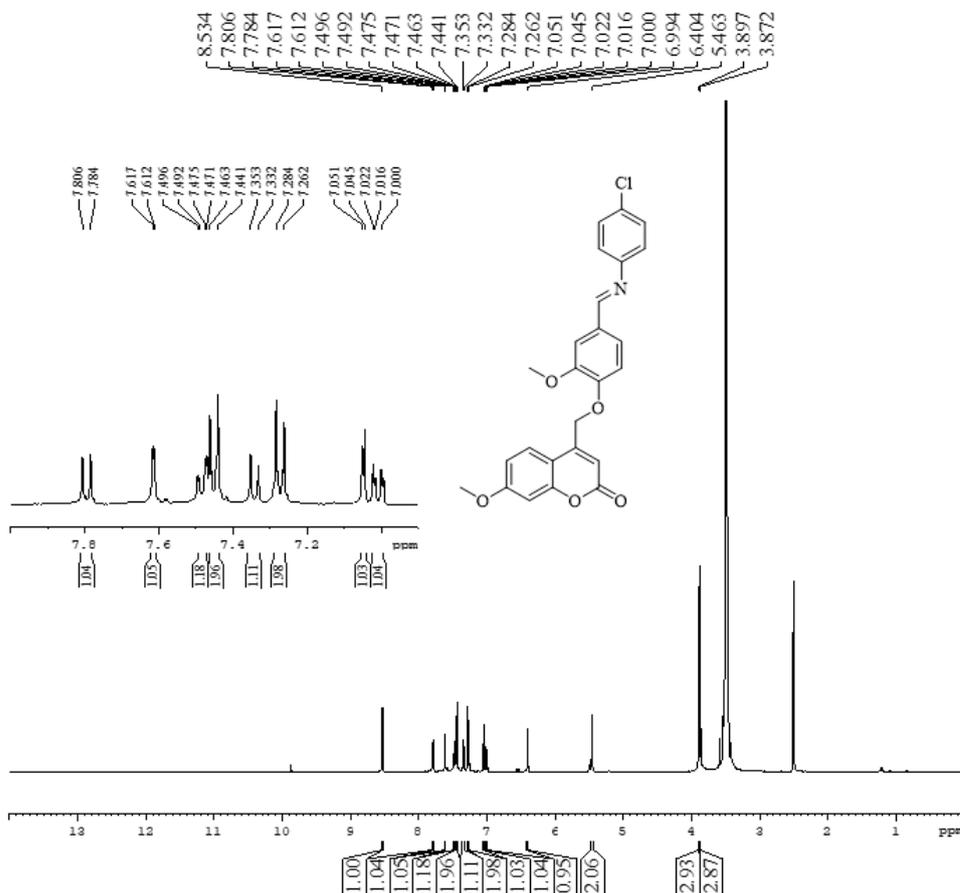


MS Zoomed Spectrum



HRMS of (*E*)-4-((4-(((4-chlorophenyl)imino)methyl)phenoxy)methyl)-7-methoxy-2*H*-chromen-2-one (13c)





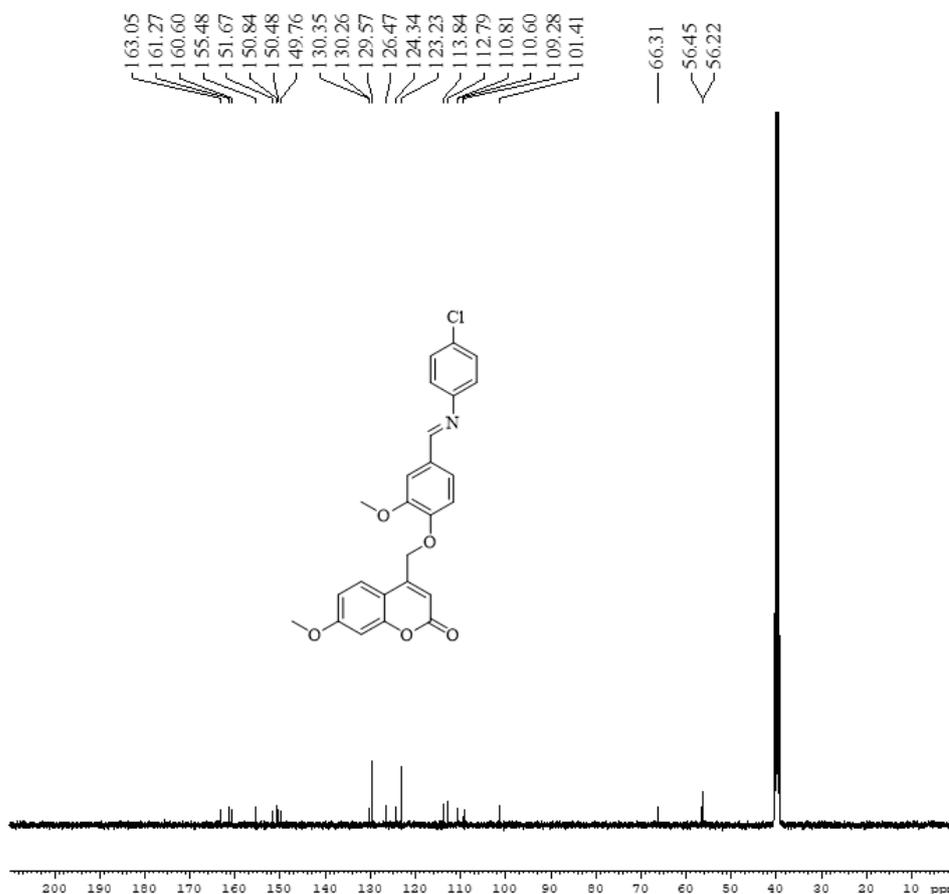
¹H NMR spectrum of (*E*)-4-((4-(((4-chlorophenyl)imino)methyl)-2-methoxy phenoxy)methyl)-7-methoxy-2H-chromen-2-one (**13d**)



```

NAME      S870MC-DMSO
EXPNO     1
PROCNO    1
Date_     20200909
Time      9.16
INSTRUM   spect
PROBHD    5 mm PABBO BB-
PULPROG   sg30
TD         65536
SOLVENT   DMSO
NS         16
DS         2
SWH        8223.685 Hz
FIDRES     0.126483 Hz
AQ         3.9846387 sec
RG         114
DW         60.800 usec
DE         6.50 usec
TE         300.0 K
D1         1.00000000 sec
TDO        1

===== CHANNEL f1 =====
NUC1      1H
P1         6.00 usec
PL1        -6.00 dB
SFO1      400.1324710 MHz
SI         32768
SF         400.1300000 MHz
WDW        EM
SSB        0
LB         0.30 Hz
GB         0
PC         1.00
  
```



¹³C NMR spectrum of (*E*)-4-((4-(((4-chlorophenyl)imino)methyl)-2-methoxy phenoxy)methyl)-7-methoxy-2H-chromen-2-one (**13d**)



```

NAME      S870MC-DMSO
EXPNO     2
PROCNO    1
Date_     20200909
Time      17.20
INSTRUM   spect
PROBHD    5 mm PABBO BB-
PULPROG   sgpg30
TD         65536
SOLVENT   DMSO
NS         1984
DS         4
SWH        24038.461 Hz
FIDRES     0.366798 Hz
AQ         1.3631988 sec
RG         803
DW         20.800 usec
DE         6.50 usec
TE         300.0 K
D1         2.00000000 sec
D11        0.03000000 sec
TDO        1

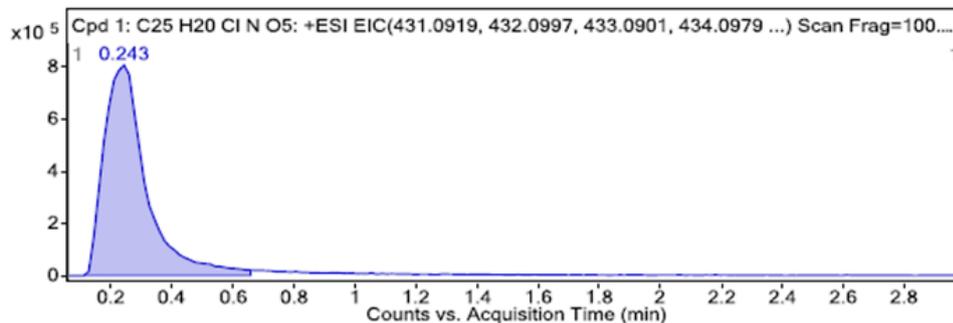
===== CHANNEL f1 =====
NUC1      13C
P1         12.00 usec
PL1        -1.00 dB
SFO1      100.6228298 MHz

===== CHANNEL f2 =====
CPDPRG2   waltz65
NUC2      1H
PCPD2     80.00 usec
PL2        -6.00 dB
PL12       19.50 dB
PL13       19.50 dB
SFO2      400.1316005 MHz
SI         32768
SF         100.6127690 MHz
WDW        EM
SSB        0
LB         1.00 Hz
GB         0
PC         1.40
  
```

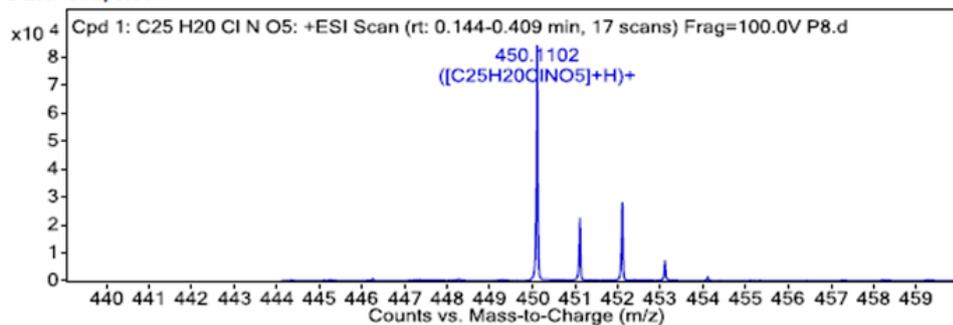
Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C ₂₅ H ₂₀ ClN ₅ O ₅	0.243	449.1028	87486	C ₂₅ H ₂₀ ClN ₅ O ₅	449.103	-0.47

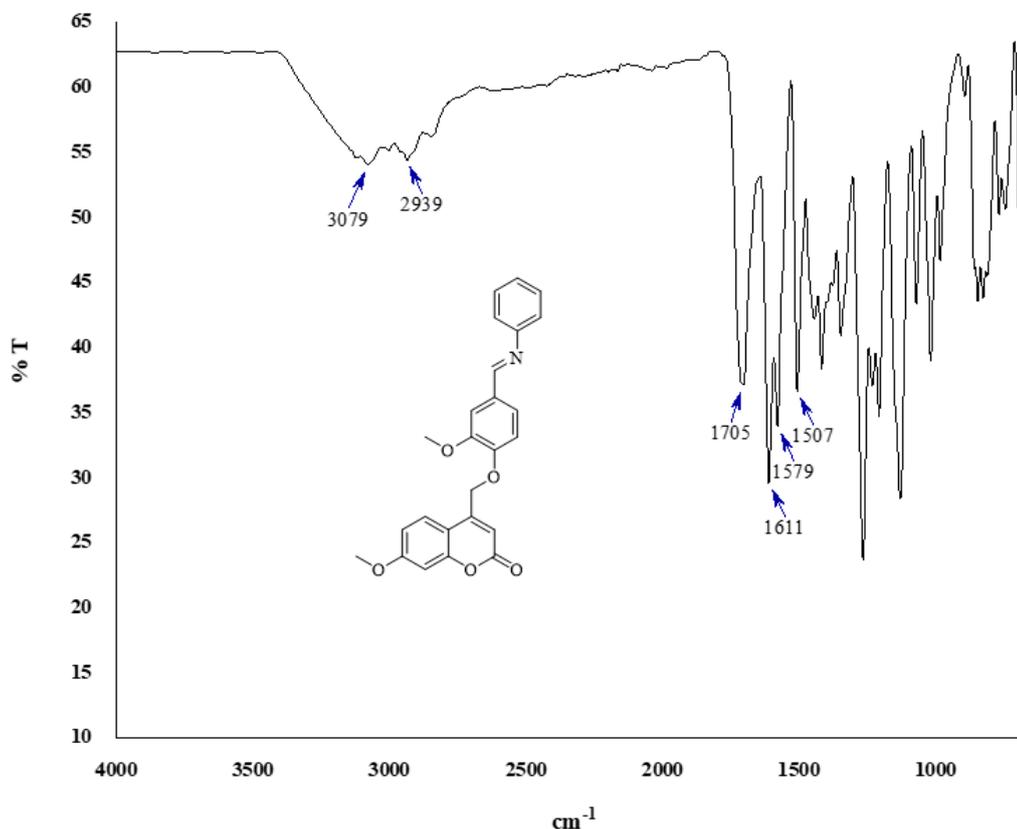
Compound Label	m/z	RT	Algorithm	Mass
Cpd 1: C ₂₅ H ₂₀ ClN ₅ O ₅	450.1102	0.243	Find By Formula	449.1028

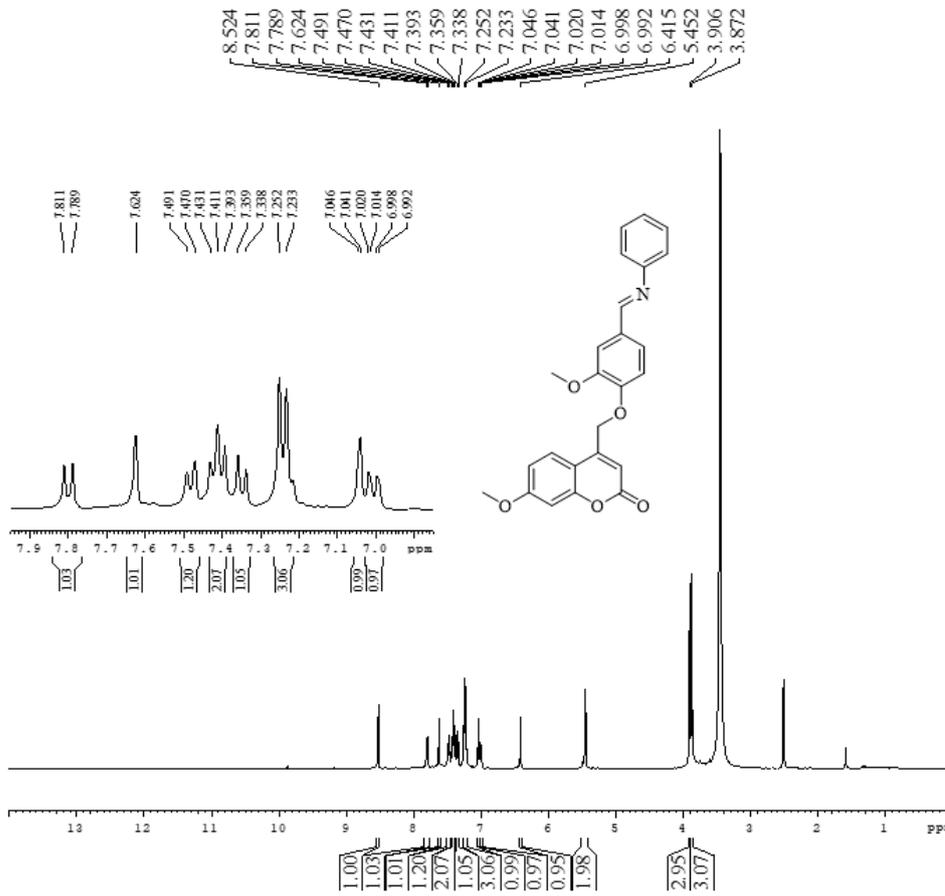


MS Zoomed Spectrum



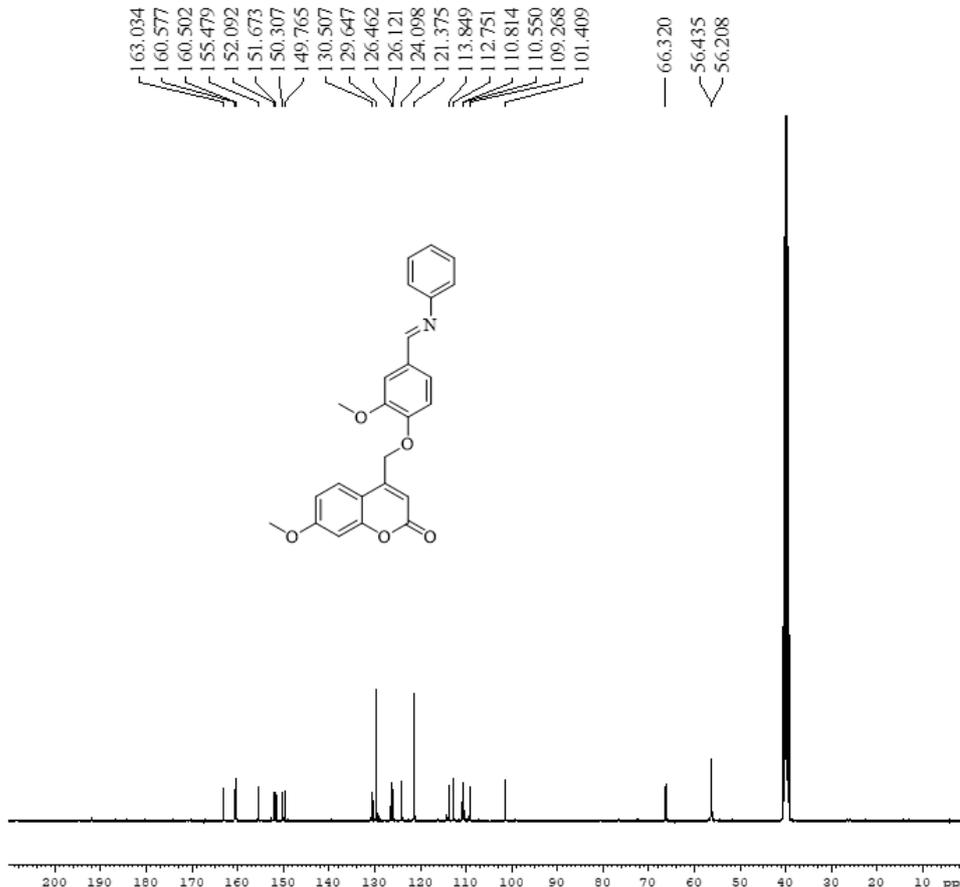
HRMS of (*E*)-4-((4-(((4-chlorophenyl)imino)methyl)-2-methoxy phenoxy)methyl)-7-methoxy-2*H*-chromen-2-one (13d)





```
NAME S970MC-DMSO
EXPNO 1
PROCNO 1
Date_ 20201007
Time 16.43
INSTRUM spect
PROBHD 5 mm FAPBBO BB-
PULPROG zg30
TD 65536
SOLVENT DMSO
NS 16
DS 2
SWH 8223.685 Hz
FIDRES 0.125482 Hz
AQ 3.9846387 sec
RG 144
DW 60.800 usec
DE 6.50 usec
TE 300.0 K
D1 1.00000000 sec
TDO 1
===== CHANNEL f1 =====
NUC1 1H
P1 6.00 usec
PL1 -6.00 dB
SFO1 400.1324710 MHz
SI 32768
SF 400.13200000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00
```

¹H NMR spectrum of (*E*)-7-methoxy-4-((2-methoxy-4-((phenylimino)methyl) phenoxy)methyl)-2*H*-chromen-2-one (13e)



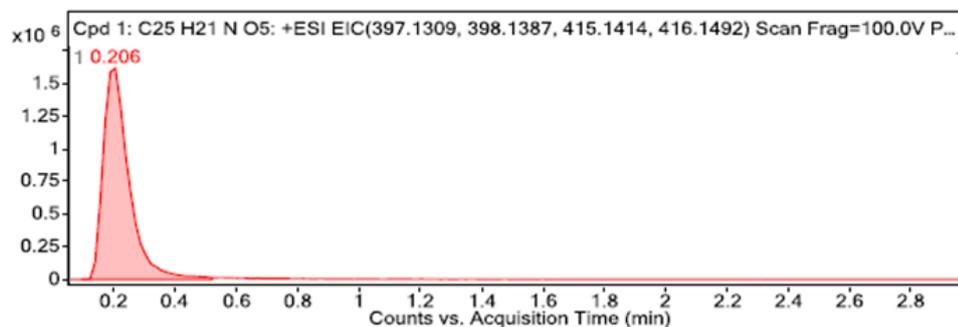
```
NAME S970MC-DMSO
EXPNO 2
PROCNO 1
Date_ 20201008
Time 15.27
INSTRUM spect
PROBHD 5 mm FAPBBO BB-
PULPROG zgpg30
TD 65536
SOLVENT DMSO
NS 20480
DS 4
SWH 24038.461 Hz
FIDRES 0.366798 Hz
AQ 1.3631988 sec
RG 203
DW 20.800 usec
DE 6.50 usec
TE 300.0 K
D1 2.00000000 sec
D11 0.03000000 sec
TDO 1
===== CHANNEL f1 =====
NUC1 13C
P1 12.00 usec
PL1 -1.00 dB
SFO1 100.6228298 MHz
===== CHANNEL f2 =====
CPDPRG2 waltz65
NUC2 1H
PCPD2 80.00 usec
PL2 -6.00 dB
PL12 19.50 dB
PL13 19.50 dB
SFO2 400.1316005 MHz
SI 32768
SF 100.6127690 MHz
WDW EM
SSB 0
LB 1.00 Hz
GB 0
PC 1.40
```

¹³C NMR spectrum of (*E*)-7-methoxy-4-((2-methoxy-4-((phenylimino)methyl) phenoxy)methyl)-2*H*-chromen-2-one (13e)

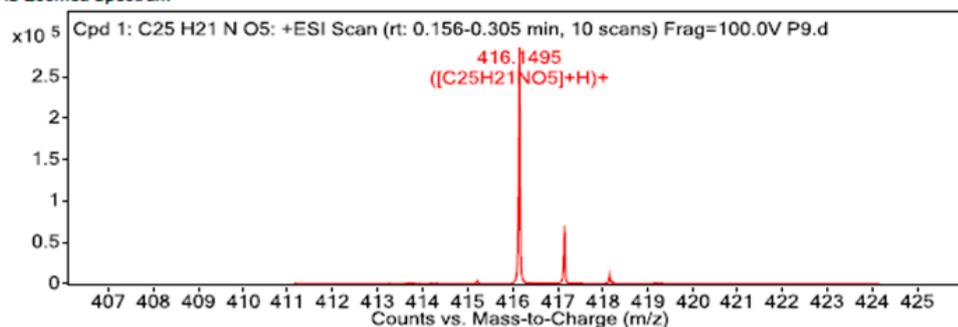
Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C ₂₅ H ₂₁ N O ₅	0.206	415.1421	287628	C ₂₅ H ₂₁ N O ₅	415.142	0.37

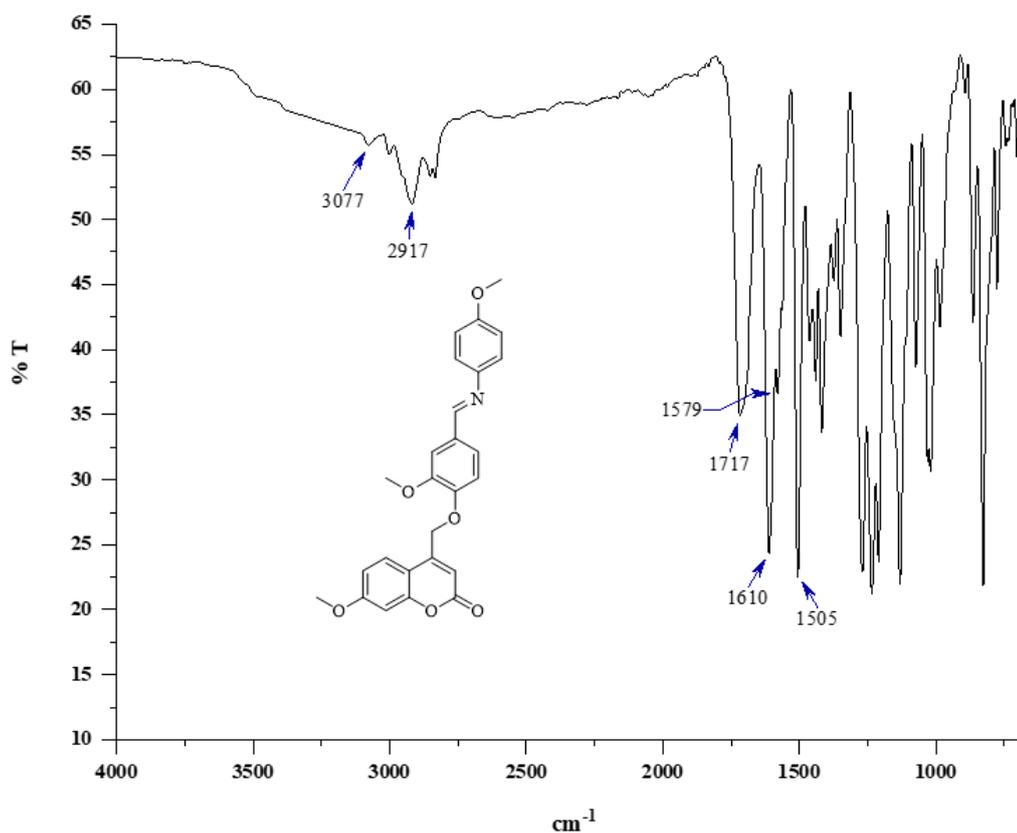
Compound Label	m/z	RT	Algorithm	Mass
Cpd 1: C ₂₅ H ₂₁ N O ₅	416.1495	0.206	Find By Formula	415.1421

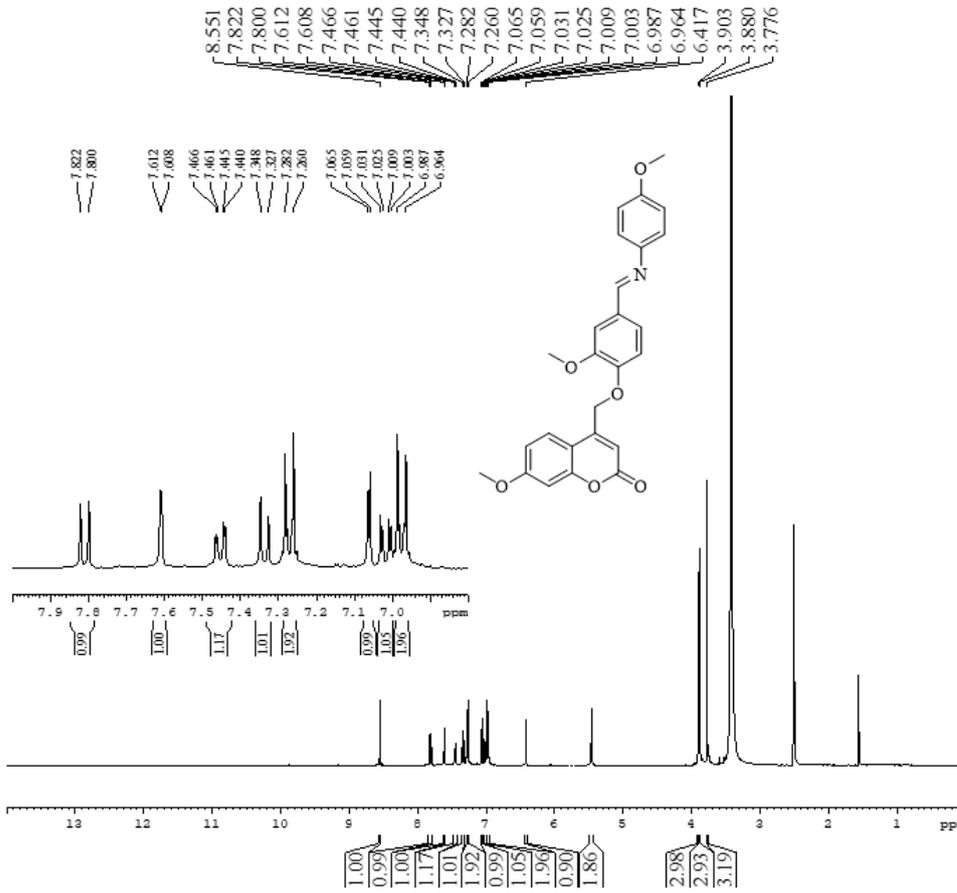


MS Zoomed Spectrum



HRMS of (*E*)-7-methoxy-4-((2-methoxy-4-((4-methoxyphenyl)imino) methyl)phenoxy)methyl)-2*H*-chromen-2-one (**13e**)

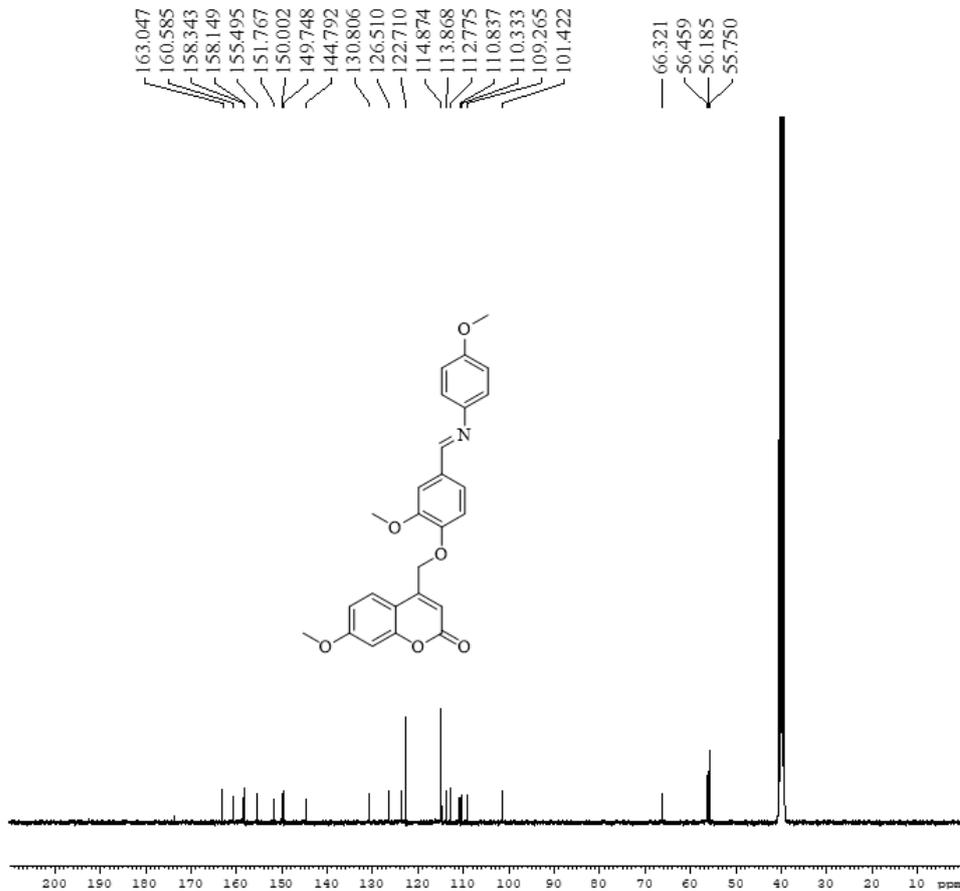




```
NAME      S1070MC-DMSO
EXPNO     3
PROCNO    1
Date_     20201007
Time      16.23
INSTRUM   spect
PROBHD    5 mm PABBO BB-
PULPROG   zg30
TD         65526
SOLVENT   DMSO
NS         16
DS         2
SWH        8223.685 Hz
FIDRES     0.125489 Hz
AQ         3.9846287 sec
RG         203
DW         60.800 usec
DE         6.50 usec
TE         300.0 K
D1         1.00000000 sec
TD0        1
```

```
===== CHANNEL f1 =====
NUC1      1H
P1         6.00 usec
PL1        -6.00 dB
SFO1      400.1324710 MHz
SI         32768
SF         400.1300000 MHz
WDW        EM
SSB         0
LB         0.30 Hz
GB         0
PC         1.00
```

¹H NMR spectrum of (*E*)-7-methoxy-4-((2-methoxy-4-(((4-methoxyphenyl)imino) methyl)phenoxy)methyl)-2H-chromen-2-one (**13f**)



```
NAME      S1070MC-DMSO
EXPNO     3
PROCNO    1
Date_     20201007
Time      17.15
INSTRUM   spect
PROBHD    5 mm PABBO BB-
PULPROG   zgpg30
TD         65526
SOLVENT   DMSO
NS         16284
DS         4
SWH        24038.461 Hz
FIDRES     0.366798 Hz
AQ         1.3631988 sec
RG         203
DW         20.800 usec
DE         6.50 usec
TE         300.0 K
D1         2.00000000 sec
D11        0.03000000 sec
TD0        1
```

```
===== CHANNEL f1 =====
NUC1      13C
P1         12.00 usec
PL1        -1.00 dB
SFO1      100.6228298 MHz
```

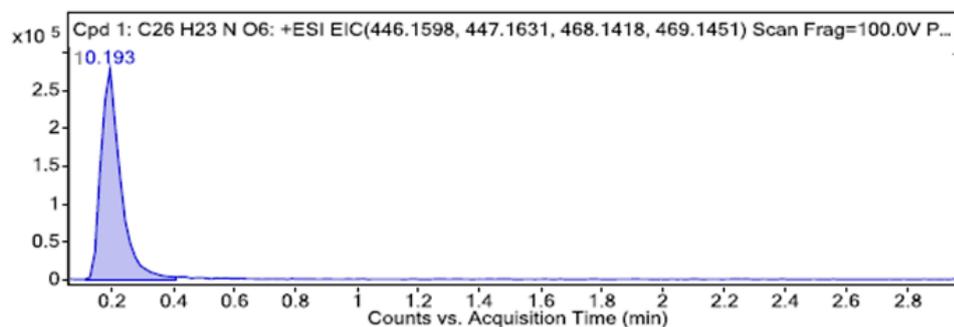
```
===== CHANNEL f2 =====
CPDPRG2   waltz65
NUC2      1H
PCPD2     80.00 usec
PL2        -6.00 dB
PL12       19.50 dB
PL13       19.50 dB
SFO2      400.1316005 MHz
SI         32768
SF         100.6127690 MHz
WDW        EM
SSB         0
LB         1.00 Hz
GB         0
PC         1.40
```

¹³C NMR spectrum of (*E*)-7-methoxy-4-((2-methoxy-4-(((4-methoxyphenyl)imino) methyl)phenoxy)methyl)-2H-chromen-2-one (**13f**)

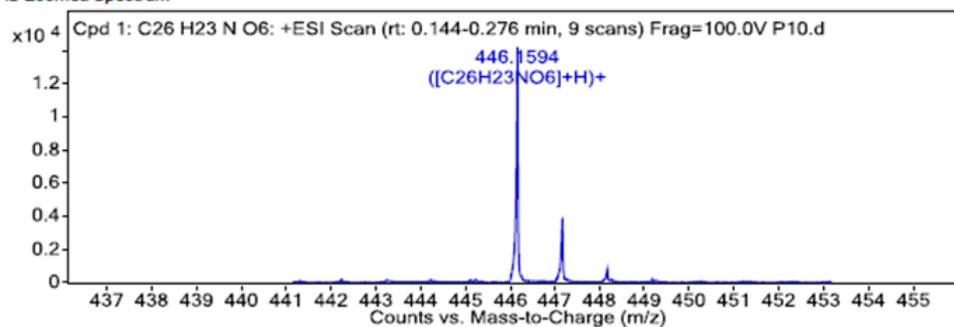
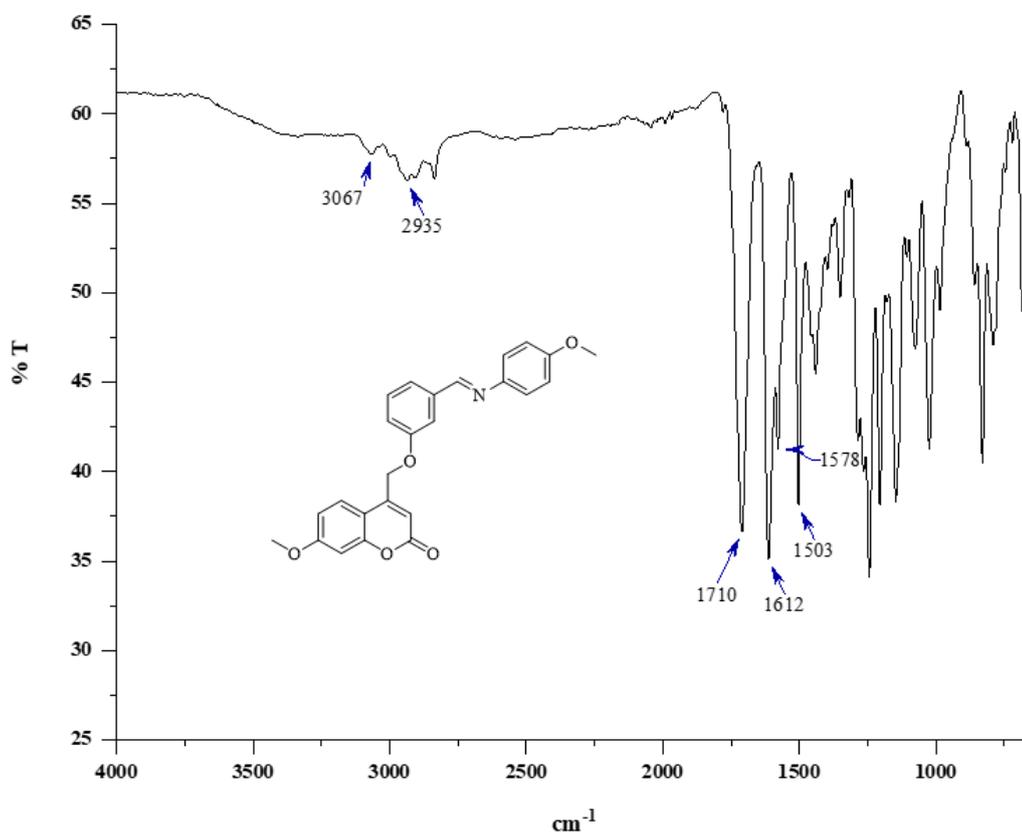
Compound Table

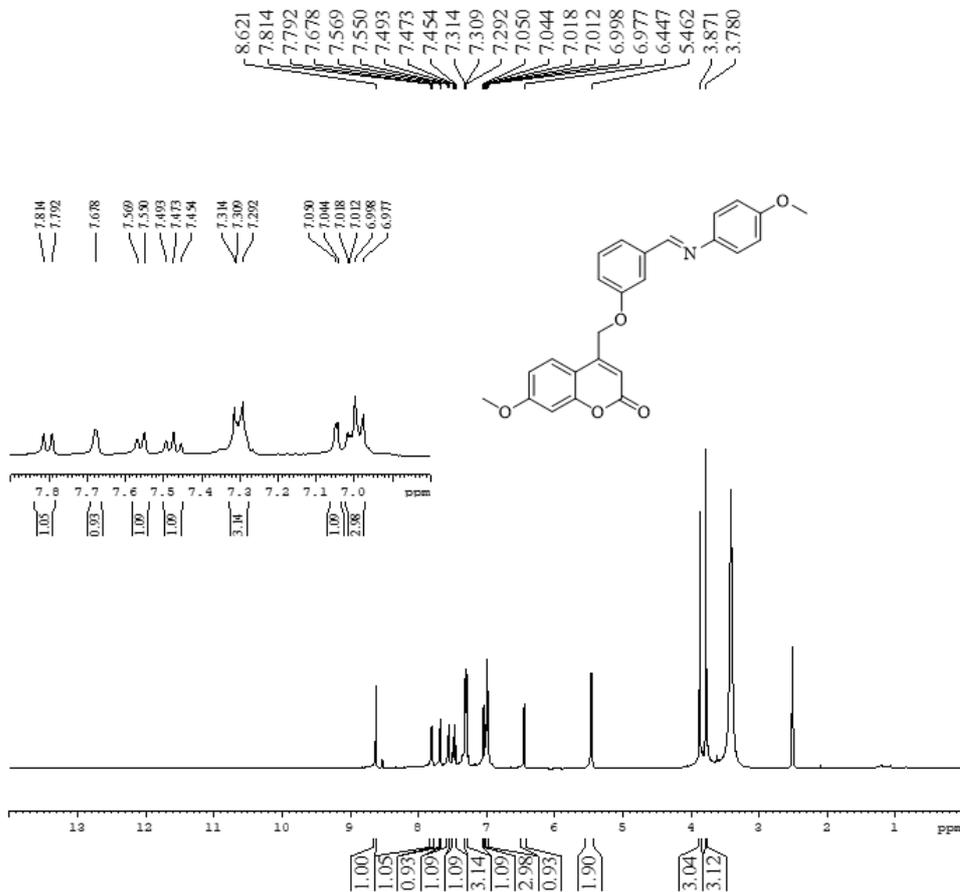
Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C ₂₆ H ₂₃ N O ₆	0.193	445.1519	14309	C ₂₆ H ₂₃ N O ₆	445.1525	-1.53

Compound Label	<i>m/z</i>	RT	Algorithm	Mass
Cpd 1: C ₂₆ H ₂₃ N O ₆	446.1594	0.193	Find By Formula	445.1519



MS Zoomed Spectrum

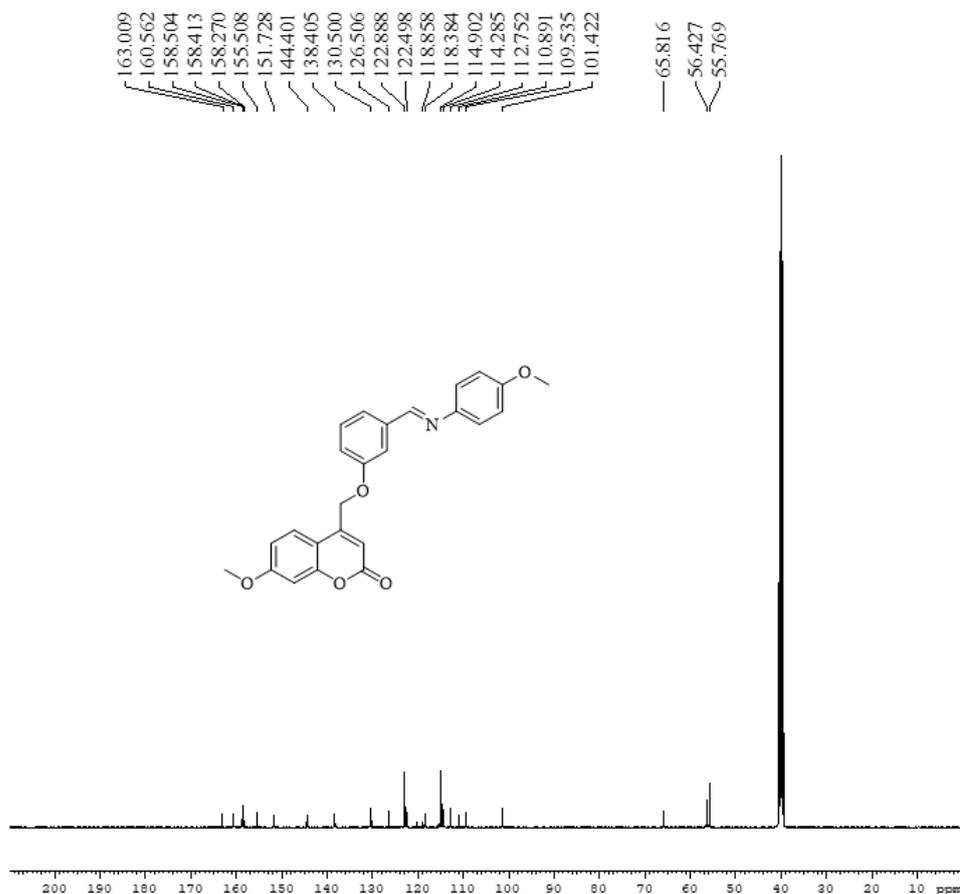
HRMS of (*E*)-7-methoxy-4-((2-methoxy-4-(((4-methoxyphenyl)imino) methyl)phenoxy)methyl)-2*H*-chromen-2-one (13f)IR spectrum of (*E*)-7-methoxy-4-((2-methoxy-4-(((4-methoxyphenyl)imino) methyl) phenoxy)methyl)-2*H*-chromen-2-one (13g)



```
NAME S1970MC-DMSO
EXPNO 1
PROCNO 1
Date_ 20201008
Time 15.04
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT DMSO
NS 16
DS 4
SWH 8223.665 Hz
FIDRES 0.125482 Hz
AQ 3.9846387 sec
RG 128
DW 60.800 usec
DE 6.50 usec
TE 300.0 K
D1 1.00000000 sec
TD0 1
```

```
===== CHANNEL f1 =====
NUC1 1H
P1 6.00 usec
PL1 -6.00 dB
SFO1 400.1324710 MHz
SI 32768
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00
```

¹H NMR spectrum of *(E)*-7-methoxy-4-((3-(((4-methoxyphenyl)imino)methyl) phenoxy)methyl)-2*H*-chromen-2-one (13g)



```
NAME S1970MC-DMSO
EXPNO 2
PROCNO 1
Date_ 20201011
Time 16.58
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zgpg30
TD 65536
SOLVENT DMSO
NS 18432
DS 4
SWH 24038.461 Hz
FIDRES 0.366798 Hz
AQ 1.3621988 sec
RG 203
DW 20.800 usec
DE 6.50 usec
TE 300.0 K
D1 2.00000000 sec
D11 0.03000000 sec
TD0 1
```

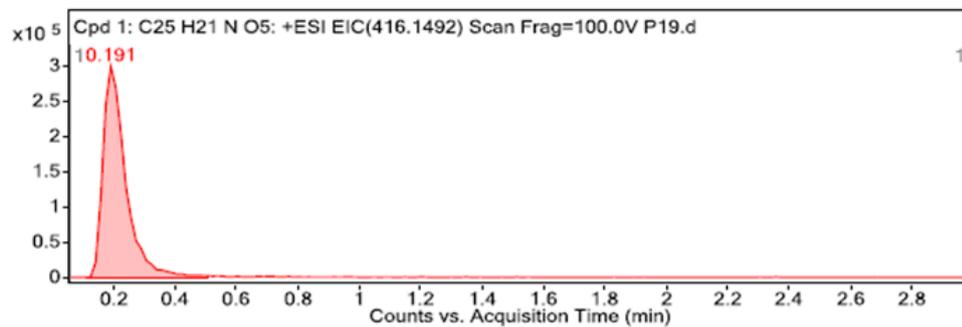
```
===== CHANNEL f1 =====
NUC1 13C
P1 12.00 usec
PL1 -1.00 dB
SFO1 100.6228298 MHz
===== CHANNEL f2 =====
CPDPRG2 waltz65
NUC2 1H
PCPD2 80.00 usec
PL2 -6.00 dB
PL12 19.50 dB
PL13 19.50 dB
SFO2 400.1316005 MHz
SI 32768
SF 100.6127690 MHz
WDW EM
SSB 0
LB 1.00 Hz
GB 0
PC 1.40
```

¹³C NMR spectrum of *(E)*-7-methoxy-4-((3-(((4-methoxyphenyl)imino)methyl) phenoxy)methyl)-2*H*-chromen-2-one (13g)

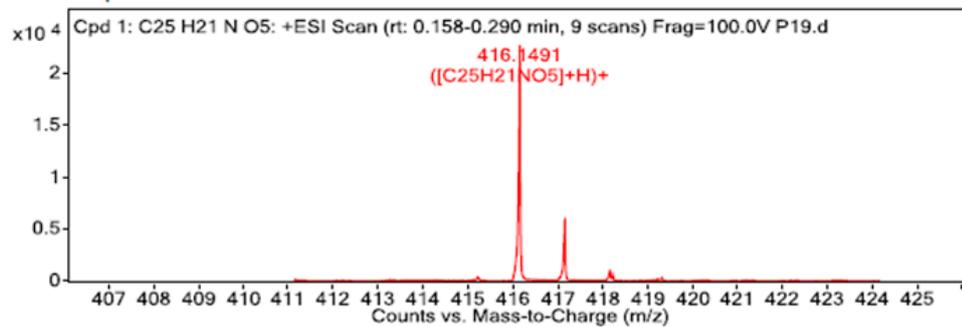
Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C ₂₅ H ₂₁ N O ₅	0.191	415.1417	22797	C ₂₅ H ₂₁ N O ₅	415.142	-0.69

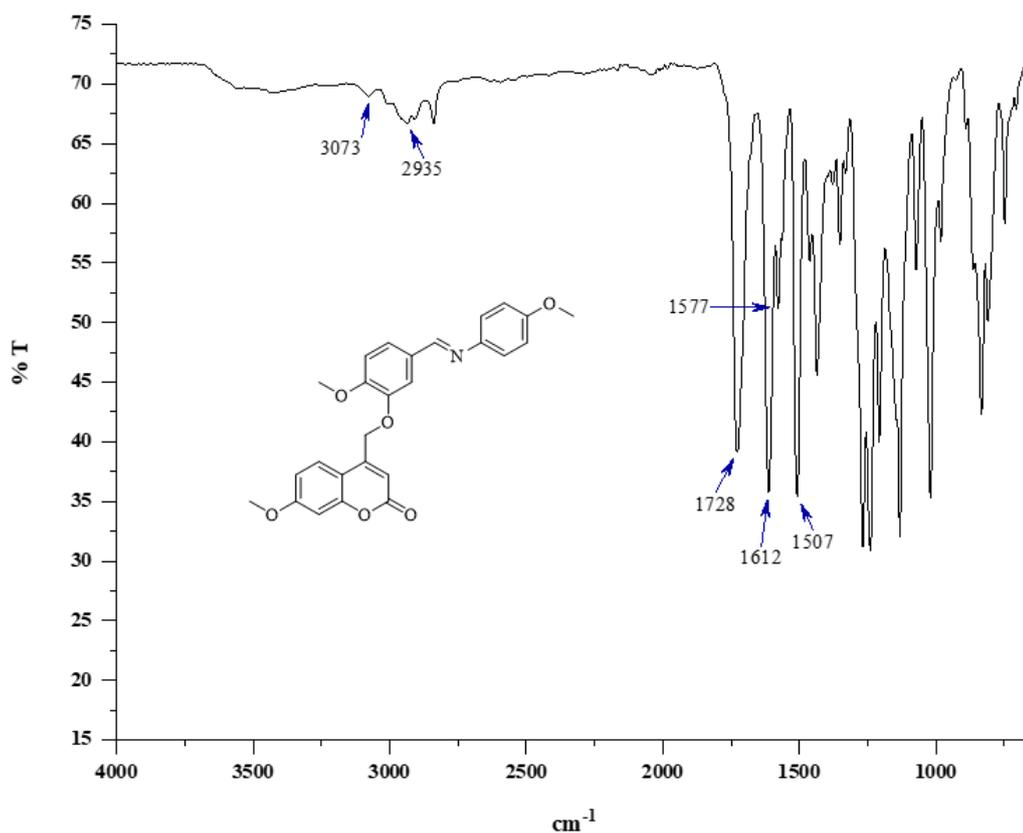
Compound Label	m/z	RT	Algorithm	Mass
Cpd 1: C ₂₅ H ₂₁ N O ₅	416.1491	0.191	Find By Formula	415.1417

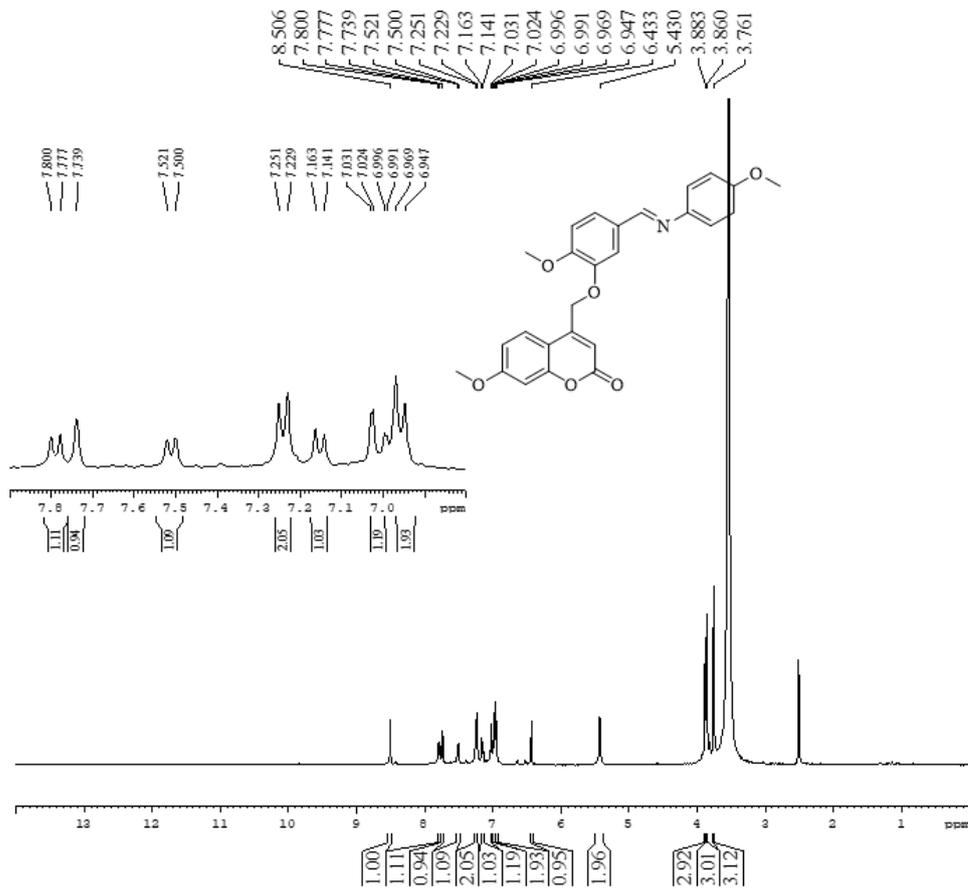


MS Zoomed Spectrum



HRMS of (*E*)-7-methoxy-4-((3-(((4-methoxyphenyl)imino)methyl)phenoxy)methyl)-2*H*-chromen-2-one (**13g**)





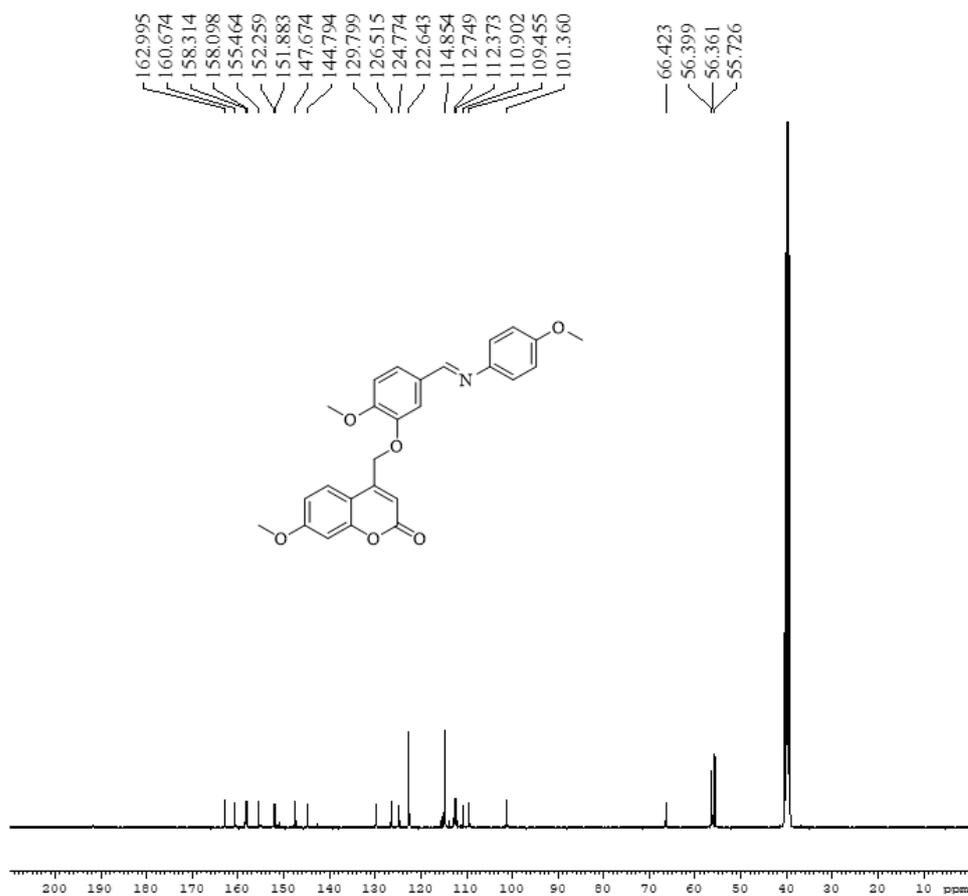
```

NAME      S2070MC-DMSO
EXPNO     3
PROCNO    1
Date_     20201011
Time      16.08
INSTRUM   spect
PROBHD    5 mm PABBO BB-
PULPROG   zg30
TD         65536
SOLVENT   DMSO
NS         100
DS         2
SWH        8223.688 Hz
FIDRES     0.125482 Hz
AQ         3.9846387 sec
RG          90.5
DW         60.800 usec
DE         6.50 usec
TE         300.0 K
D1         1.00000000 sec
TD0        1

===== CHANNEL f1 =====
NUC1      1H
P1         6.00 usec
PL1        -6.00 dB
SFO1      400.1324710 MHz
SI         32768
SF         400.1300000 MHz
WDW        EM
SSB         0
LB         0.30 Hz
GB         0
PC         1.00

```

¹H NMR spectrum of (*E*)-7-methoxy-4-((2-methoxy-5-(((4-methoxyphenyl)imino) methyl)phenoxy)methyl)-2H-chromen-2-one (**13h**)



```

NAME      S2070MC-DMSO
EXPNO     2
PROCNO    1
Date_     20201015
Time      14.49
INSTRUM   spect
PROBHD    5 mm PABBO BB-
PULPROG   zgpg30
TD         65536
SOLVENT   DMSO
NS         25600
DS         4
SWH        24038.461 Hz
FIDRES     0.366798 Hz
AQ         1.3621988 sec
RG          203
DW         20.800 usec
DE         6.50 usec
TE         300.0 K
D1         2.00000000 sec
D11        0.03000000 sec
TD0        1

===== CHANNEL f1 =====
NUC1      13C
P1         12.00 usec
PL1        -1.00 dB
SFO1      100.6228298 MHz

===== CHANNEL f2 =====
CFDPRG2   waltz16
NUC2      1H
PCPD2     80.00 usec
PL2        -6.00 dB
PL12       19.50 dB
PL13       19.50 dB
SFO2      400.1316005 MHz
SI         32768
SF         100.6127650 MHz
WDW        EM
SSB         0
LB         1.00 Hz
GB         0
PC         1.40

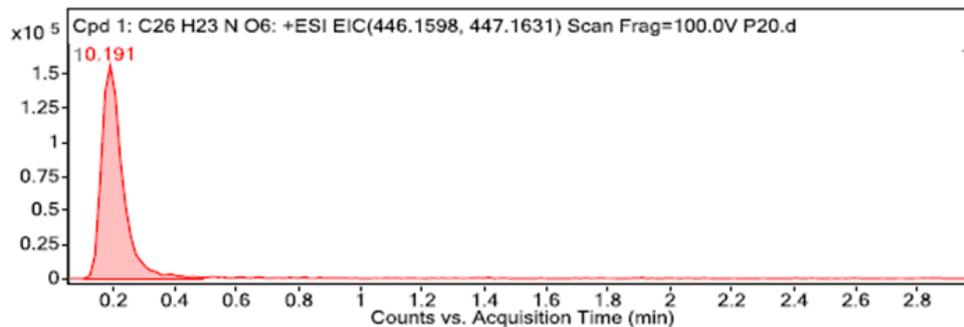
```

¹³C NMR spectrum of (*E*)-7-methoxy-4-((2-methoxy-5-(((4-methoxyphenyl)imino) methyl)phenoxy)methyl)-2H-chromen-2-one (**13h**)

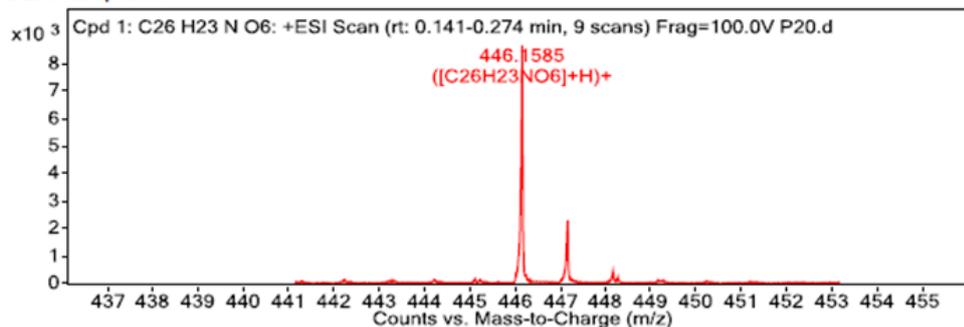
Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C26 H23 N O6	0.191	445.1512	8662	C26 H23 N O6	445.1525	-3

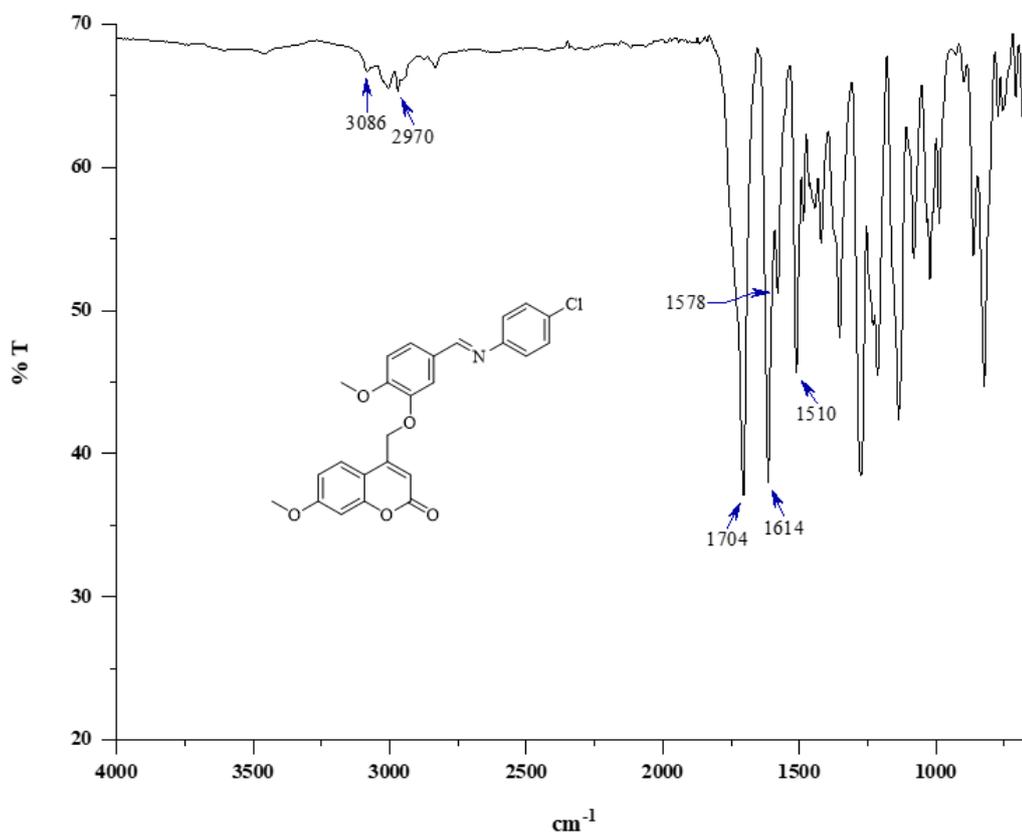
Compound Label	m/z	RT	Algorithm	Mass
Cpd 1: C26 H23 N O6	446.1585	0.191	Find By Formula	445.1512



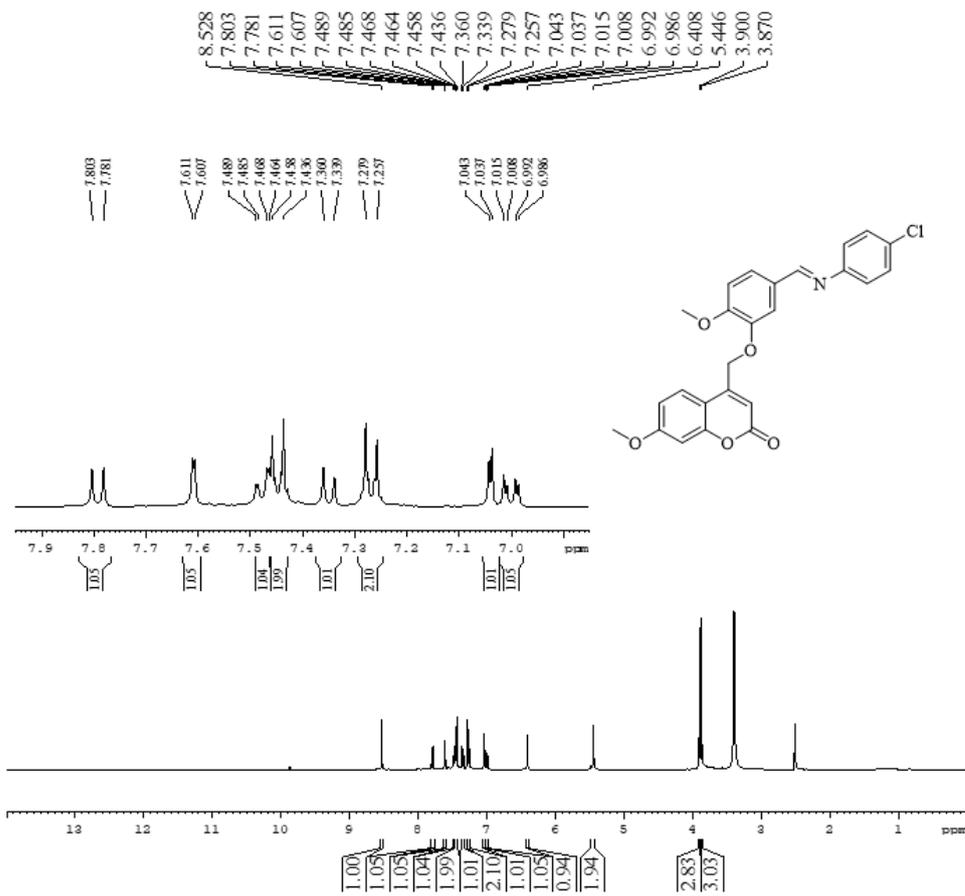
MS Zoomed Spectrum



HRMS of (*E*)-7-methoxy-4-((2-methoxy-5-(((4-methoxyphenyl)imino) methyl)phenoxy)methyl)-2*H*-chromen-2-one (13h)



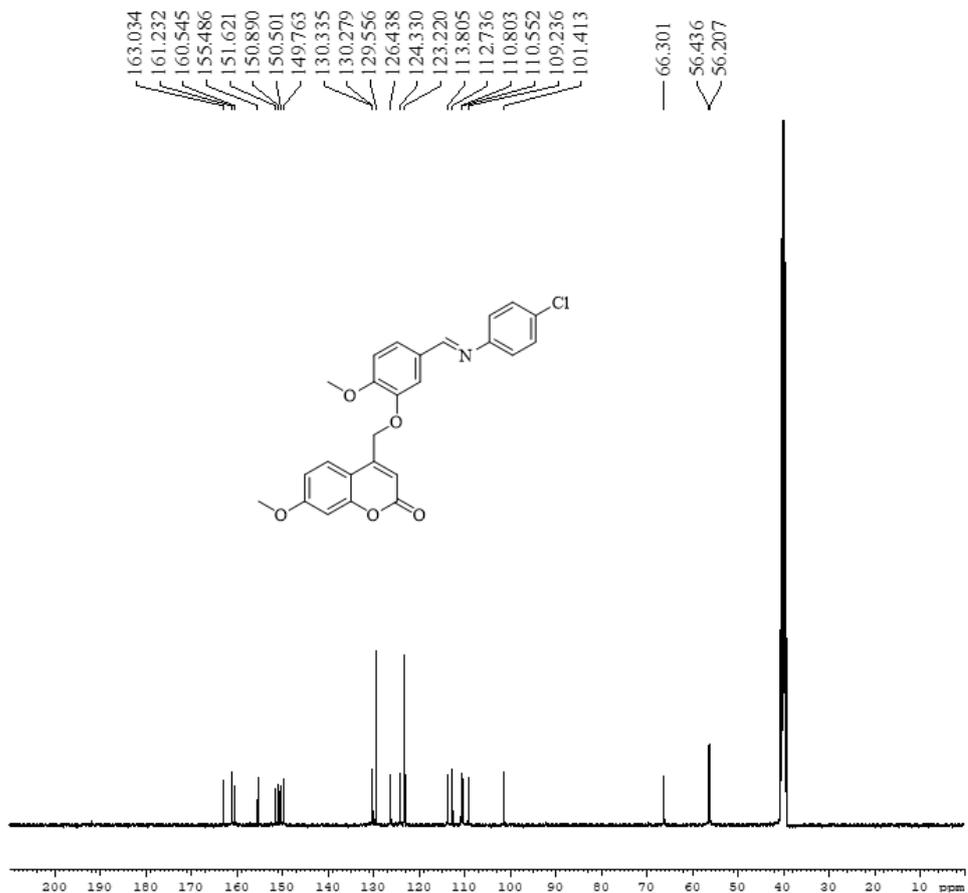
IR spectrum of (*E*)-4-((5-(((4-chlorophenyl)imino)methyl)-2-methoxy phenoxy)methyl)-7-methoxy-2*H*-chromen-2-one (13i)



```
NAME      S2370MC-DMSO
EXPNO     2
PROCNO    1
Date_     20201004
Time      16.27
INSTRUM   spect
PROBHD    5 mm FAPBO BB-
PULPROG   zg30
TD         65526
SOLVENT   DMSO
NS         16
DS         2
SWH        8222.685 Hz
FIDRES     0.125483 Hz
AQ         3.9846387 sec
RG         144
DW         60.800 usec
DE         6.50 usec
TE         300.0 K
D1         1.00000000 sec
TD0        1
```

```
===== CHANNEL f1 =====
NUC1      1H
P1         6.00 usec
PL1        -6.00 dB
SFO1      400.1324710 MHz
SI         32768
SF         400.1300000 MHz
WDW        EM
SSB        0
LB         0.20 Hz
GB         0
PC         1.00
```

¹H NMR spectrum of (*E*)-4-((5-(((4-chlorophenyl)imino)methyl)-2-methoxy phenoxy)methyl)-7-methoxy-2H-chromen-2-one (**13i**)



```
NAME      S2370MC-DMSO
EXPNO     1
PROCNO    1
Date_     20201004
Time      17.11
INSTRUM   spect
PROBHD    5 mm FAPBO BB-
PULPROG   zgpg30
TD         65526
SOLVENT   DMSO
NS         15872
DS         4
SWH        24028.461 Hz
FIDRES     0.366798 Hz
AQ         1.3631988 sec
RG         202
DW         20.800 usec
DE         6.50 usec
TE         300.0 K
D1         2.00000000 sec
D11        0.02000000 sec
TD0        1
```

```
===== CHANNEL f1 =====
NUC1      13C
P1         12.00 usec
PL1        -1.00 dB
SFO1      100.6228298 MHz
```

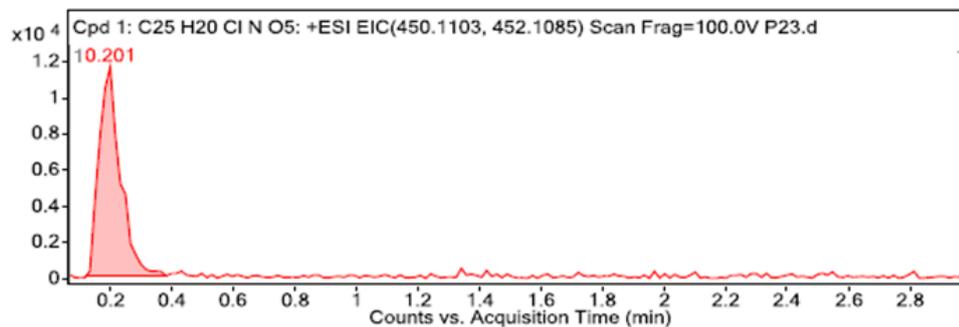
```
===== CHANNEL f2 =====
CPDPRG2   waltz65
NUC2      1H
PCPD2     80.00 usec
PL2        -6.00 dB
PL12      19.50 dB
PL13      19.50 dB
SFO2      400.1316005 MHz
SI         32768
SF         100.6127690 MHz
WDW        EM
SSB        0
LB         1.00 Hz
GB         0
PC         1.40
```

¹³C NMR spectrum of (*E*)-4-((5-(((4-chlorophenyl)imino)methyl)-2-methoxy phenoxy)methyl)-7-methoxy-2H-chromen-2-one (**13i**)

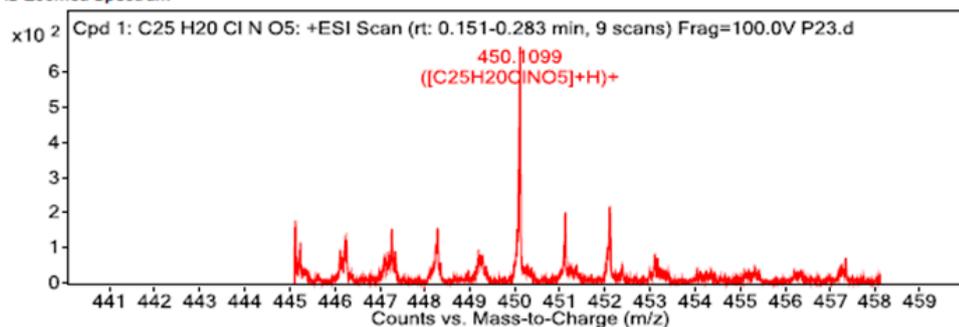
Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C25 H20 Cl N O5	0.201	449.1029	691	C25 H20 Cl N O5	449.103	-0.19

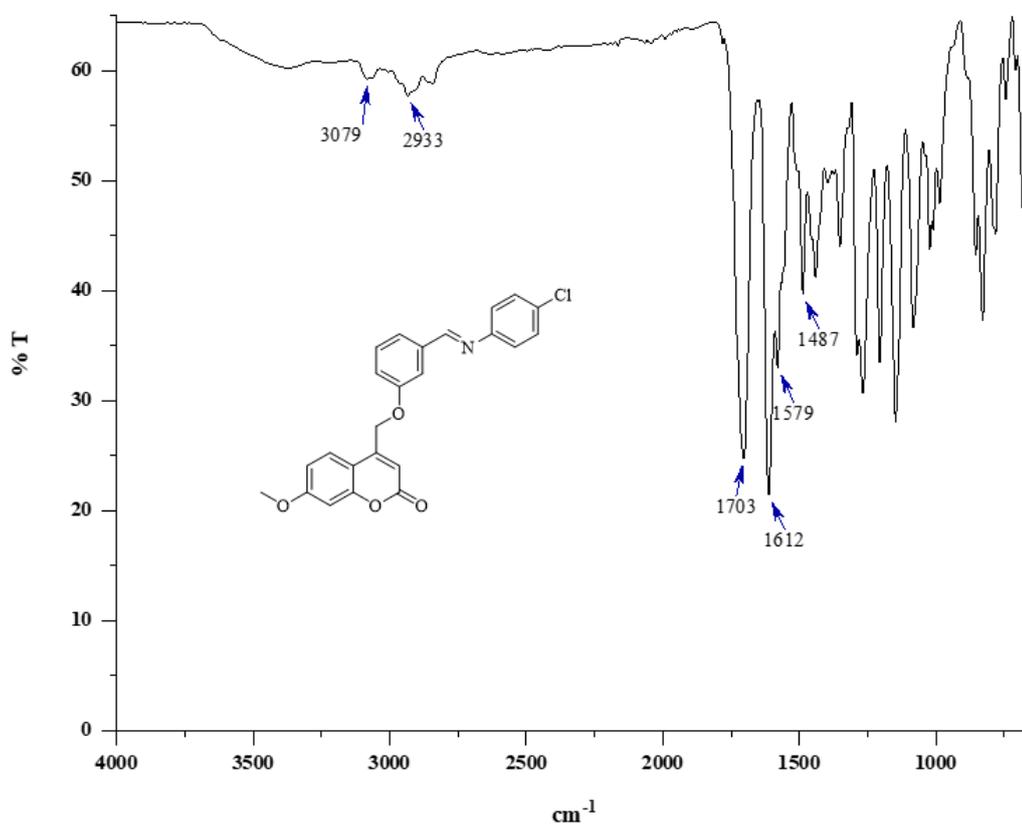
Compound Label	m/z	RT	Algorithm	Mass
Cpd 1: C25 H20 Cl N O5	450.1099	0.201	Find By Formula	449.1029

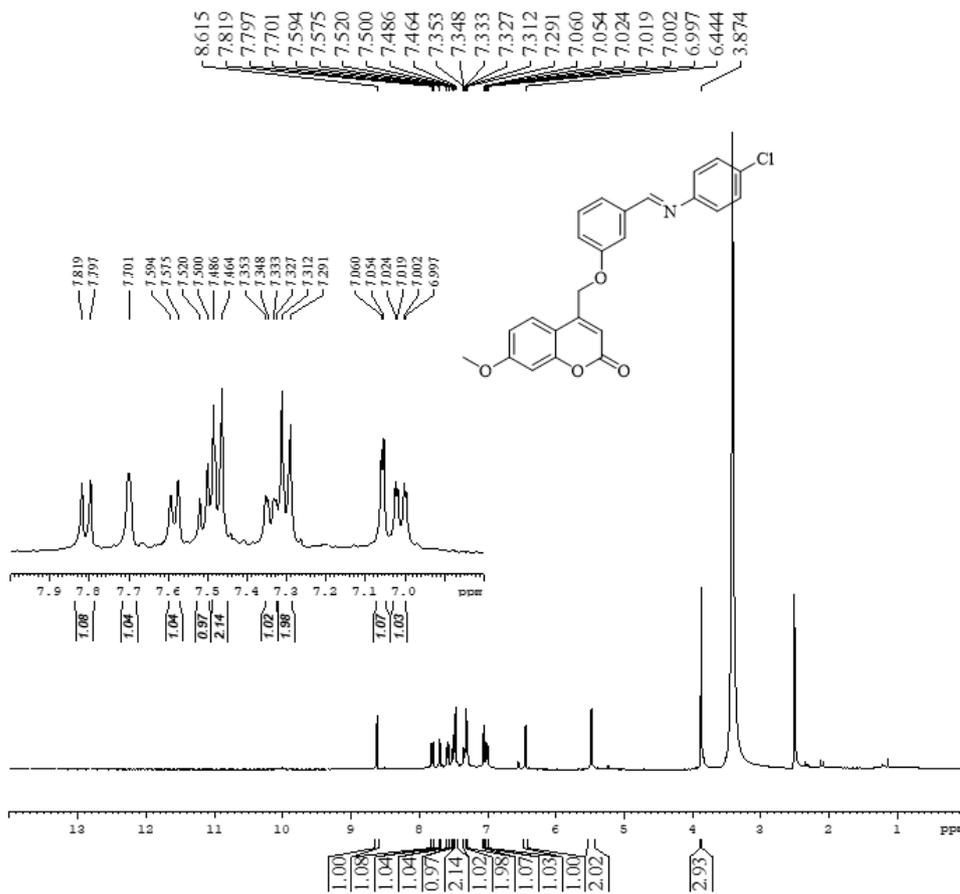


MS Zoomed Spectrum



HRMS of (*E*)-4-((5-(((4-chlorophenyl)imino)methyl)-2-methoxy phenoxy)methyl)-7-methoxy-2*H*-chromen-2-one (13i)





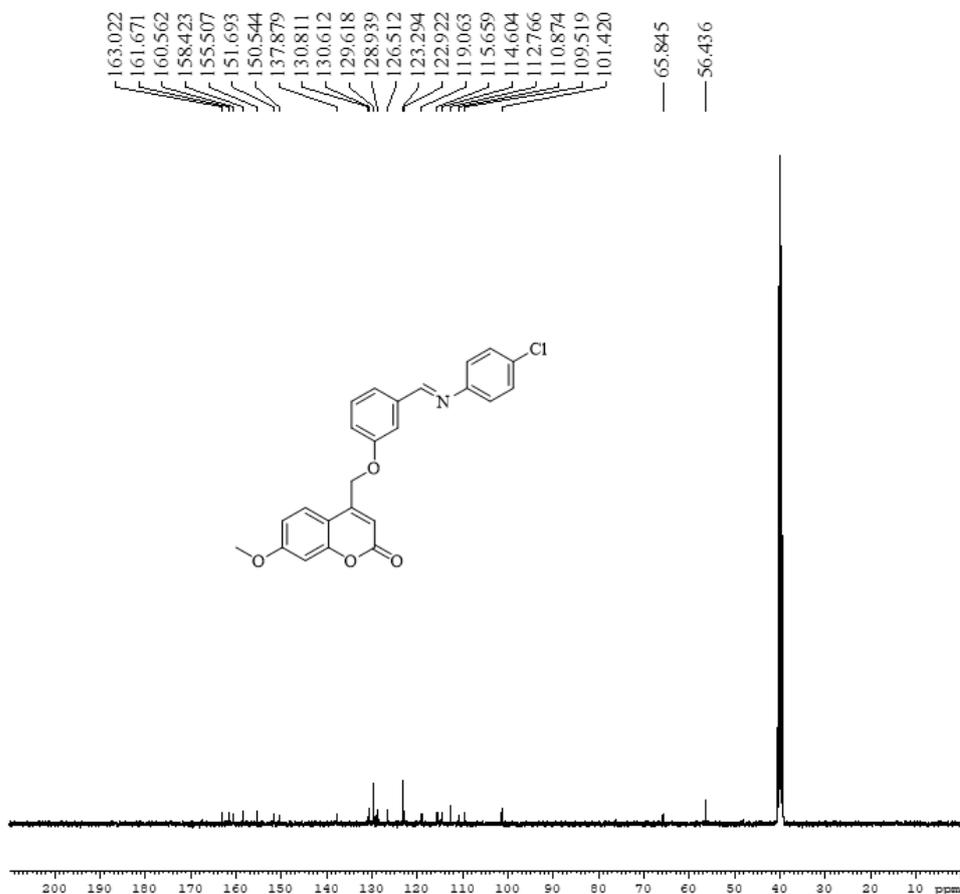
BRUKER

```

NAME      S2470MC-DMSO
EXPNO     1
PROCNO    1
Date_     20201018
Time      11.12
INSTRUM   spect
PROBHD    5 mm PABBO BB-
PULPROG   sg30
TD         65526
SOLVENT   DMSO
NS         16
DS         2
SWH        8222.688 Hz
FIDRES     0.125489 Hz
AQ         3.9846387 sec
RG         161
DW         60.800 usec
DE         6.50 usec
TE         300.0 K
D1         1.00000000 sec
TD0        1

===== CHANNEL f1 =====
NUC1      1H
P1         6.00 usec
PL1        -6.00 dB
SFO1      400.1324710 MHz
SI         32768
SF         400.1300000 MHz
WDW        EM
SSB        0
LB         0.30 Hz
GB         0
PC         1.00
  
```

¹H NMR spectrum of (E)-4-((3-(((4-chlorophenyl)imino)methyl)phenoxy)methyl)-7-methoxy-2H-chromen-2-one (13j)



BRUKER

```

NAME      S2470MC-DMSO
EXPNO     5
PROCNO    1
Date_     20201018
Time      16.23
INSTRUM   spect
PROBHD    5 mm PABBO BB-
PULPROG   waltz16
TD         65526
SOLVENT   DMSO
NS         1056
DS         4
SWH        24038.461 Hz
FIDRES     0.366798 Hz
AQ         1.3631988 sec
RG         203
DW         20.800 usec
DE         6.50 usec
TE         300.0 K
D1         2.00000000 sec
D11        0.03000000 sec
TD0        1

===== CHANNEL f1 =====
NUC1      13C
P1         12.00 usec
PL1        -1.00 dB
SFO1      100.6228298 MHz

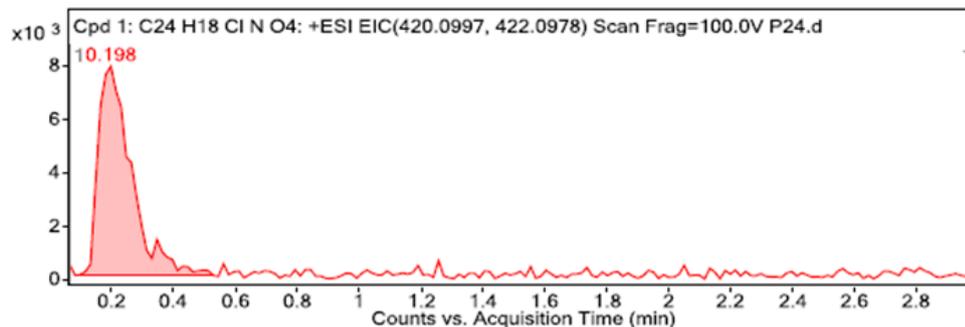
===== CHANNEL f2 =====
CPDPRG2   waltz16
NUC2      1H
PCPD2     80.00 usec
PL2        -6.00 dB
PL12       19.50 dB
PL13       19.50 dB
SFO2      400.1316005 MHz
SI         32768
SF         100.6127690 MHz
WDW        EM
SSB        0
LB         1.00 Hz
GB         0
PC         1.40
  
```

¹³C NMR spectrum of (E)-4-((3-(((4-chlorophenyl)imino)methyl)phenoxy)methyl)-7-methoxy-2H-chromen-2-one (13j)

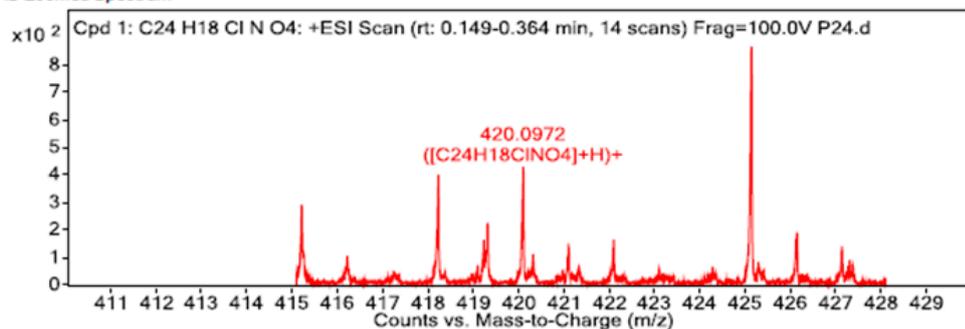
Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C24 H18 Cl N O4	0.198	419.0908	429	C24 H18 Cl N O4	419.0924	-3.79

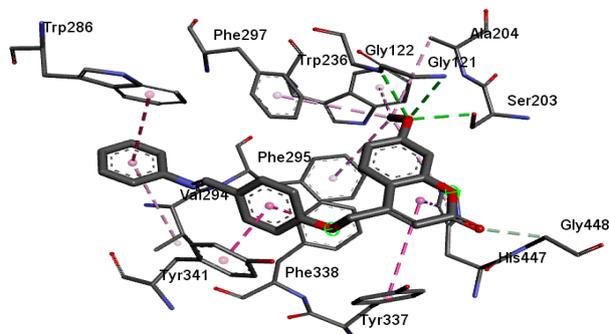
Compound Label	m/z	RT	Algorithm	Mass
Cpd 1: C24 H18 Cl N O4	420.0972	0.198	Find By Formula	419.0908



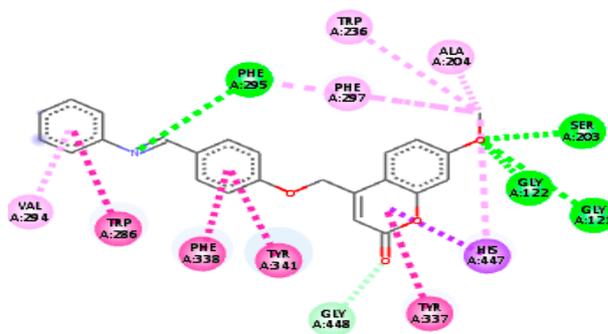
MS Zoomed Spectrum



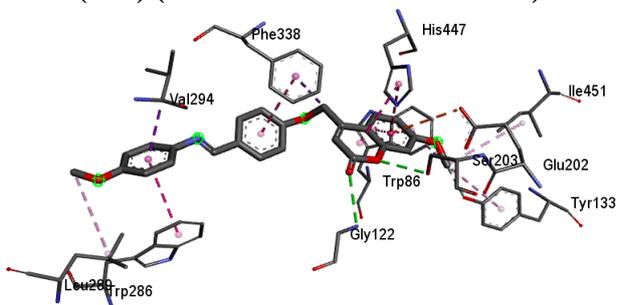
HRMS of (*E*)-4-((3-(((4-chlorophenyl)imino)methyl)phenoxy)methyl)-7-methoxy-2*H*-chromen-2-one (13j)



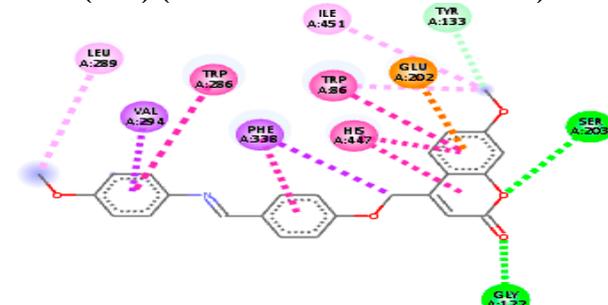
(13a) (3D LIGAND-RECEPTOR)



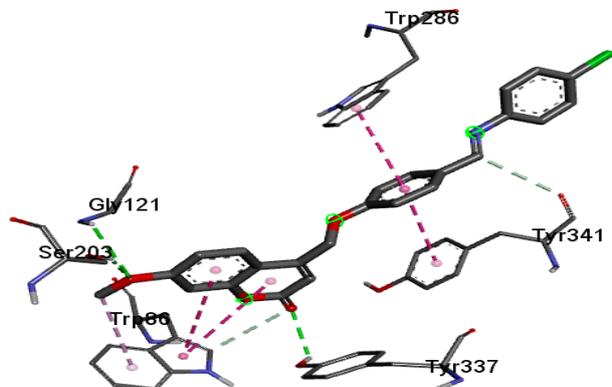
(13a) (2D LIGAND-RECEPTOR)



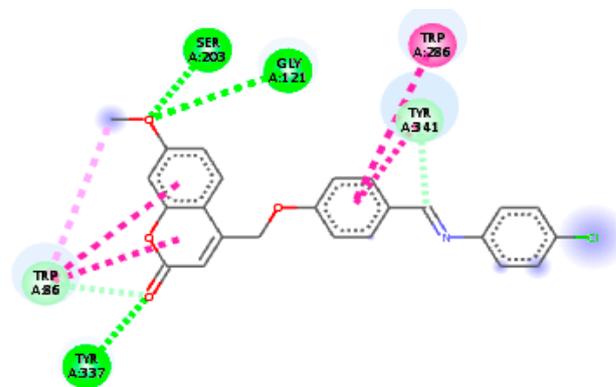
(13b) (3D LIGAND-RECEPTOR)



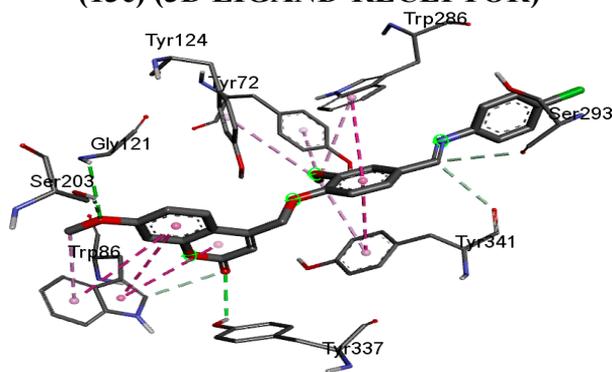
(13b) (2D LIGAND-RECEPTOR)



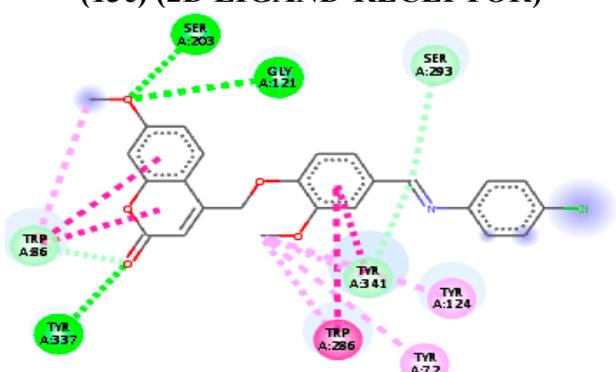
(13c) (3D LIGAND-RECEPTOR)



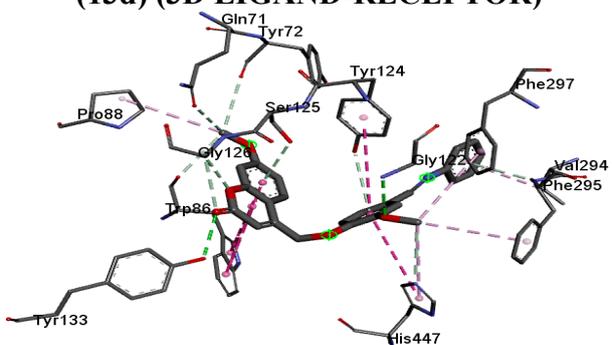
(13c) (2D LIGAND-RECEPTOR)



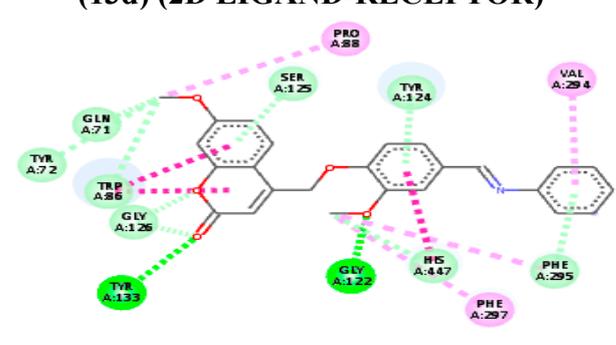
(13d) (3D LIGAND-RECEPTOR)



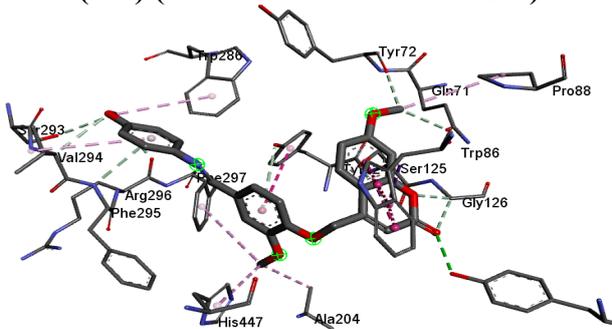
(13d) (2D LIGAND-RECEPTOR)



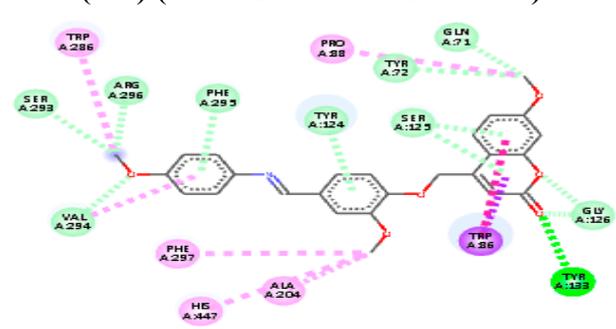
(13e) (3D LIGAND-RECEPTOR)



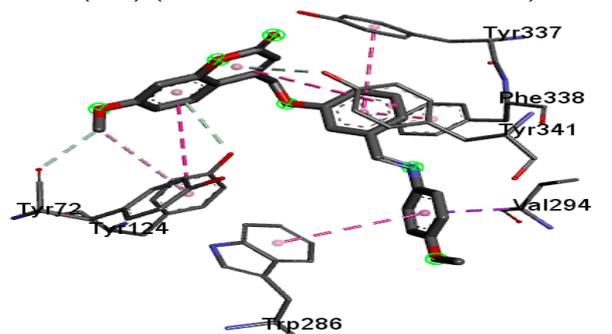
(13e) (2D LIGAND-RECEPTOR)



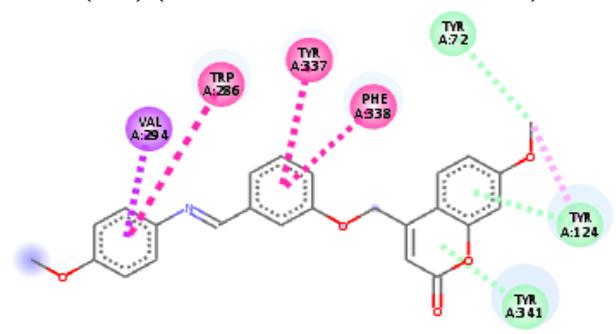
(13f) (3D LIGAND-RECEPTOR)



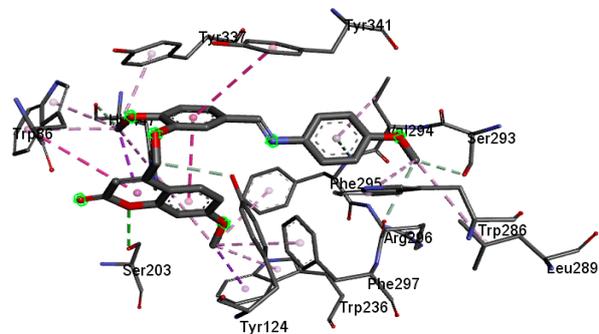
(13f) (2D LIGAND-RECEPTOR)



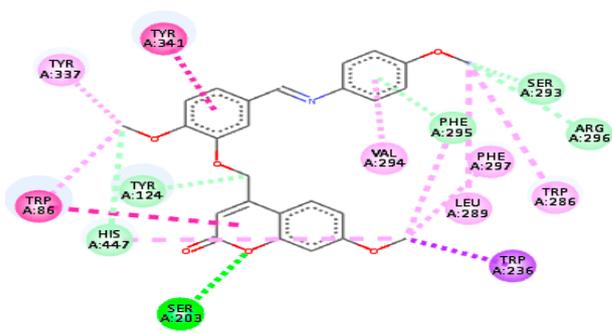
(13g) (3D LIGAND-RECEPTOR)



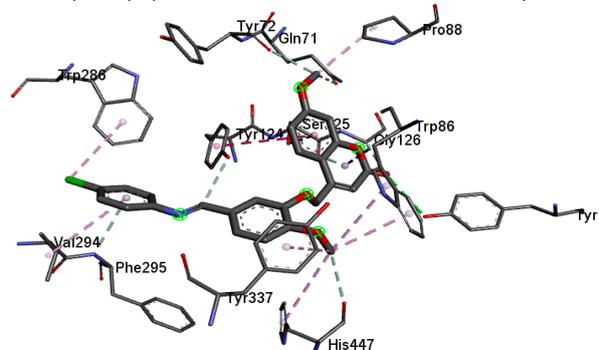
(13g) (2D LIGAND-RECEPTOR)



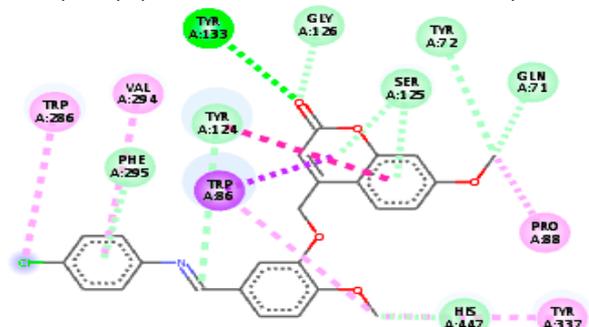
(13h) (3D LIGAND-RECEPTOR)



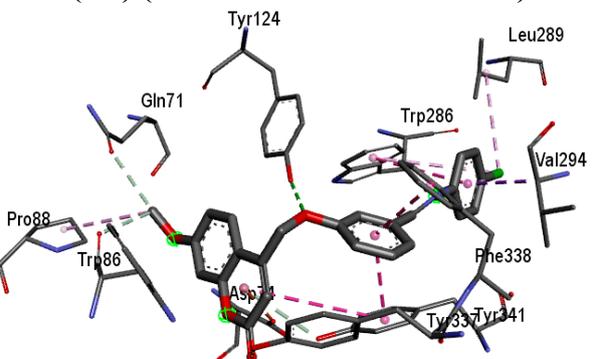
(13h) (2D LIGAND-RECEPTOR)



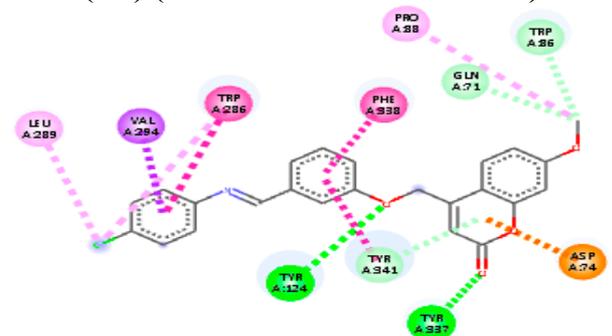
(13i) (3D LIGAND-RECEPTOR)



(13i) (2D LIGAND-RECEPTOR)



(13j) (3D LIGAND-RECEPTOR)



(13j) (2D LIGAND-RECEPTOR)

Interactions

	Conventional Hydrogen Bond
	Carbon Hydrogen Bond
	Pi-Donor Hydrogen Bond
	Pi-Anion
	Pi-Donor Hydrogen Bond

	Pi-Pi Stacked
	Pi-Pi T-shaped
	Alkyl
	Pi-Alkyl
	Pi-Sigma

Figure S1. The proposed binding mode of newly synthesized hybrids (13a-j) docked in the active site of acetylcholinesterase (4EY7) protein; (2D and 3D ligand-receptor interactions).

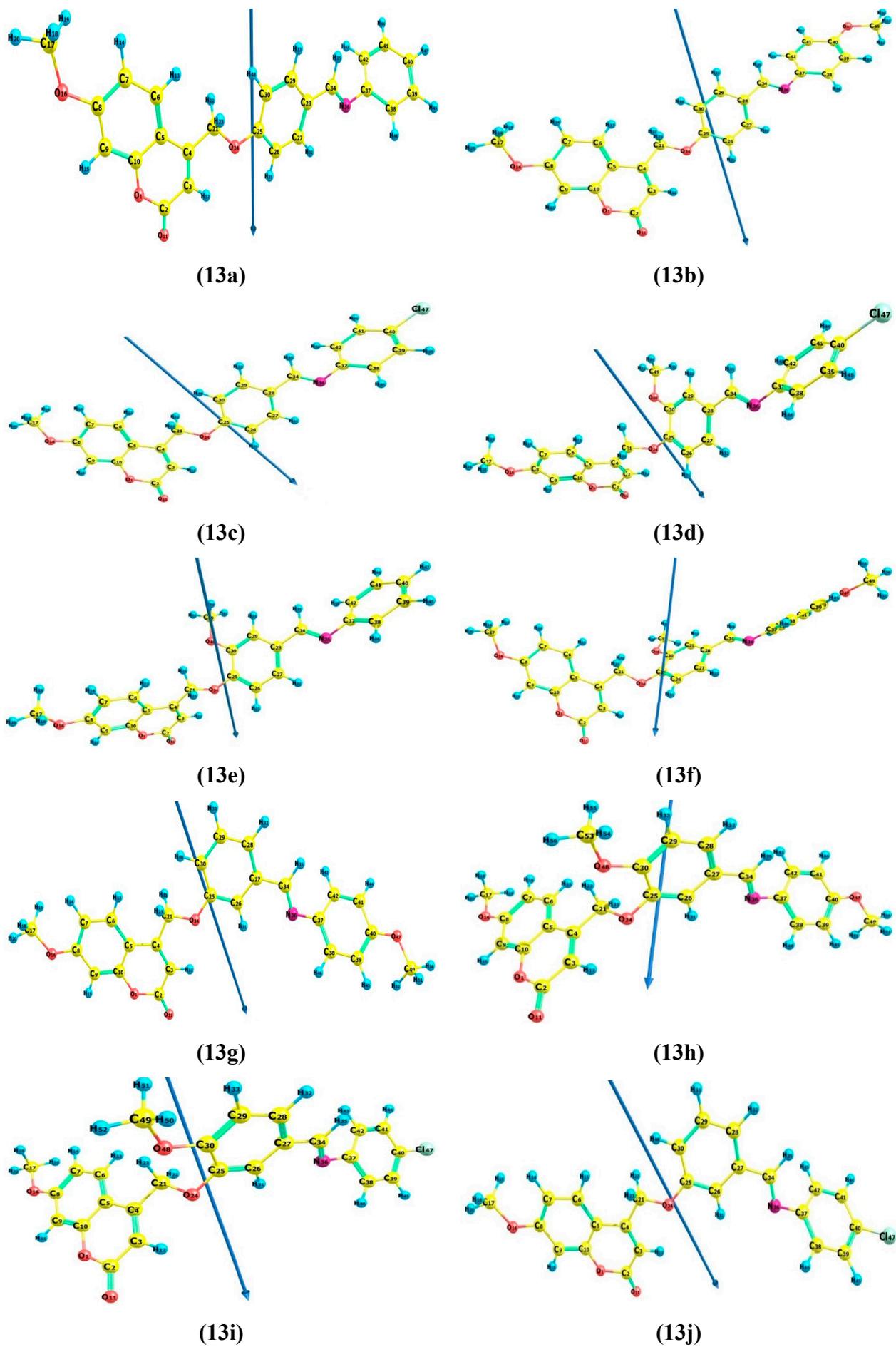
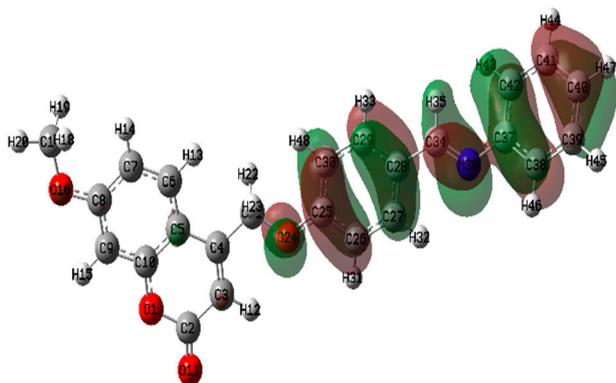
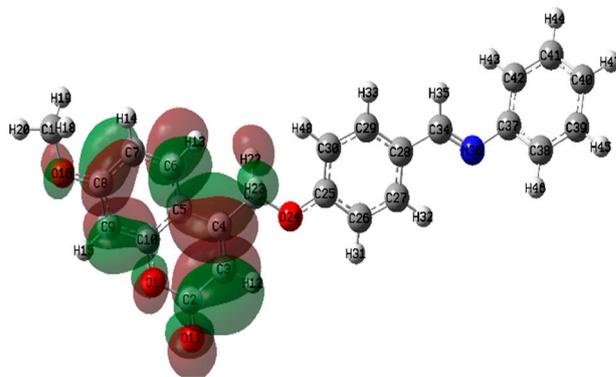


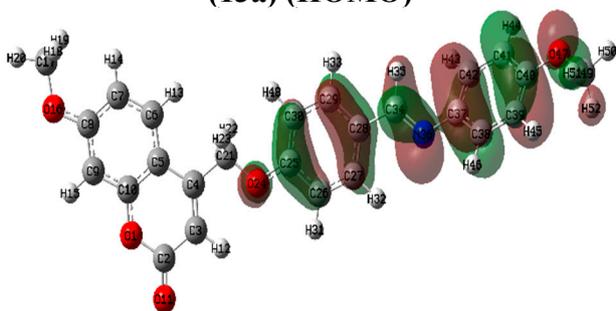
Figure S2. The optimized geometry, numbering system, vector of dipole moment of the synthesized compounds (13a-j) using wb97xd/6-311++g(d,p) level of calculation.



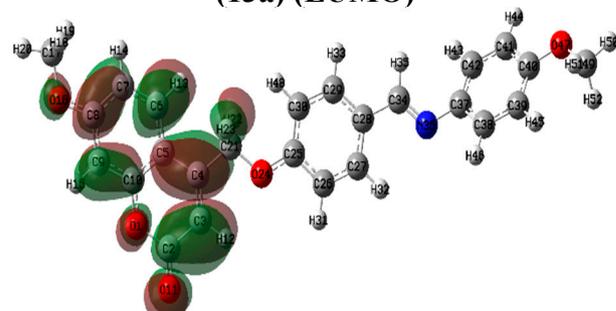
(13a) (HOMO)



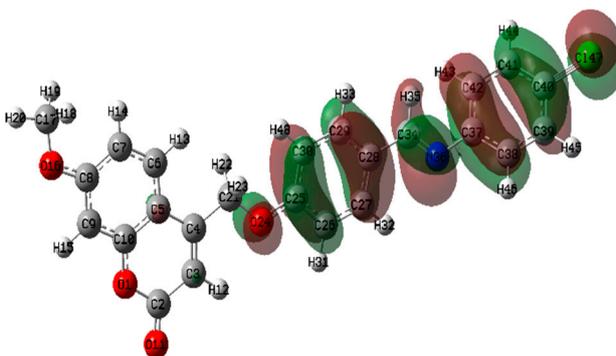
(13a) (LUMO)



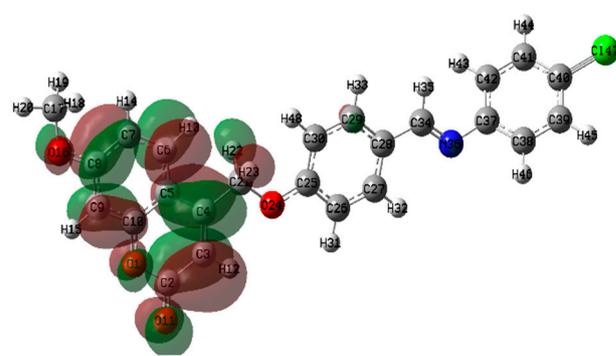
(13b) (HOMO)



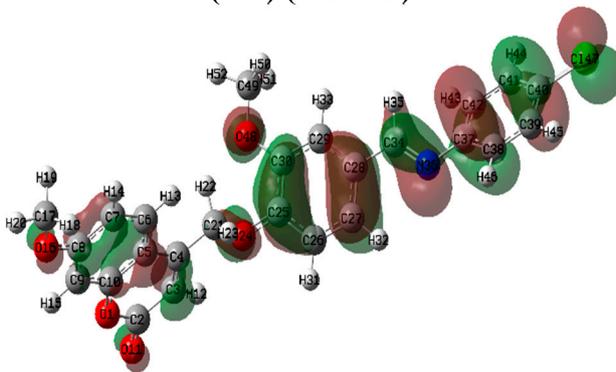
(13b) (LUMO)



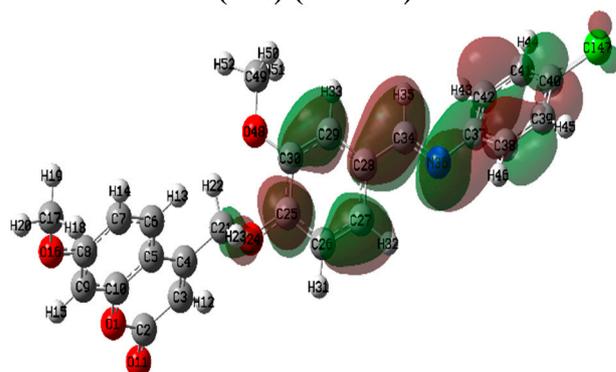
(13c) (HOMO)



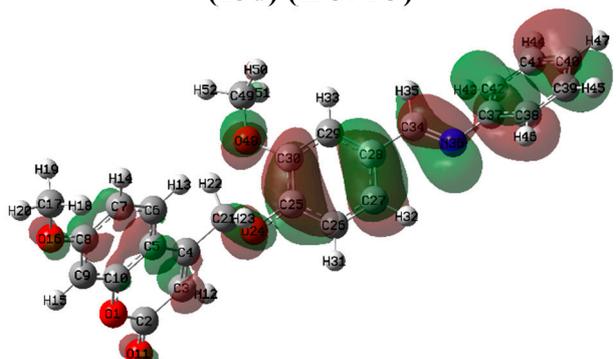
(13c) (LUMO)



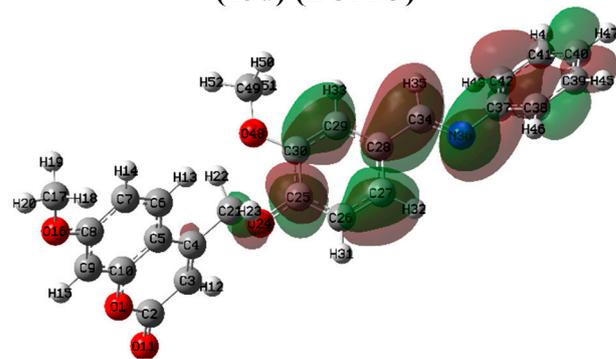
(13d) (HOMO)



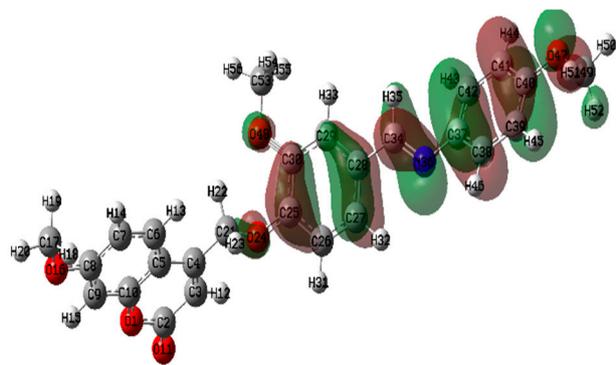
(13d) (LUMO)



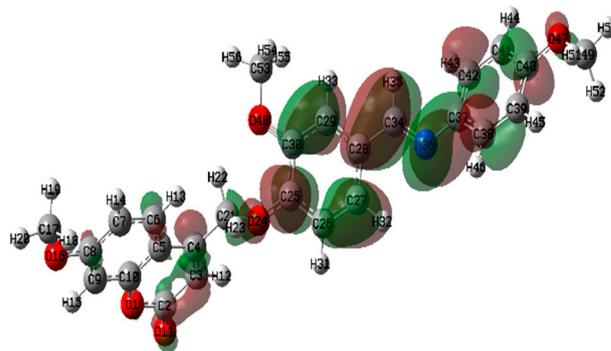
(13e) (HOMO)



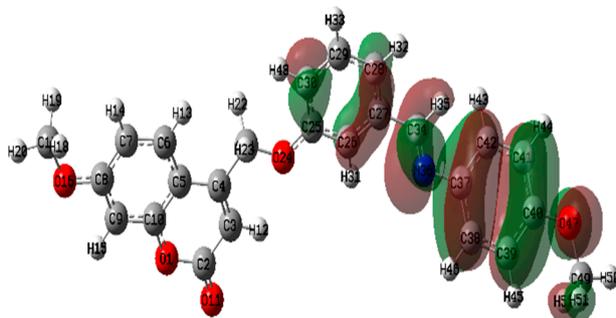
(13e) (LUMO)



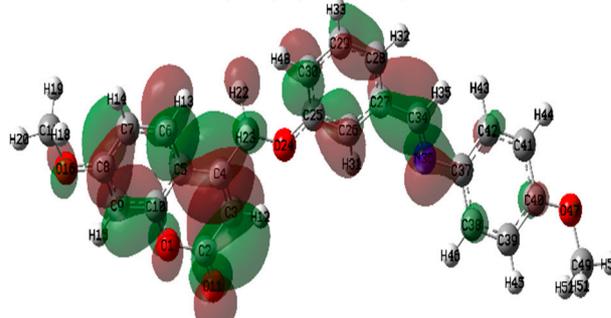
(13f) (HOMO)



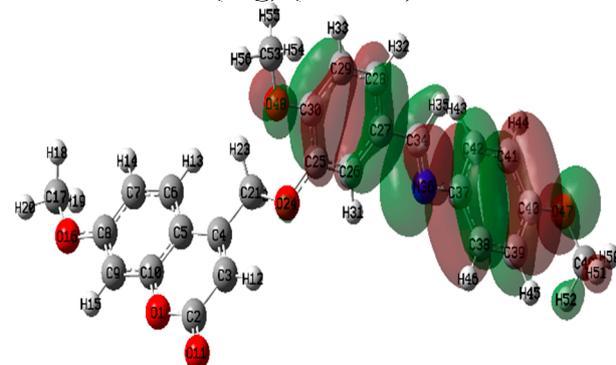
(13f) (LUMO)



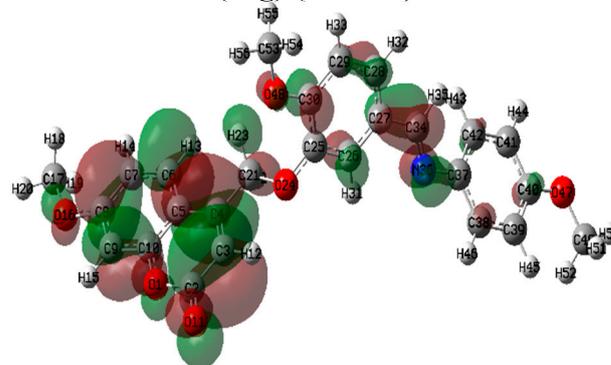
(13g) (HOMO)



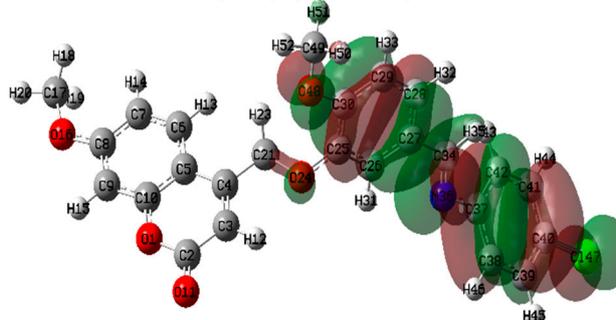
(13g) (LUMO)



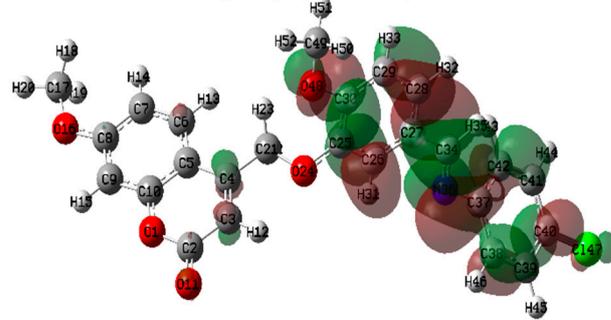
(13h) (HOMO)



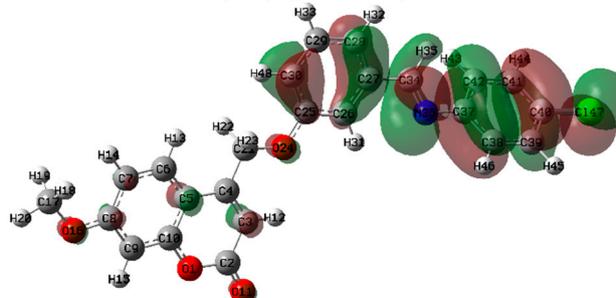
(13h) (LUMO)



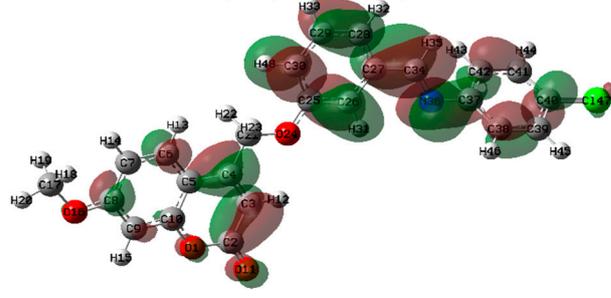
(13i) (HOMO)



(13i) (LUMO)

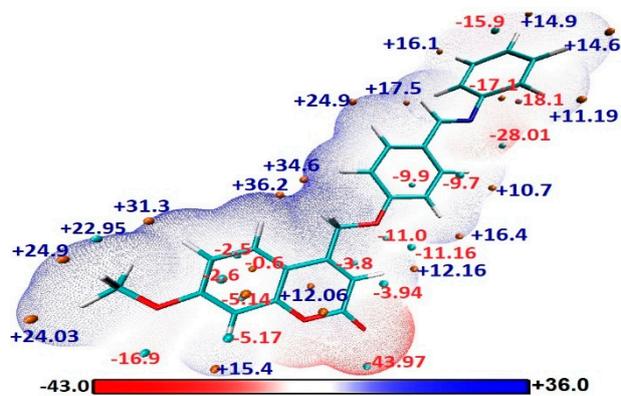


(13j) (HOMO)

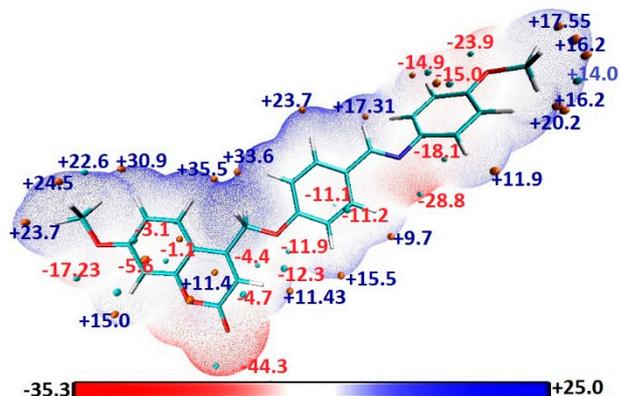


(13j) (LUMO)

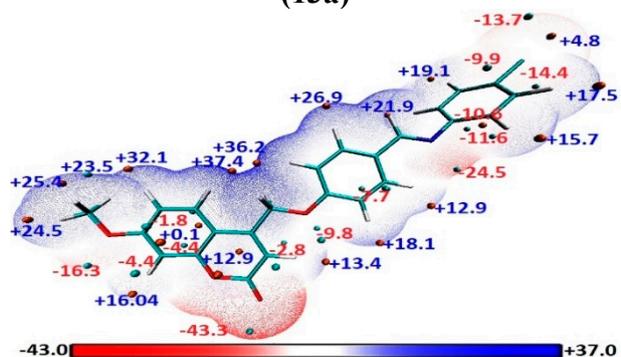
Figure S3. Frontier molecular orbitals of the synthesized compounds (13a-j) using wb97xd/6-311++g(d,p) level of calculation.



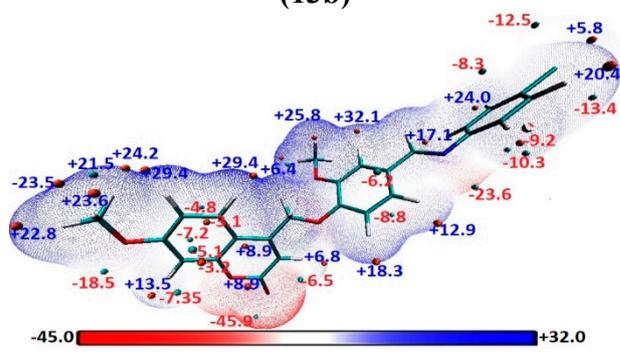
(13a)



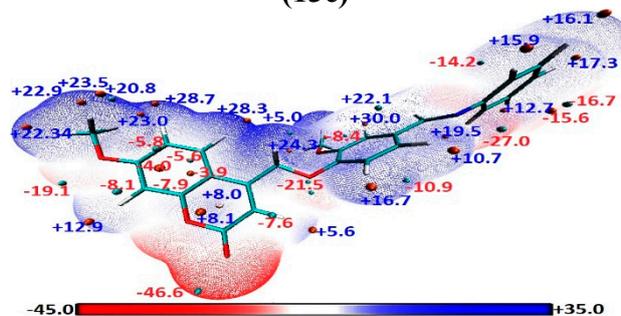
(13b)



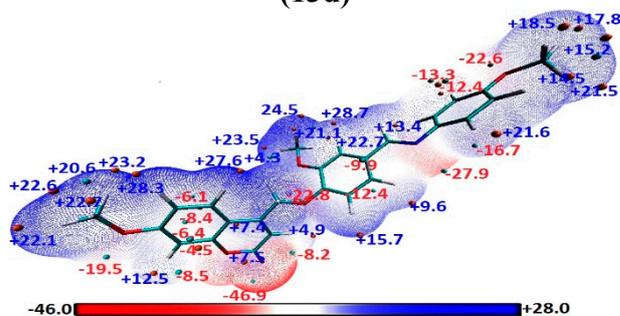
(13c)



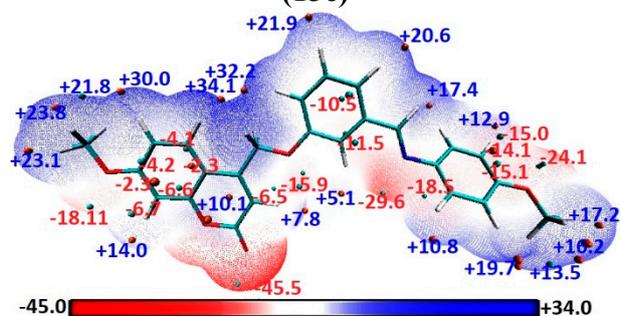
(13d)



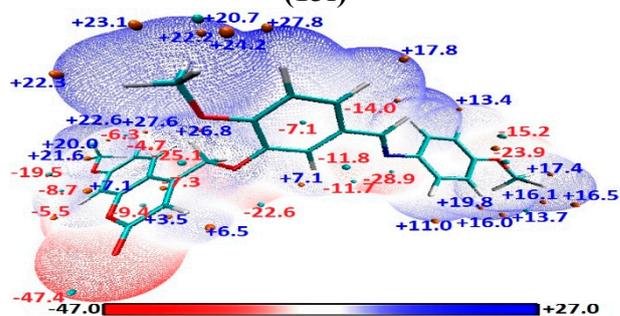
(13e)



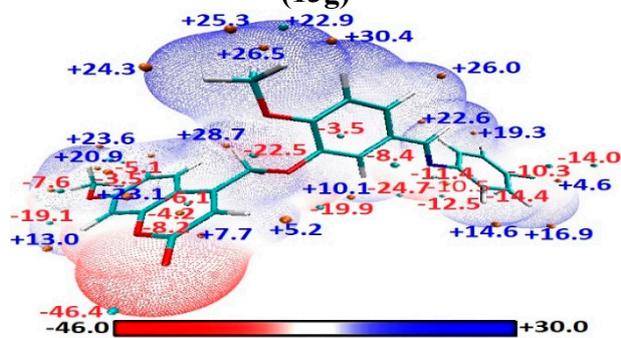
(13f)



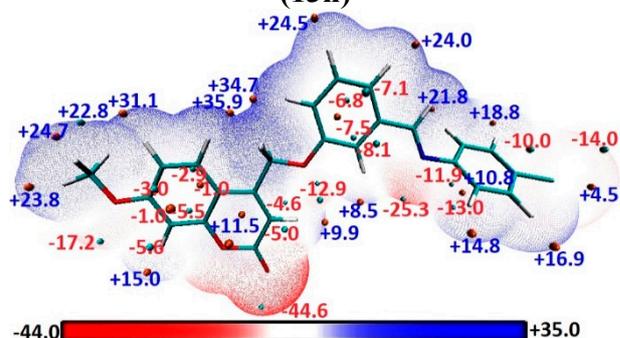
(13g)



(13h)



(13i)



(13j)

Figure S4. MEP surfaces of the synthesized compounds (13a-j) using wb97xd/6-311++g(d,p) level of calculation.

Table S1. The selected bond length (Å), bond angles and dihedral angles, (degree) of the newly synthesized compounds (**13a-j**) using wb97xd/6-311++g(d,p) level of level of theory.

	Exp. [6]	13a	13b	13c	13d	13e	13f	13g	13h	13i	13j
R(O1,C2)	1.383	1.380	1.380	1.380	1.381	1.381	1.381	1.381	1.382	1.382	1.381
R(O1,C10)	1.360	1.356	1.356	1.356	1.356	1.356	1.356	1.356	1.356	1.356	1.356
R(C2,C3)	1.450	1.457	1.457	1.458	1.456	1.456	1.456	1.458	1.456	1.456	1.458
R(C2,O11)	1.204	1.198	1.198	1.198	1.198	1.198	1.198	1.198	1.198	1.198	1.198
R(C4,C21)		1.501	1.501	1.501	1.499	1.499	1.499	1.502	1.499	1.499	1.502
R(C8,C9)	1.379	1.391	1.391	1.391	1.390	1.390	1.390	1.390	1.390	1.390	1.390
R(C8,O16)	1.388	1.349	1.349	1.348	1.350	1.350	1.350	1.349	1.350	1.350	1.349
R(C9,C10)	1.381	1.383	1.383	1.383	1.384	1.384	1.384	1.384	1.384	1.384	1.383
R(O16,C17)	1.363	1.414	1.414	1.414	1.413	1.413	1.413	1.413	1.413	1.413	1.414
R(C21,O24)		1.404	1.404	1.405	1.414	1.414	1.414	1.403	1.413	1.413	1.403
R(O24,C25)		1.357	1.358	1.356	1.365	1.365	1.366	1.361	1.369	1.367	1.360
R(C30,A48)		1.082	1.082	1.082	1.355	1.356	1.356	1.082	1.350	1.349	1.082
R(C34,N36)		1.270	1.270	1.271	1.270	1.269	1.270	1.269	1.270	1.271	1.269
R(N36,C37)		1.407	1.408	1.405	1.405	1.407	1.408	1.408	1.408	1.405	1.406
R(C40,A47)		1.084	1.358	1.749	1.749	1.084	1.357	1.357	1.358	1.749	1.749
A(C2,O1,C10)	122.1	122.5	122.5	122.5	122.5	122.5	122.5	122.5	122.5	122.5	122.5
A(O1,C2,C3)	117.1	116.5	116.6	116.6	116.5	116.5	116.6	116.6	116.5	116.5	116.5
A(O1,C2,O11)	117.4	117.8	117.8	117.8	117.7	117.7	117.7	117.8	117.6	117.6	117.8
A(C4,C5,C6)	125.0	125.0	125.0	124.9	124.9	124.9	124.9	124.9	124.9	124.9	124.9
A(C4,C5,C10)	117.4	117.6	117.7	117.6	117.7	117.7	117.7	117.7	117.7	117.7	117.6
A(C7,C8,O16)		124.2	124.2	124.2	124.2	124.2	124.2	124.2	124.2	124.2	124.2
A(C8,O16,C17)	119.1	118.8	118.8	118.8	118.7	118.7	118.7	118.7	118.7	118.7	118.7
A(C4,C21,O24)		109.8	109.9	109.9	109.7	109.7	109.7	109.9	110.1	110.0	109.8
A(C21,O24,C25)		118.6	118.5	118.5	116.0	115.9	115.9	118.5	114.5	114.8	118.6
A(O24,C25,C30)		124.5	124.4	124.4	121.2	121.2	121.2	124.2	119.8	119.9	124.2
A(C25,C26,C27)		120.1	120.1	120.1	120.9	120.9	120.9	120.0	120.6	120.6	120.0
A(C27,C28,C34)		121.5	121.5	121.5	121.5	121.5	121.6	120.0	119.7	119.8	120.1
A(C34,N36,C37)		119.1	119.3	119.1	119.2	119.1	119.6	119.7	119.3	119.2	119.2
A(N36,C37,C38)		118.3	118.4	118.4	118.2	118.2	118.1	118.0	118.5	118.4	118.3
A(C39,C40,C41)		119.5	119.5	120.9	120.9	119.6	119.5	119.5	119.5	120.9	121.0
D(C10,O1,C2,O11)	180.0	179.9	179.8	179.8	-179.9	180.0	-179.9	-179.9	179.6	179.6	-180.0
D(C2,C3,C4,C21)	179.1	-179.8	-179.5	-179.6	180.0	180.0	-180.0	179.9	179.5	179.5	-179.9
D(C5,C4,C21,O24)		179.8	179.3	-179.5	179.5	179.3	-179.9	179.0	-177.8	-177.6	178.8
D(C4,C5,C6,C7)	-179.1	-179.9	-179.8	179.9	-179.9	-179.8	179.9	180.0	-179.9	180.0	179.9
D(C4,C5,C10,C9)	179.6	179.9	179.9	-179.9	179.9	179.8	-179.9	-180.0	179.9	-179.9	-179.9
D(O16,C8,C9,C10)	178.6	-179.9	-179.8	-179.9	-179.9	-179.8	-180.0	179.9	-179.9	180.0	179.9
D(C7,C8,O16,C17)		0.3	0.0	0.4	-1.2	-1.4	-0.8	0.3	-0.3	0.1	0.4
D(C21,O24,C25,C26)		179.2	178.7	-179.9	115.1	114.8	115.0	176.5	-107.0	-108.0	177.9
D(C26,C27,C28,C34)		180.0	-180.0	-179.8	180.0	-179.9	179.8	180.0	180.0	179.9	-180.0
D(C27,C28,C34,N36)		-2.8	-3.9	-3.3	-1.1	-1.0	-0.6	179.9	179.5	179.5	179.8
D(C28,C34,N36,C37)		177.6	177.5	177.4	177.5	177.4	177.7	178.3	177.3	177.2	177.9
D(C34,N36,C37,C42)		-47.6	-42.2	-47.6	-44.7	-45.1	-38.5	-39.5	-41.8	-46.2	-46.8
D(N36,C37,C38,C39)		179.9	-179.7	179.9	-180.0	179.9	-179.6	-179.4	-179.8	179.6	180.0

Values are mean \pm SD triplicate assays.

Table S2: Mean absolute errors computed for selected bond lengths (Å) and angles (degree) of synthesized compounds (**13a-j**) versus **7-Acetoxy-coumarin** calculated at long-range corrections wb97xd/6-311++g(d,p) level of theory. The X-ray crystal structure data have been taken from ref. [1]

	13a	13b	13c	13d	13e	13f	13g	13h	13i	13j
R(O1,C2)	0.003	0.002	0.003	0.002	0.001	0.001	0.002	0.001	0.001	0.002
R(O1,C10)	0.004	0.004	0.004	0.004	0.004	0.004	0.004	0.004	0.004	0.004
R(C2,C3)	0.007	0.007	0.008	0.006	0.006	0.006	0.008	0.006	0.006	0.008
R(C2,O11)	0.007	0.007	0.007	0.006	0.006	0.006	0.007	0.006	0.006	0.007
R(C8,C9)	0.012	0.012	0.012	0.011	0.011	0.011	0.011	0.011	0.011	0.011
R(C8,O16)	0.039	0.039	0.039	0.038	0.038	0.038	0.039	0.038	0.038	0.039
R(C9,C10)	0.002	0.002	0.002	0.003	0.003	0.003	0.002	0.003	0.003	0.002
R(O16,C17)	0.051	0.051	0.051	0.050	0.050	0.050	0.050	0.050	0.050	0.051
A(C2,O1,C10)	0.436	0.427	0.434	0.385	0.389	0.375	0.423	0.402	0.414	0.444
A(O1,C2,C3)	0.562	0.551	0.549	0.554	0.557	0.548	0.540	0.555	0.561	0.555
A(O1,C2,O11)	0.450	0.430	0.462	0.306	0.289	0.273	0.370	0.231	0.263	0.395
A(C4,C5,C6)	0.014	0.059	0.069	0.124	0.140	0.132	0.082	0.142	0.123	0.068
A(C4,C5,C10)	0.250	0.277	0.271	0.301	0.308	0.313	0.293	0.317	0.301	0.272
A(C8,O16,C17)	0.337	0.331	0.325	0.398	0.374	0.433	0.407	0.410	0.392	0.388
D(C10,O1,C2,O11)	0.037	0.137	0.148	0.042	0.025	0.040	0.035	0.315	0.352	0.004
D(C2,C3,C4,C21)	0.697	0.422	0.545	0.907	0.955	0.904	0.819	0.415	0.468	0.855
D(C4,C5,C6,C7)	0.778	0.736	0.843	0.812	0.718	0.841	0.911	0.786	0.890	0.864
D(C4,C5,C10,C9)	0.333	0.322	0.338	0.341	0.250	0.390	0.417	0.370	0.359	0.362
D(O16,C8,C9,C10)	1.267	1.200	1.314	1.266	1.187	1.353	1.281	1.314	1.354	1.267

Table S3. Values of the Fukui functions and dual descriptor of compounds (13a-e) by using wb97xd/6-311++g(d,p) level of theory

Atom	(13a)			(13b)			(13c)			(13d)			(13e)		
	f(-)	f(+)	Δf	f(-)	f(+)	Δf	f(-)	f(+)	Δf	f(-)	f(+)	Δf	f(-)	f(+)	Δf
O1	0.000	0.021	0.021	0.000	0.021	0.021	0.000	0.021	0.020	0.001	0.000	-0.001	0.001	0.000	-0.001
C2	0.000	0.084	0.084	0.000	0.084	0.084	0.000	0.084	0.083	0.001	0.001	0.000	0.001	0.001	0.001
C3	0.004	0.199	0.195	0.000	0.198	0.198	0.003	0.198	0.194	0.011	0.007	-0.003	0.007	0.009	0.002
C4	0.001	0.209	0.208	0.000	0.209	0.209	0.001	0.207	0.206	0.016	-0.003	-0.019	0.018	0.000	-0.017
C6	0.001	0.166	0.165	0.000	0.167	0.167	0.001	0.164	0.163	0.004	-0.004	-0.008	0.003	-0.003	-0.006
C8	0.003	0.082	0.080	0.000	0.083	0.083	0.003	0.081	0.078	0.010	-0.001	-0.011	0.008	-0.001	-0.008
C9	0.001	0.052	0.051	0.000	0.052	0.052	0.001	0.051	0.050	0.004	0.001	-0.003	0.003	0.002	-0.002
C10	0.001	0.035	0.034	0.000	0.035	0.035	0.001	0.035	0.034	0.005	0.002	-0.002	0.004	0.003	-0.001
O11	0.002	0.065	0.063	0.000	0.065	0.065	0.002	0.065	0.063	0.007	0.000	-0.007	0.006	0.001	-0.005
O16	0.002	0.011	0.009	0.000	0.011	0.011	0.002	0.011	0.009	0.009	0.000	-0.009	0.007	0.000	-0.007
C21	0.001	-0.039	-0.039	0.000	-0.038	-0.039	0.001	-0.040	-0.041	-0.001	0.004	0.005	-0.003	0.002	0.004
O24	0.051	0.001	-0.050	0.021	0.001	-0.019	0.043	0.001	-0.042	0.029	0.006	-0.023	0.033	0.006	-0.027
C25	0.073	0.004	-0.069	0.036	0.004	-0.033	0.061	0.004	-0.057	0.093	0.112	0.019	0.109	0.117	0.009
C26	0.027	0.000	-0.027	0.009	0.000	-0.009	0.022	0.000	-0.022	0.010	0.010	0.000	0.010	0.013	0.004
C27	0.008	0.001	-0.008	0.011	0.001	-0.010	0.008	0.001	-0.008	0.073	0.085	0.012	0.086	0.089	0.003
C28	0.096	0.000	-0.096	0.047	0.000	-0.047	0.083	0.001	-0.082	0.068	0.082	0.014	0.077	0.087	0.010
C29	0.037	0.002	-0.035	0.025	0.002	-0.023	0.031	0.004	-0.027	-0.008	0.075	0.083	-0.010	0.073	0.083
C30	0.040	0.003	-0.038	0.015	0.003	-0.013	0.034	0.002	-0.031	0.042	0.023	-0.019	0.052	0.025	-0.027
C34	0.021	0.002	-0.020	0.052	0.001	-0.051	0.021	0.005	-0.016	0.027	0.238	0.211	0.029	0.229	0.200
N36	0.141	0.001	-0.141	0.103	0.000	-0.103	0.128	0.002	-0.126	0.104	0.164	0.060	0.114	0.168	0.053
C37	0.129	0.000	-0.129	0.144	0.000	-0.144	0.125	0.000	-0.125	0.092	0.051	-0.041	0.093	0.043	-0.050
C38	0.089	0.000	-0.089	0.092	0.000	-0.092	0.078	0.000	-0.077	0.060	0.016	-0.045	0.067	0.017	-0.050
C39	0.016	0.000	-0.016	0.055	0.000	-0.055	0.023	0.000	-0.024	0.018	0.001	-0.017	0.012	0.006	-0.007
C40	0.138	0.000	-0.138	0.133	0.000	-0.133	0.128	0.001	-0.128	0.101	0.052	-0.049	0.107	0.050	-0.058
C41	0.032	0.000	-0.032	0.078	0.000	-0.078	0.039	0.000	-0.039	0.034	0.017	-0.017	0.029	0.007	-0.022
C42	0.058	0.000	-0.058	0.038	0.000	-0.038	0.057	0.000	-0.056	0.050	0.032	-0.018	0.052	0.029	-0.023
A47	0.001	0	-0.001	0.1063	0	-0.1063	0.0796	0.0001	-0.0795	0.0622	0.0066	-0.0556	0.0008	0.0007	-0.0001
A48	0.0005	0.0001	-0.0004	0.0002	0.0003	0.0001	0.0004	0.0001	-0.0003	0.0333	0.0006	-0.0327	0.0402	0.0006	-0.0396

A = H, O or Cl which different atom in derivatives series; Values are mean \pm SD triplicate assays.

Table S4. Values of the Fukui functions and dual descriptor of compounds (**13f-j**) by using wb97xd/6-311++g(d,p) level of theory.

Atom	(13f)			(13g)			(13h)			(13i)			(13j)		
	f(-)	f(+)	Δf												
O1	0.000	0.001	0.001	0.000	0.016	0.016	0.000	0.017	0.017	0.000	0.001	0.001	0.000	0.004	0.003
C2	0.000	0.005	0.005	0.000	0.063	0.063	0.000	0.073	0.073	0.001	0.003	0.002	0.000	0.013	0.013
C3	0.000	0.017	0.017	0.000	0.153	0.153	-0.001	0.163	0.164	-0.001	0.005	0.007	0.008	0.036	0.027
C4	0.001	0.009	0.008	0.000	0.162	0.162	0.000	0.183	0.183	0.000	-0.001	-0.001	0.002	0.038	0.037
C6	0.001	0.004	0.004	0.000	0.121	0.121	0.000	0.152	0.152	0.001	0.009	0.008	0.001	0.023	0.021
C8	0.000	0.002	0.002	0.000	0.062	0.062	0.000	0.073	0.073	0.000	0.002	0.002	0.005	0.012	0.008
C9	0.000	0.004	0.004	0.000	0.038	0.038	0.000	0.048	0.048	0.000	0.001	0.001	0.002	0.007	0.005
C10	0.000	0.004	0.004	0.000	0.028	0.028	0.000	0.029	0.029	0.000	-0.004	-0.003	0.002	0.007	0.005
O11	0.000	0.004	0.004	0.000	0.049	0.049	0.000	0.056	0.056	0.000	0.002	0.002	0.004	0.011	0.007
O16	0.000	0.001	0.000	0.000	0.009	0.009	0.000	0.010	0.010	0.000	0.000	0.000	0.004	0.002	-0.002
C21	0.007	-0.001	-0.008	0.000	-0.040	-0.040	0.002	-0.036	-0.039	0.005	0.013	0.008	0.000	-0.017	-0.017
O24	0.014	0.006	-0.008	0.001	0.002	0.001	0.002	0.001	-0.002	0.007	0.001	-0.006	0.006	0.001	-0.005
C25	0.051	0.116	0.065	-0.001	0.003	0.005	0.011	0.005	-0.006	0.029	0.004	-0.025	0.005	0.007	0.003
C26	0.003	0.015	0.012	-0.003	0.031	0.034	0.006	0.021	0.015	0.007	0.087	0.080	-0.017	0.098	0.116
C27	0.038	0.083	0.045	0.023	0.026	0.003	0.046	0.009	-0.036	0.083	0.074	-0.009	0.031	0.078	0.047
C28	0.042	0.087	0.044	0.028	0.007	-0.021	0.030	0.004	-0.026	0.042	0.069	0.027	0.039	0.032	-0.007
C29	0.007	0.068	0.062	0.003	0.014	0.010	0.014	-0.002	-0.016	0.031	0.018	-0.013	0.003	0.037	0.034
C30	0.018	0.023	0.004	0.029	0.042	0.013	0.042	0.030	-0.012	0.074	0.115	0.041	0.045	0.116	0.072
C34	0.055	0.215	0.160	0.079	0.055	-0.024	0.057	0.032	-0.025	0.035	0.235	0.200	0.050	0.197	0.147
N36	0.096	0.162	0.066	0.094	0.040	-0.053	0.103	0.024	-0.079	0.125	0.157	0.032	0.126	0.137	0.011
C37	0.135	0.041	-0.095	0.159	0.005	-0.154	0.132	0.005	-0.128	0.105	0.051	-0.054	0.151	0.036	-0.115
C38	0.083	0.023	-0.059	0.091	0.008	-0.083	0.088	0.007	-0.081	0.071	0.016	-0.055	0.084	0.015	-0.068
C39	0.053	-0.002	-0.055	0.065	-0.001	-0.065	0.053	0.000	-0.053	0.022	-0.001	-0.023	0.034	-0.004	-0.037
C40	0.128	0.043	-0.085	0.145	0.009	-0.136	0.130	0.006	-0.124	0.120	0.051	-0.069	0.159	0.039	-0.120
C41	0.078	0.009	-0.069	0.090	0.002	-0.088	0.079	0.002	-0.077	0.038	0.019	-0.019	0.050	0.012	-0.039
C42	0.047	0.022	-0.025	0.042	0.004	-0.038	0.044	0.000	-0.044	0.060	0.025	-0.035	0.068	0.018	-0.049
A47	0.102	0.0055	-0.0965	0.1224	0.0012	-0.1212	0.1034	0.0007	-0.1027	0.0731	0.0062	-0.0669	0.1083	0.0045	-0.1038
A48	0.0109	0.0007	-0.0102	0.0002	0.0007	0.0005	0.0249	0.0032	-0.0217	0.0532	0.016	-0.0372	0.0003	0.0016	0.0013

A = H, O or Cl which different atom in derivatives series; Values are mean \pm SD triplicate assays.

Table S5. Values of the condensed local softnesses (Hartree*e) of compounds (**13a-j**) by using wb97xd/6-311++g(d,p) level of theory from CDFT point of view.

Atoms	13a		13b		13c		13d		13e		13f		13g		13h		13i		13j	
	s+/s-	s-/s+	s+/s-	s-/s+	s+/s-	s-/s+	s+/s-	s-/s+	s+/s-	s-/s+	s+/s-	s-/s+	s+/s-	s-/s+	s+/s-	s-/s+	s+/s-	s-/s+	s+/s-	s-/s+
O1	0.5	2.0	0.1	7.1	0.2	5.9	0.0	-51.1	1.1	0.9	0.8	1.3	0.6	1.8	0.1	7.3	0.2	4.5	2.7	0.4
C2	0.2	5.0	0.0	23.9	0.1	19.0	0.0	26.6	1.0	1.0	0.7	1.5	0.2	4.4	0.0	29.1	0.1	15.7	4.2	0.2
C3	0.0	31.6	0.0	1993.5	0.0	186.6	0.0	-35.7	1.2	0.8	0.4	2.8	-1.2	-0.8	0.0	-144.2	0.0	-163.1	225.3	0.0
C8	0.5	2.1	0.1	9.4	0.1	7.7	0.1	9.8	1.1	0.9	0.8	1.3	0.6	1.6	0.1	9.9	0.2	5.8	5.1	0.2
C9	0.5	2.0	0.1	7.2	0.2	6.0	0.1	8.3	1.1	0.9	0.8	1.3	0.6	1.7	0.1	7.7	0.2	4.7	3.2	0.3
C10	0.6	1.7	0.1	8.2	0.2	6.6	0.1	6.9	1.1	0.9	0.8	1.3	0.5	2.1	0.1	10.2	0.2	5.7	9.1	0.1
O11	0.4	2.6	0.1	10.6	0.1	8.5	0.1	10.7	1.1	0.9	0.7	1.3	0.3	3.5	0.1	11.0	0.2	6.4	5.8	0.2
O16	0.6	1.6	0.2	5.2	0.2	4.3	0.1	18.6	1.1	0.9	0.8	1.2	0.7	1.5	0.2	5.3	0.3	3.4	7.3	0.1
C21	0.6	1.7	0.2	4.2	0.3	3.0	0.1	14.4	1.0	1.0	0.6	1.7	0.4	2.5	0.1	10.5	0.2	4.4	0.9	1.2
O24	2.1	0.5	-2.4	-0.4	-3.3	-0.3	-1.6	-0.6	1.8	0.6	1.0	1.0	0.7	1.5	-2.6	-0.4	12.9	0.1	0.5	1.8
C25	1.0	1.0	-3.1	-0.3	-4.0	-0.3	-2.0	-0.5	1.1	0.9	0.7	1.5	0.5	2.0	-1.7	-0.6	33.9	0.0	0.1	7.4
C26	1.2	0.8	20.1	0.0	25.1	0.0	2.6	0.4	0.8	1.2	0.6	1.8	0.3	3.6	7.4	0.1	1.0	1.0	0.1	15.5
C27	0.7	1.5	1.6	0.6	2.1	0.5	5.8	0.2	1.2	0.8	0.6	1.7	0.0	-23.0	0.9	1.1	1.8	0.5	0.3	3.3
C28	1.1	0.9	0.5	2.2	1.7	0.6	-0.3	-3.1	1.4	0.7	0.2	4.2	0.6	1.6	2.3	0.4	1.8	0.6	0.8	1.3
C29	0.9	1.1	4.5	0.2	5.5	0.2	53.7	0.0	0.4	2.4	0.4	2.6	0.4	2.3	2.3	0.4	1.7	0.6	0.3	3.3
C30	1.3	0.8	-13.8	-0.1	-19.8	-0.1	0.3	3.6	1.4	0.7	0.6	1.5	0.5	1.9	-7.6	-0.1	2.7	0.4	0.3	3.8
C34	0.5	2.2	-27.5	0.0	-60.1	0.0	2.3	0.4	0.3	3.9	0.5	2.2	0.6	1.7	-48.5	0.0	0.9	1.1	0.1	7.5
N36	0.9	1.1	3.6	0.3	5.2	0.2	1.0	1.0	0.7	1.5	0.4	2.3	0.4	2.3	8.5	0.1	1.9	0.5	0.2	4.1
C37	9.6	0.1	-22.3	0.0	-16.1	-0.1	1.8	0.5	4.5	0.2	22.7	0.0	67.7	0.0	-16.4	-0.1	1321.0	0.0	3.5	0.3
C38	2.6	0.4	18.7	0.1	20.1	0.0	42.8	0.0	1.6	0.6	2.0	0.5	2.3	0.4	58.7	0.0	5.7	0.2	0.7	1.4
C39	1.5	0.7	9.5	0.1	6.6	0.2	1.8	0.5	1.1	0.9	1.9	0.5	2.3	0.4	10.9	0.1	2.9	0.3	0.6	1.6
C40	2.1	0.5	9.6	0.1	9.7	0.1	-0.1	-7.3	1.4	0.7	1.7	0.6	2.0	0.5	9.6	0.1	3.6	0.3	0.8	1.3
C41	1.8	0.5	12.6	0.1	9.3	0.1	2.0	0.5	1.2	0.8	2.4	0.4	2.8	0.4	10.8	0.1	3.4	0.3	0.8	1.3
C42	2.5	0.4	29.7	0.0	28.1	0.0	36.4	0.0	1.6	0.6	1.8	0.6	2.1	0.5	33.4	0.0	5.1	0.2	0.8	1.3
A47	1.5	0.7	15.4	0.1	8.0	0.1	-2.2	-0.5	1.1	0.9	3.4	0.3	4.1	0.2	14.1	0.1	3.8	0.3	0.9	1.1

A48	1.2	0.9	-10.4	-0.1	-13.7	-0.1	-1.7	-0.6	2.1	0.5	1.0	1.0	0.5	1.9	-2.3	-0.4	23.8	0.0	0.2	5.5
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Table S6. Values of the condensed local electrophilicity (EIP)/nucleophilicity (NuP) index (e^*eV) of compounds (**13a-j**) by using wb97xd/6-311++g(d,p) level of theory from CDFT point of view.

Atoms	13a		13b		13c		13d		13e		13f		13g		13h		13i		13j	
	EIP	NuP																		
O1	-0.01	-0.01	-0.01	-0.05	-0.01	-0.03	0.00	-0.04	-0.01	-0.01	-0.01	-0.01	0.00	-0.01	0.00	-0.05	-0.01	-0.03	-0.02	-0.01
C2	0.00	-0.02	0.00	-0.10	0.00	-0.07	0.00	-0.06	-0.01	0.00	0.00	-0.01	0.00	-0.01	0.00	-0.10	0.00	-0.05	-0.02	0.00
C3	0.00	-0.02	0.00	-0.15	0.00	-0.10	0.00	-0.09	0.00	0.00	0.00	0.00	0.00	0.00	0.00	-0.15	0.00	-0.06	-0.08	0.00
C8	-0.01	-0.02	-0.01	-0.08	-0.01	-0.05	0.00	-0.05	-0.01	-0.01	-0.01	-0.01	-0.01	-0.01	-0.01	-0.09	-0.01	-0.04	-0.06	-0.01
C9	-0.01	-0.02	-0.01	-0.06	-0.01	-0.04	0.00	-0.04	-0.01	-0.01	-0.01	-0.01	-0.01	-0.01	-0.01	-0.07	-0.01	-0.03	-0.03	-0.01
C10	0.00	-0.01	0.00	-0.03	0.00	-0.02	0.00	-0.02	0.00	0.00	0.00	0.00	0.00	0.00	0.00	-0.03	0.00	-0.01	-0.03	0.00
O11	-0.01	-0.03	-0.01	-0.15	-0.01	-0.10	0.00	-0.10	-0.02	-0.01	-0.01	-0.02	0.00	-0.02	-0.01	-0.16	-0.01	-0.07	-0.09	-0.01
O16	-0.01	-0.01	-0.01	-0.04	-0.01	-0.03	0.00	-0.03	-0.01	-0.01	-0.01	-0.01	-0.01	-0.01	-0.01	-0.04	-0.01	-0.02	-0.06	-0.01
C21	-0.01	-0.01	0.00	-0.02	-0.01	-0.01	0.00	-0.01	-0.01	-0.01	0.00	-0.01	0.00	-0.01	0.00	-0.02	0.00	-0.01	0.00	0.00
O24	-0.04	-0.02	-0.03	0.02	-0.04	0.01	0.00	0.02	-0.04	-0.02	-0.02	-0.03	-0.01	-0.02	-0.01	0.00	-0.01	0.00	-0.01	-0.01
C25	-0.05	-0.05	-0.04	0.02	-0.06	0.01	-0.01	0.02	-0.08	-0.07	-0.05	-0.09	-0.02	-0.04	-0.02	0.02	-0.03	0.00	-0.01	-0.02
C26	-0.03	-0.03	-0.02	0.00	-0.03	0.00	0.00	0.00	-0.03	-0.03	-0.02	-0.05	-0.02	-0.08	-0.01	0.00	-0.02	-0.02	0.00	-0.05
C27	-0.03	-0.04	-0.02	-0.02	-0.03	-0.01	0.00	0.00	-0.07	-0.05	-0.03	-0.07	0.00	-0.06	-0.01	-0.01	-0.04	-0.02	-0.01	-0.03
C28	-0.03	-0.03	-0.01	-0.02	-0.03	-0.01	0.00	-0.02	-0.05	-0.03	-0.01	-0.05	-0.02	-0.05	-0.03	-0.02	-0.04	-0.02	-0.03	-0.03
C29	-0.04	-0.04	-0.03	-0.01	-0.04	-0.01	0.00	0.00	-0.02	-0.04	-0.02	-0.06	-0.02	-0.07	-0.02	-0.01	-0.03	-0.02	-0.02	-0.04
C30	-0.04	-0.03	-0.02	0.00	-0.04	0.00	0.00	-0.01	-0.05	-0.03	-0.02	-0.04	-0.04	-0.10	-0.04	0.01	-0.05	-0.02	-0.02	-0.06
C34	-0.04	-0.09	-0.05	0.00	-0.05	0.00	0.00	-0.01	-0.03	-0.11	-0.05	-0.15	-0.06	-0.14	-0.05	0.00	-0.04	-0.04	-0.02	-0.09
N36	-0.08	-0.08	-0.05	-0.02	-0.08	-0.01	0.00	-0.02	-0.07	-0.10	-0.04	-0.14	-0.04	-0.13	-0.04	-0.01	-0.07	-0.03	-0.03	-0.08
C37	-0.05	0.00	-0.08	0.00	-0.06	0.00	-0.01	-0.03	-0.03	-0.01	-0.07	0.00	-0.09	0.00	-0.07	0.01	-0.04	0.00	-0.02	0.00
C38	-0.06	-0.02	-0.06	0.00	-0.06	0.00	-0.01	0.00	-0.05	-0.03	-0.06	-0.04	-0.07	-0.04	-0.06	0.00	-0.05	-0.01	-0.02	-0.02
C39	-0.04	-0.02	-0.05	-0.01	-0.04	0.00	-0.01	-0.02	-0.03	-0.03	-0.05	-0.03	-0.05	-0.03	-0.04	-0.01	-0.03	-0.01	-0.02	-0.02
C40	-0.09	-0.04	-0.08	-0.01	-0.08	-0.01	0.00	-0.06	-0.08	-0.05	-0.07	-0.06	-0.08	-0.05	-0.07	-0.01	-0.06	-0.02	-0.03	-0.03
C41	-0.04	-0.02	-0.06	-0.01	-0.04	0.00	-0.01	-0.02	-0.04	-0.03	-0.06	-0.03	-0.07	-0.03	-0.05	-0.01	-0.04	-0.01	-0.02	-0.02

C42	-0.05	-0.02	-0.04	0.00	-0.05	0.00	-0.01	0.00	-0.04	-0.02	-0.04	-0.03	-0.04	-0.03	-0.04	0.00	-0.04	-0.01	-0.02	-0.02
A47	-0.04	-0.03	-0.08	-0.01	-0.14	-0.01	-0.05	0.17	-0.04	-0.03	-0.07	-0.03	-0.08	-0.03	-0.07	-0.01	-0.12	-0.03	-0.07	-0.05
A48	-0.02	-0.02	-0.02	0.00	-0.02	0.00	0.00	0.00	-0.04	-0.02	-0.02	-0.02	-0.02	-0.05	-0.03	0.02	-0.05	0.00	-0.01	-0.03

Values are mean \pm SD triplicate assays

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