Electronic Supplementary Material (ESI) for New Journal of Chemistry.

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Supporting Information

Molecular hybrid design, synthesis and biological evaluation of N-phenyl sulfonamide linked N-acyl hydrazone derivatives functioning as COX-2 inhibitors: New anti-inflammatory, anti-oxidant and anti-bacterial agents

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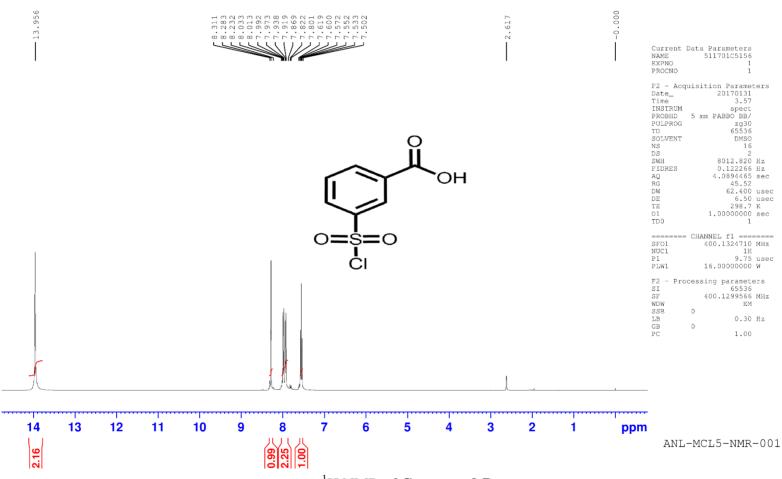
^dDepartment of Pharmacology, Bapatla College of Pharmacy, Bapatla Guntur (Dist.), Andhra Pradesh, Bapatla, India, 522 101

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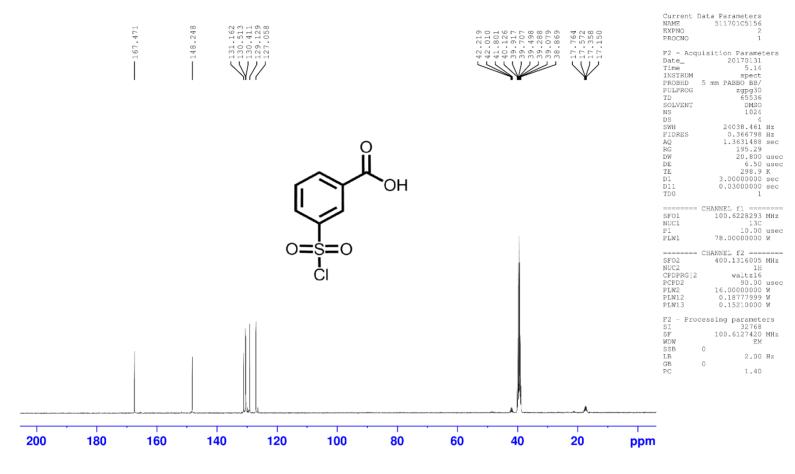
1. ¹H NMR, ¹³C NMR, HRMS, DSC

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GVB-SK-INT-1

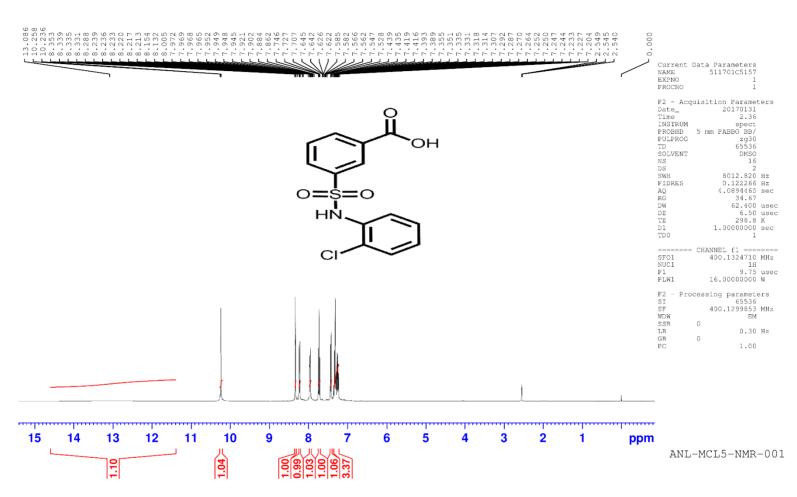


¹H NMR of **Compound-B**

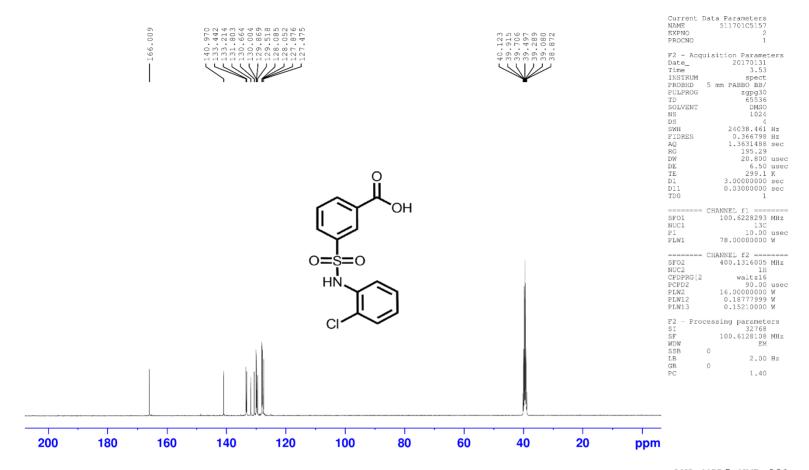


¹³CNMR of **Compound-B**in DMSO-d₆





 1 H NMR of **Compound-C**



ANL-MCL5-NMR-001

¹³CNMR of **Compound-C** in DMSO-d₆

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Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

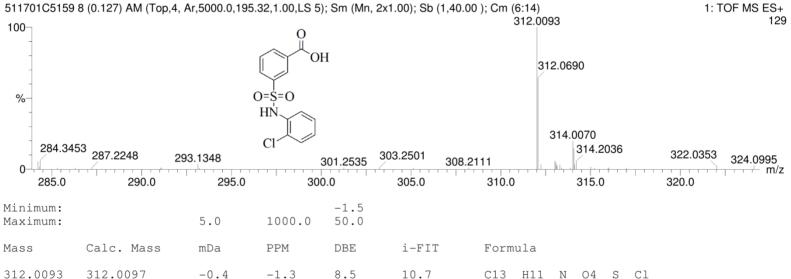
Selected filters: None

Monoisotopic Mass, Even Electron Ions

38 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass) Elements Used:

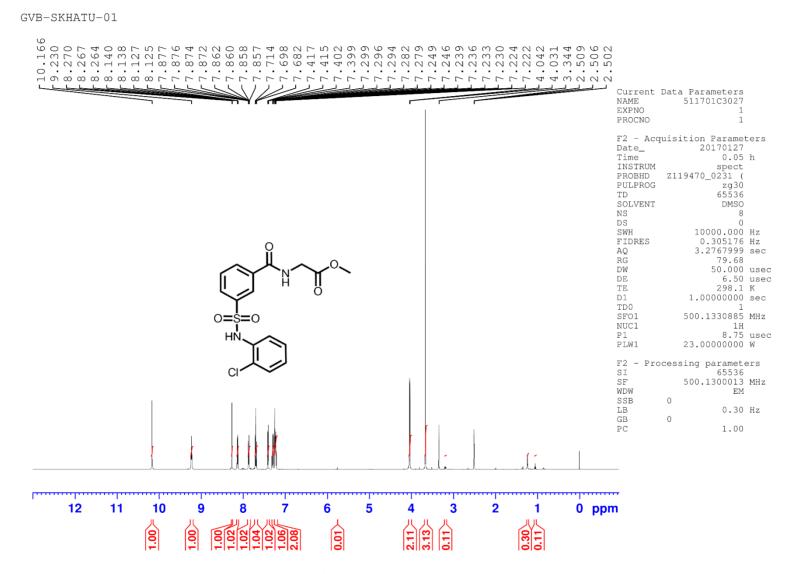
C: 0-13 H: 0-11 N: 0-1 O: 0-4 S: 0-1 CI: 0-1

GVB-SK-INT-2

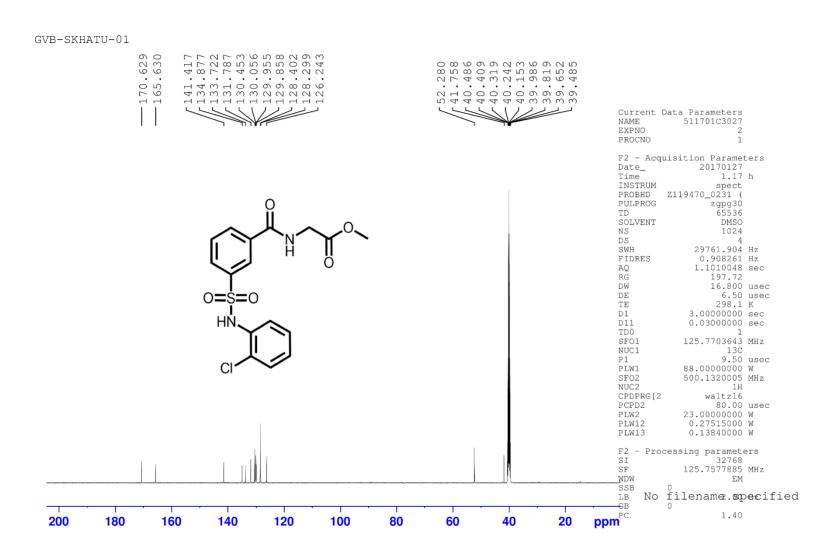


HRMS of Compound-C





¹H NMR of **Compound-D**



¹³CNMR of **Compound-D** in DMSO-d₆

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Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

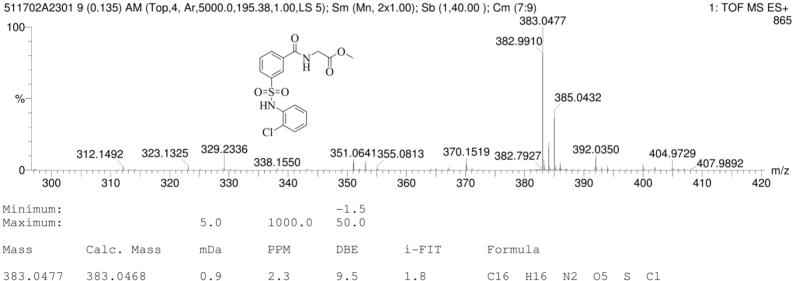
Selected filters: None

Monoisotopic Mass, Even Electron Ions

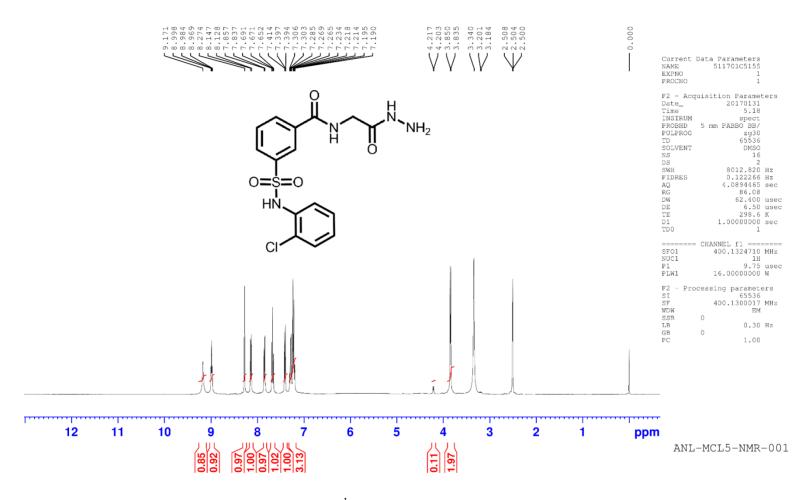
70 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass) Elements Used:

C: 0-16 H: 0-16 N: 0-2 O: 0-5 S: 0-1 Cl: 0-1

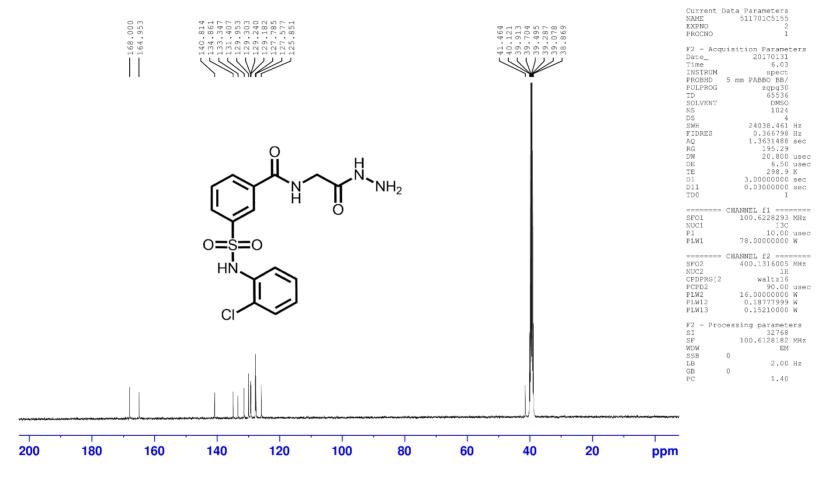
GVBSK-HATU-01



HRMS of Compound-D



 1 H NMR of **Compound-E**



ANL-MCL5-NMR-001

¹³CNMR of **Compound-E** in DMSO-d₆

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520

480

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

240

260

280

300

320

340

98 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass)

Elements Used:

C: 0-15 H: 0-16 N: 0-4 O: 0-4 S: 0-1 CI: 0-1

GVB-SK-INT-4

511701C5154 8 (0.126) AM (Top,4, Ar,5000.0,195.39,1.00,LS 5); Sm (Mn, 2x1.00); Sb (1,40.00) 1: TOF MS ES+ 383.0567 3.45e3 100-% 385.0999 195.4140 452.9772 196.4205 _{236.3634} 351.1158 424.0754 455.0133 294.2208 523.0003 576.7888 0 1 m/z ┍╗╇╬╌╒╌╌╌╌╌╌╌┼╒╠╚╌┰╌╌╌┼╒┼┼╬╬╌╌╌╌╌╌╌╬╌╌┼┼┼┼

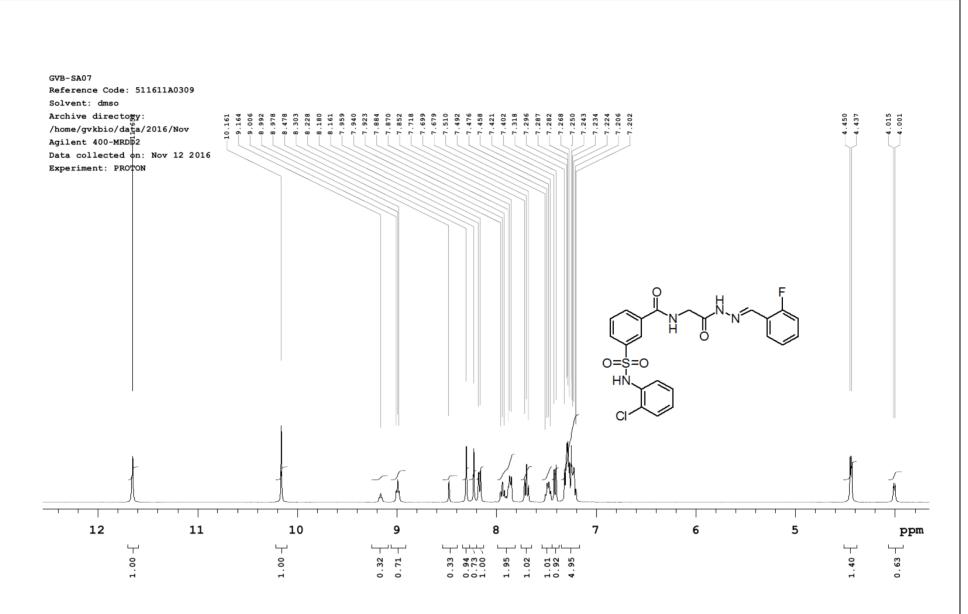
400

Minimum: -1.550.0 Maximum: 5.0 1000.0 i-FIT Mass Calc. Mass mDa PPM DBE Formula -1.4-3.79.5 266.2 383.0567 383.0581 C15 H16 N4 O4 S Cl

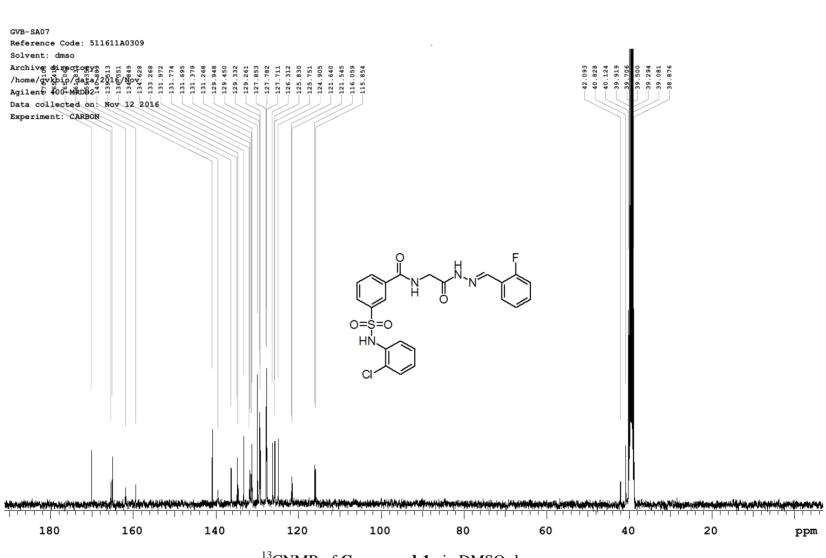
360

380

HRMS of Compound-E



¹H NMR of **Compound-1a**



 13 CNMR of **Compound-1a** in DMSO-d₆

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Single Mass Analysis

Tolerance = 10000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

178 formula(e) evaluated with 2 results within limits (up to 1 closest results for each mass)

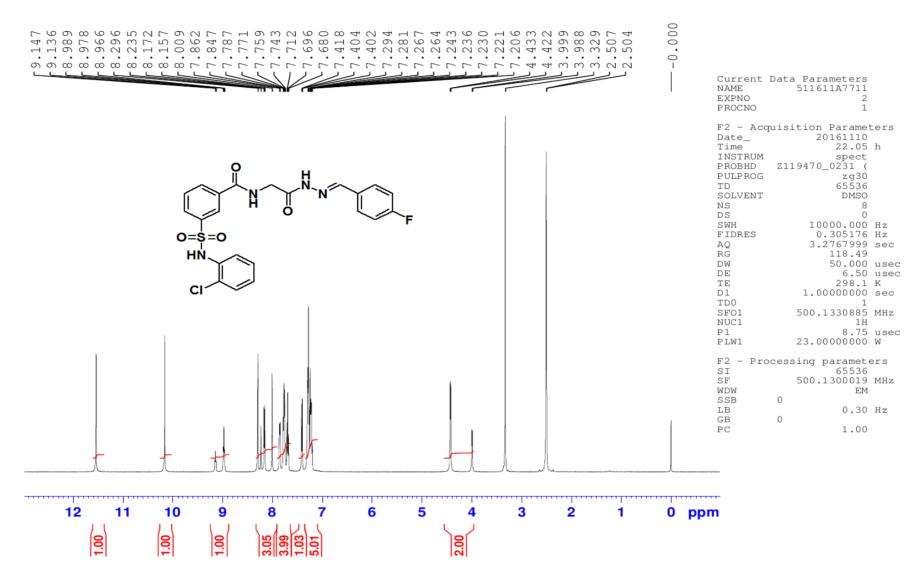
Elements Used:

C: 0-23 H: 0-19 N: 0-4 O: 0-4 F: 0-1 S: 0-1 CI: 0-1

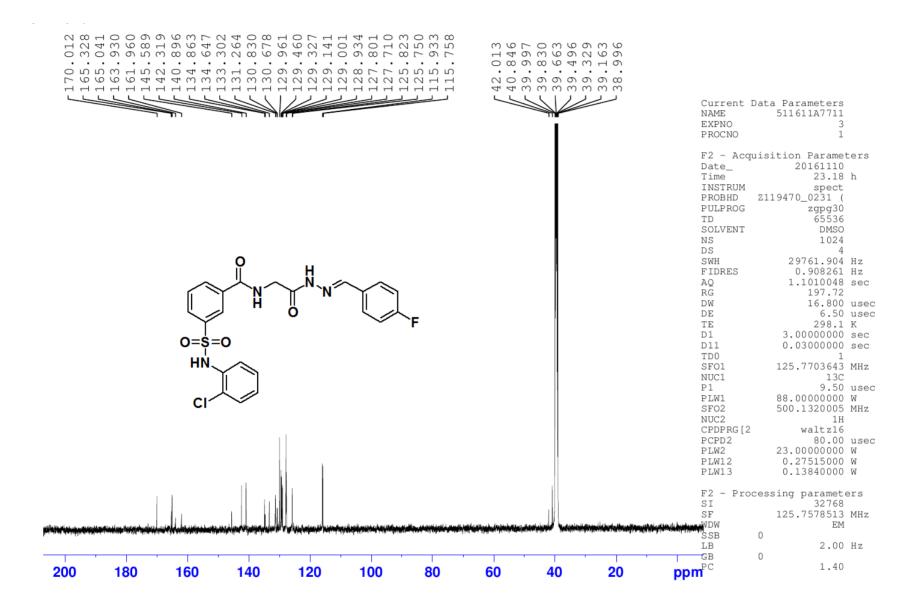
GVB-SA07

511611B4321 23 (0.341) AM (Top,4, Ar,5000.0,195.14,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00) 1: TOF MS ES+ 489.0807 3.86e3 100-491.0713 491.1785 511.0580 527.9943 294.0344 472.0714488.6638 323.0499 364.0981 435.1055 392.0964 340 360 400 420 500 300 320 380 460 480 Minimum: -1.55.0 Maximum: 10000.0 50.0 Mass Calc. Mass mDa PPM DBE i-FIT Formula 489.0807 489.0800 0.7 1.4 14.5 225054.8 C22 H19 N4 O4 F S C1

HRMS of Compound-1a



¹H NMR of **Compound-1b**



¹³CNMR of **Compound-1b** in DMSO-d₆

Page 1

Single Mass Analysis

Tolerance = 10000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

190 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

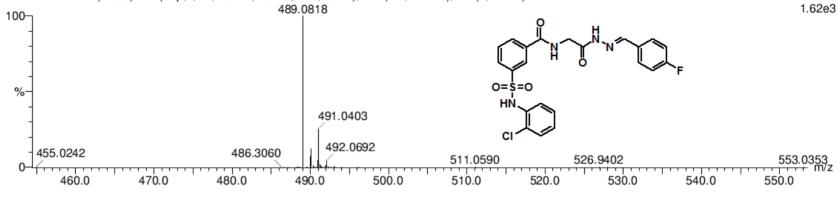
Elements Used:

C: 0-22 H: 0-19 N: 0-4 O: 0-4 F: 0-1 S: 0-1 CI: 0-1

GVB-SA01

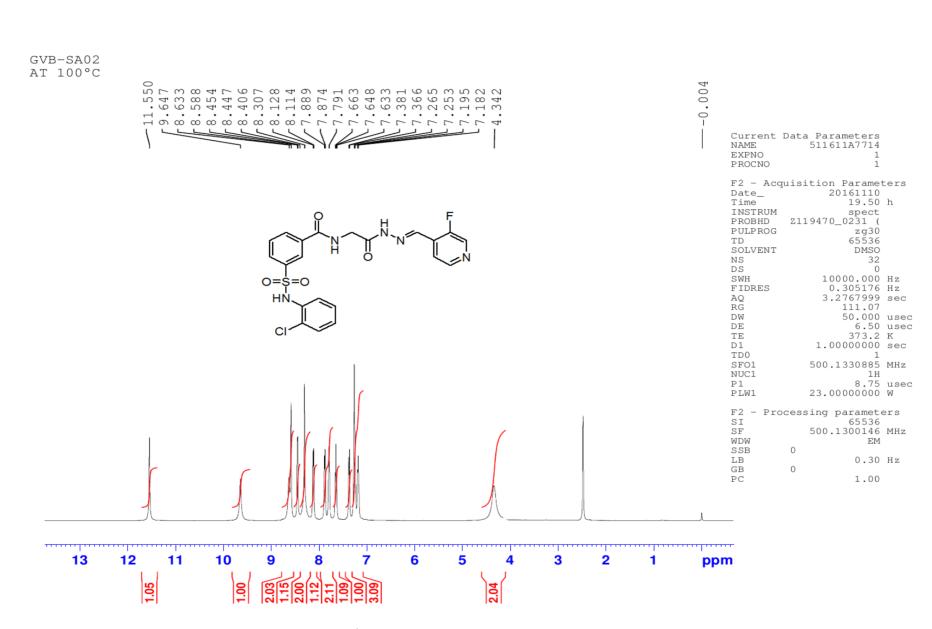
511611B4315 25 (0.360) AM (Top,4, Ar,5000.0,195.16,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00)

1: TOF MS ES+

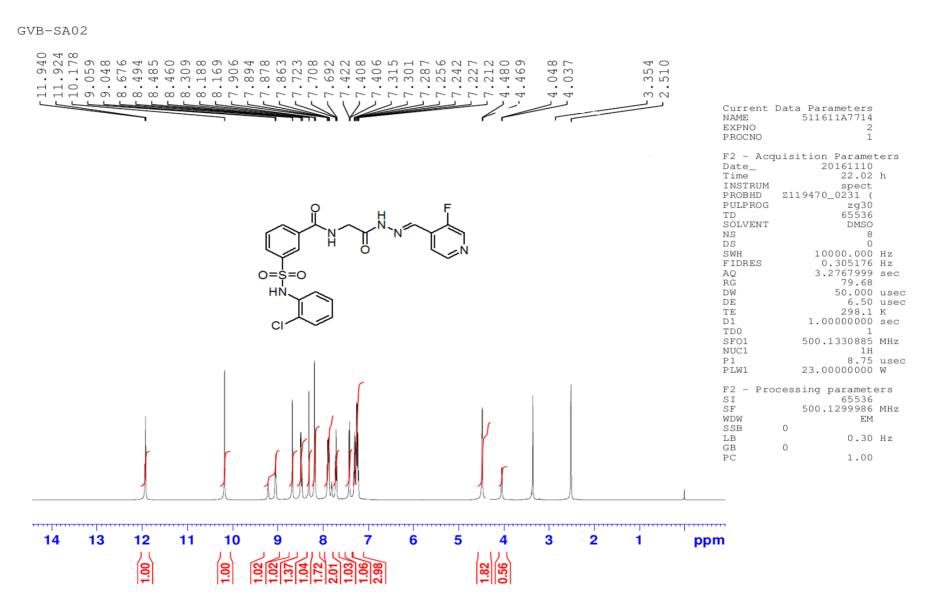


Minimum: -1.55.0 Maximum: 10000.0 50.0 Calc. Mass DBE i-FIT Mass mDa PPM Formula 489.0818 489.0800 1.8 3.7 14.5 94405.7 C22 H19 N4 O4 F S C1

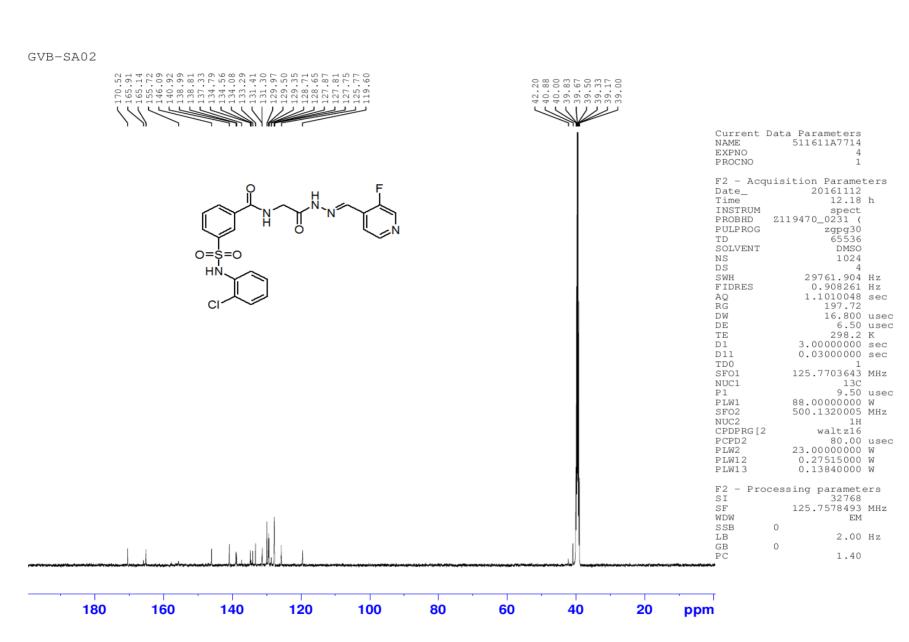
HRMS of Compound-1b



¹H NMR of Compound-1c at 100°C



¹H NMR of **Compound-1c**



 $^{13}\text{CNMR}$ of **Compound-1c** in DMSO-d₆

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1: TOF MS ES+

Single Mass Analysis

Tolerance = 10000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

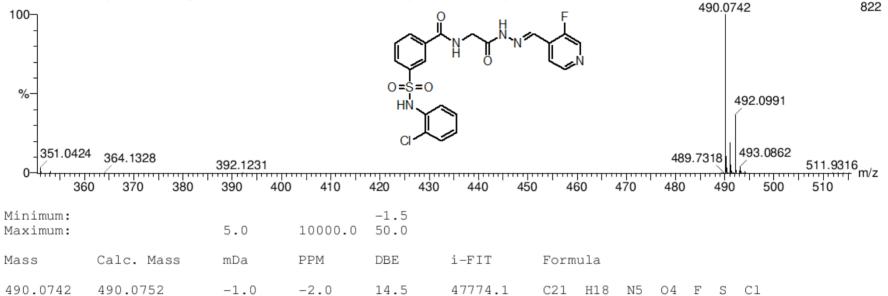
230 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

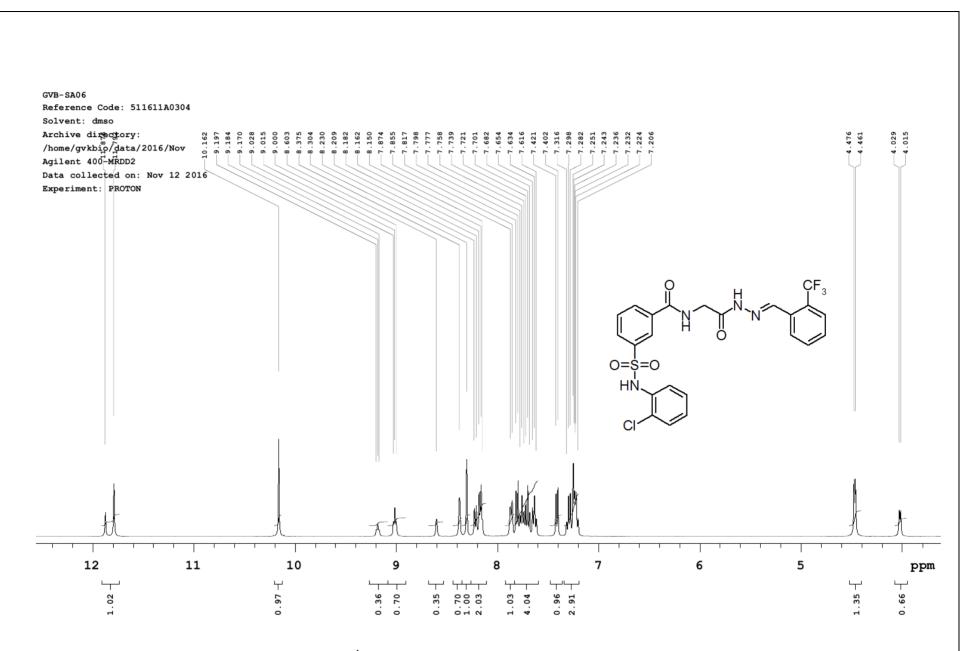
C: 0-21 H: 0-18 N: 0-5 O: 0-4 F: 0-1 S: 0-1 CI: 0-1

GVB-SA02

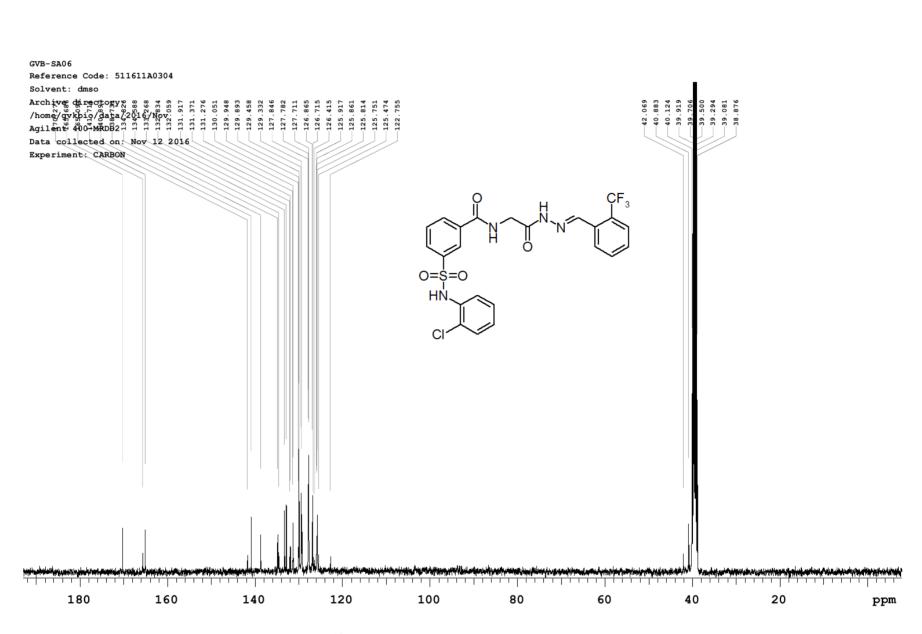
511611B4316 23 (0.343) AM (Top,4, Ar,5000.0,195.16,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00)



HRMS of Compound-1c



 $^1\mbox{H}$ NMR of Compound-1d in DMSO-d $_6$



 13 CNMR of **Compound-1d** in DMSO-d₆

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Single Mass Analysis

Tolerance = 10000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

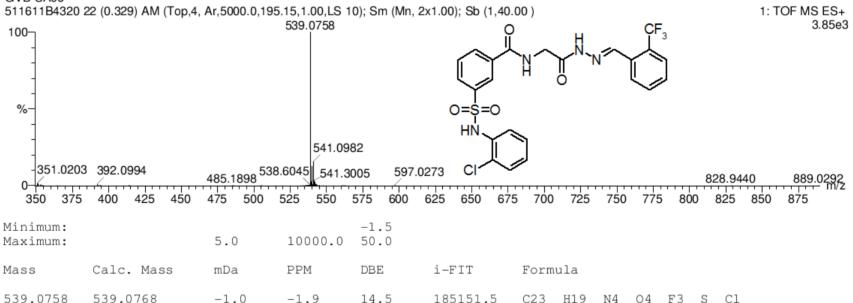
Monoisotopic Mass, Even Electron Ions

389 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

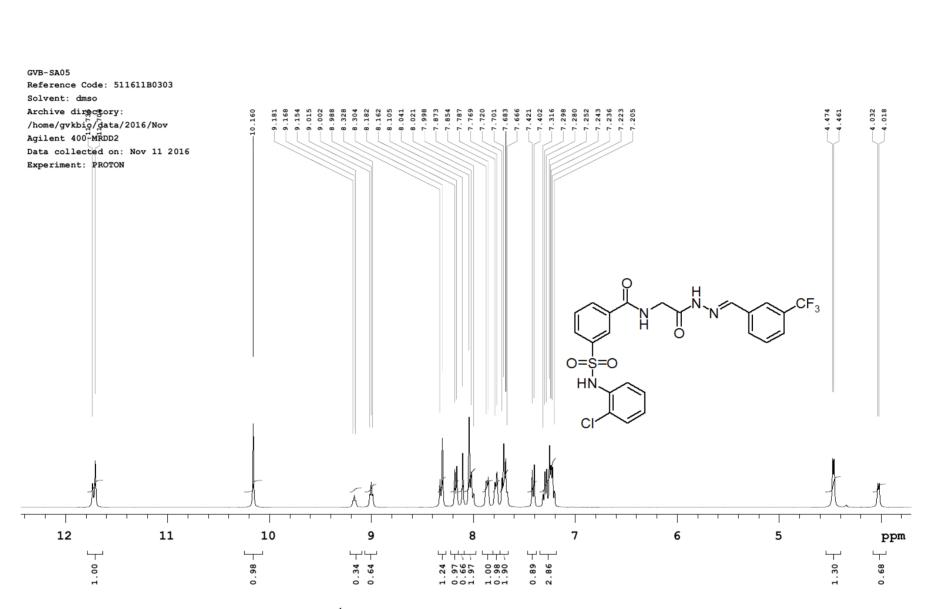
Elements Used:

C: 0-23 H: 0-19 N: 0-4 O: 0-4 F: 0-3 S: 0-1 CI: 0-1

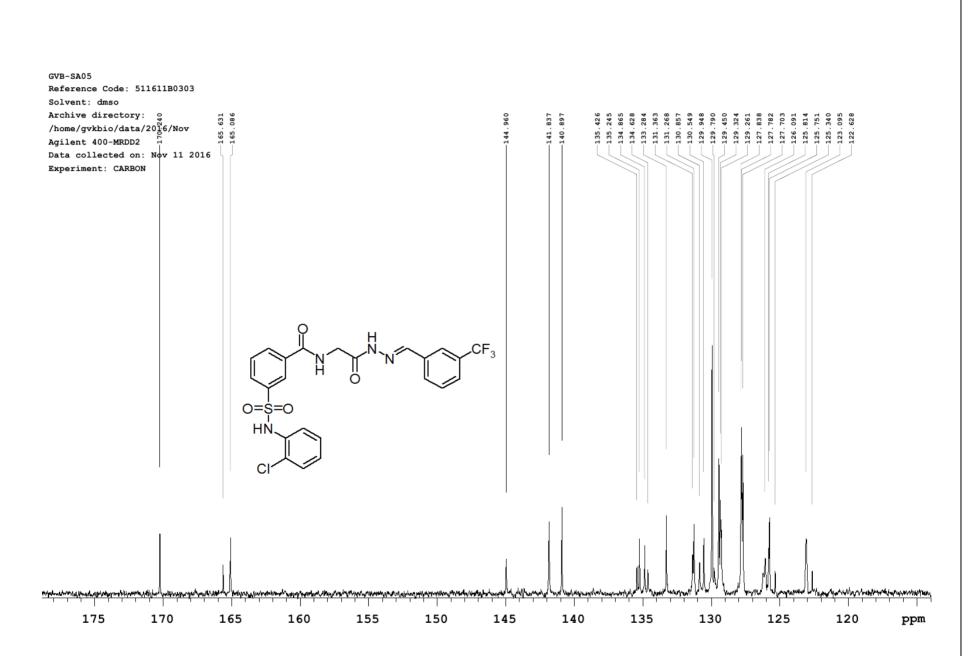
GVB-SA06



HRMS of Compound-1d



¹H NMR of **Compound-1e** in DMSO-d₆



 $^{13}CNMR$ of ${\color{red}{\bf Compound-1e}}$ in DMSO-d $_6$

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Single Mass Analysis

Tolerance = 10000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

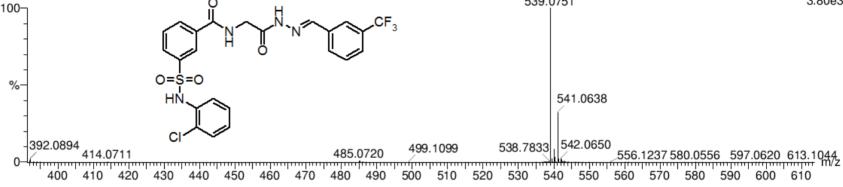
Monoisotopic Mass, Even Electron Ions

389 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

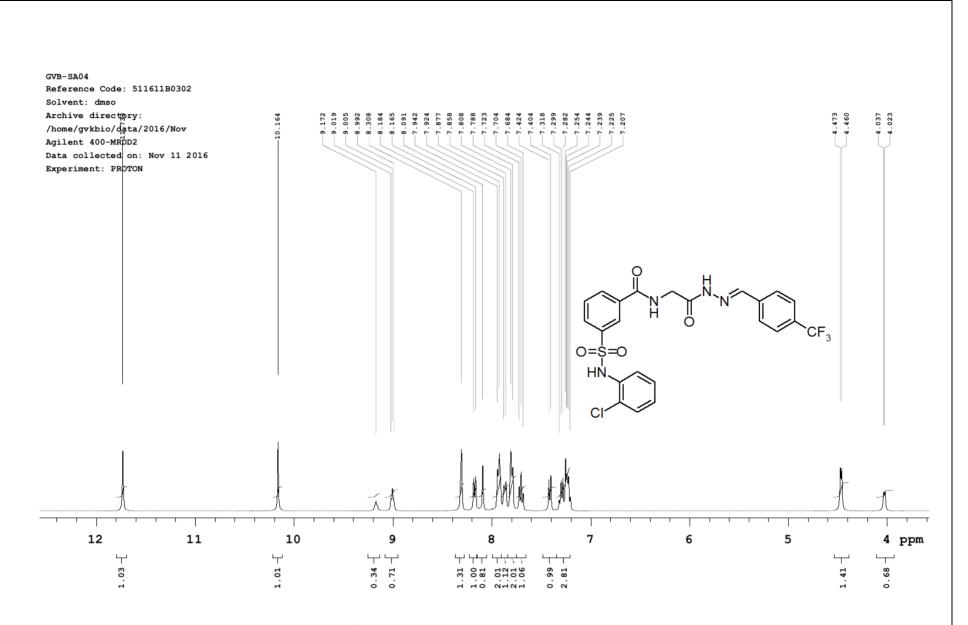
C: 0-23 H: 0-19 N: 0-4 O: 0-4 F: 0-3 S: 0-1 CI: 0-1

GVB-SA05

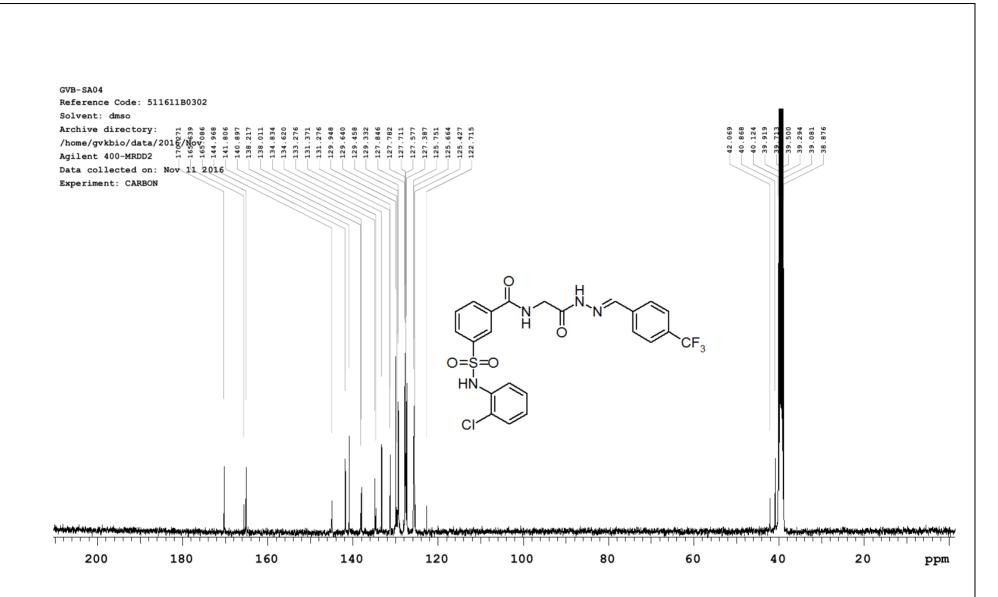


Minimum: -1.55.0 Maximum: 10000.0 50.0 PPM DBE i-FIT Calc. Mass Formula Mass mDa 539.0751 539.0768 -1.7-3.214.5 182666.4 C23 H19 N4 O4 F3 S C1

HRMS of Compound-1e



¹H NMR of **Compound-1f** in DMSO-d₆



 $^{13}\mbox{CNMR}$ of $\mbox{Compound-1f}$ in DMSO-d $_6$

Page 1

Single Mass Analysis

Tolerance = 10000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

389 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

C: 0-23 H: 0-19 N: 0-4 O: 0-4 S: 0-1 CI: 0-1 F: 0-3

GVB-SA04

511611B4318 23 (0.339) AM (Top,4, Ar,5000.0,195.17,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00)

1: TOF MS ES+

539.0767

776

477.1833 485.1155 492.0615

538.8185 541.2565

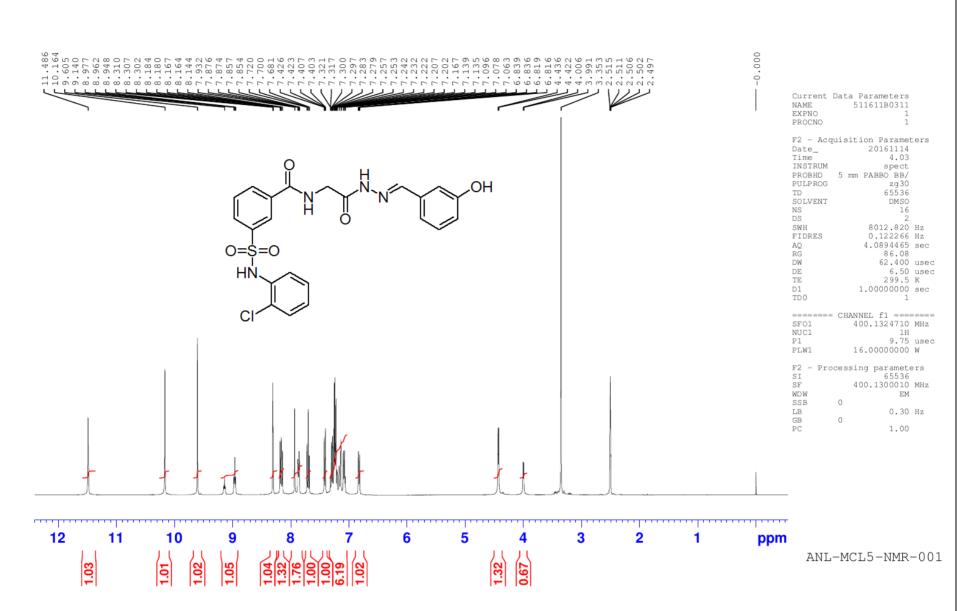
500.9357

m/z

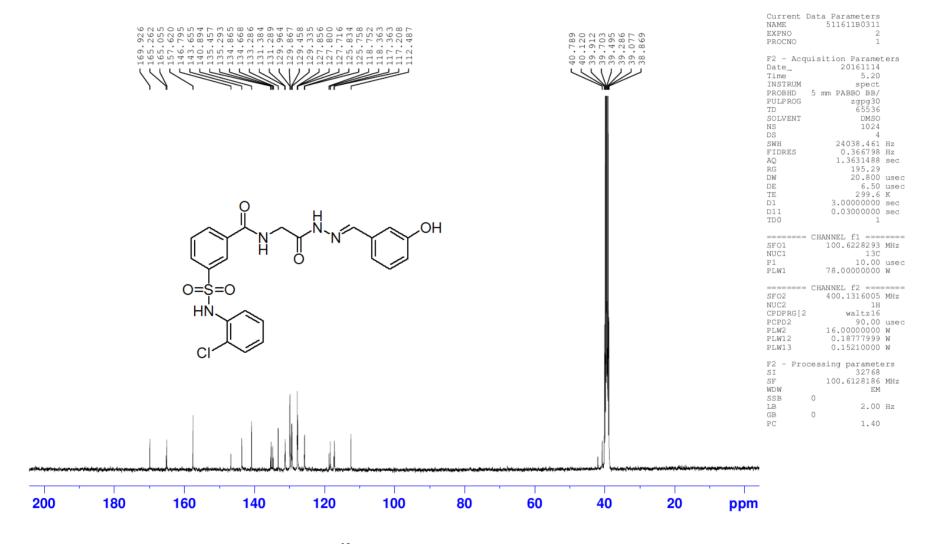
350 360 370 380 390 400 410 420 430 440 450 460 470 480 490 500 510 520 530 540 550 560 570

Minimum: -1.5Maximum: 5.0 10000.0 50.0 Mass Calc. Mass mDa PPM DBE i-FIT Formula -0.1 -0.2 14.5 539.0767 539.0768 37263.6 C23 H19 N4 O4 S C1 F3

HRMS of Compound-1f



¹H NMR of **Compound-1g** in DMSO-d₆



 13 CNMR of **Compound-1g** in DMSO- d_6

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1: TOF MS ES+

Single Mass Analysis

Tolerance = 10000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

106 formula(e) evaluated with 2 results within limits (up to 1 closest results for each mass)

Elements Used:

C: 0-23 H: 0-20 N: 0-4 O: 0-5 S: 0-1 CI: 0-1

390

400

410

GVB-SA08

511611B4322 23 (0.341) AM (Top,4, Ar,5000.0,195.18,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00) 487.0821 2.85e3 100-489.0794 490.0956 503.0338 362.0693 423.0471 486.2502 433.2201 545.0589 myz 454, 1656

450

460

470

510

520

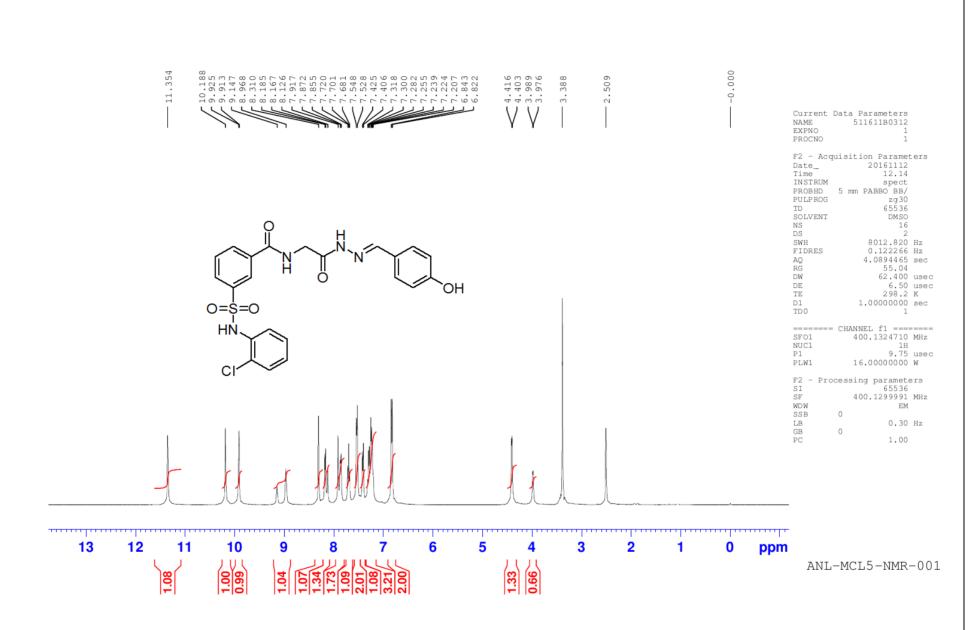
530

Minimum: -1.55.0 50.0 Maximum: 10000.0 Calc. Mass PPM i-FIT Mass mDa DBE Formula -2.2 487.0821 -4.5 14.5 487.0843 168119.6 C22 H20 N4 O5 S C1

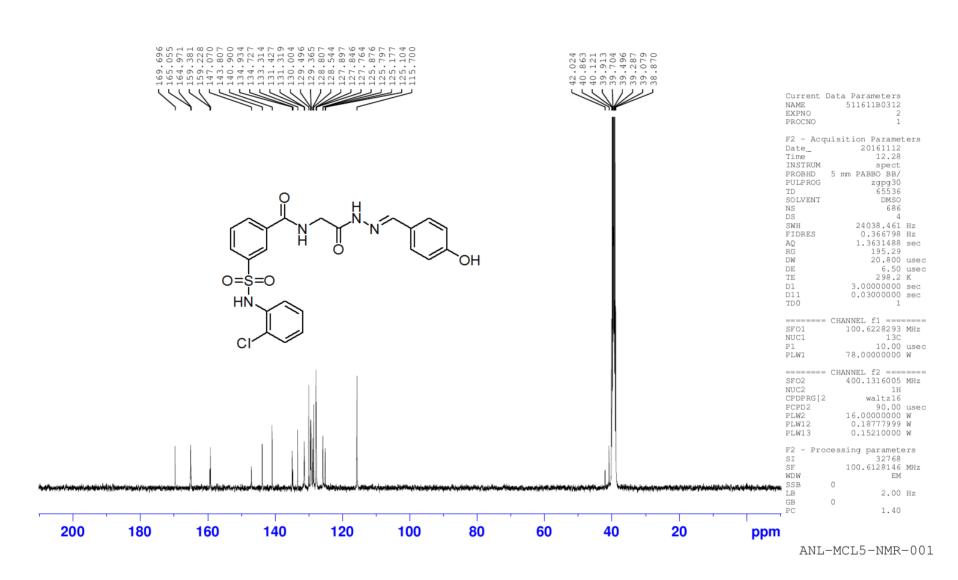
430

440

HRMS of Compound-1g



¹H NMR of **Compound-1h** in DMSO-d₆



¹³CNMR of **Compound-1h** in DMSO-d₆

Page 1

Single Mass Analysis

Tolerance = 10000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

106 formula(e) evaluated with 2 results within limits (up to 1 closest results for each mass)

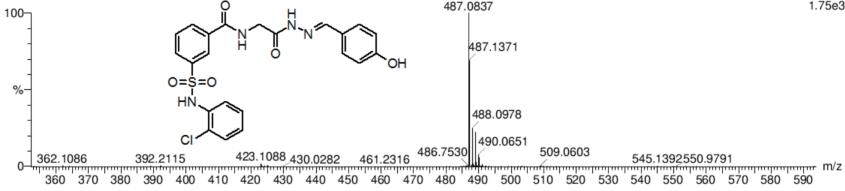
Elements Used:

C: 0-23 H: 0-20 N: 0-4 O: 0-5 S: 0-1 CI: 0-1

GVB-SA09

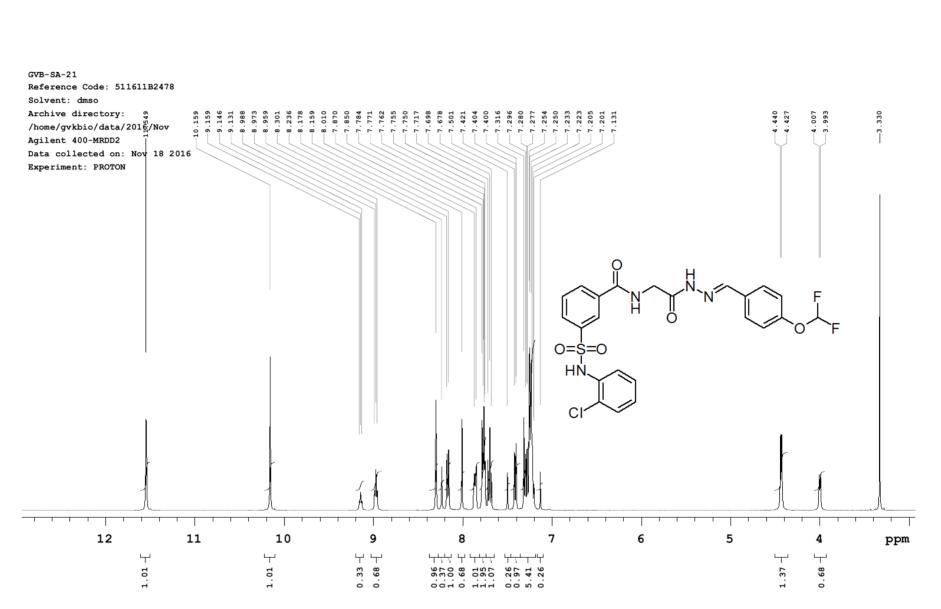
 $511611B4323\ 21\ (0.287)\ AM\ (Top,4,\ Ar,5000.0,195.16,1.00,LS\ 10);\ Sm\ (Mn,\ 2x1.00);\ Sb\ (1,40.00\);\ Cm\ (15:22)$

1: TOF MS ES+ 1.75e3

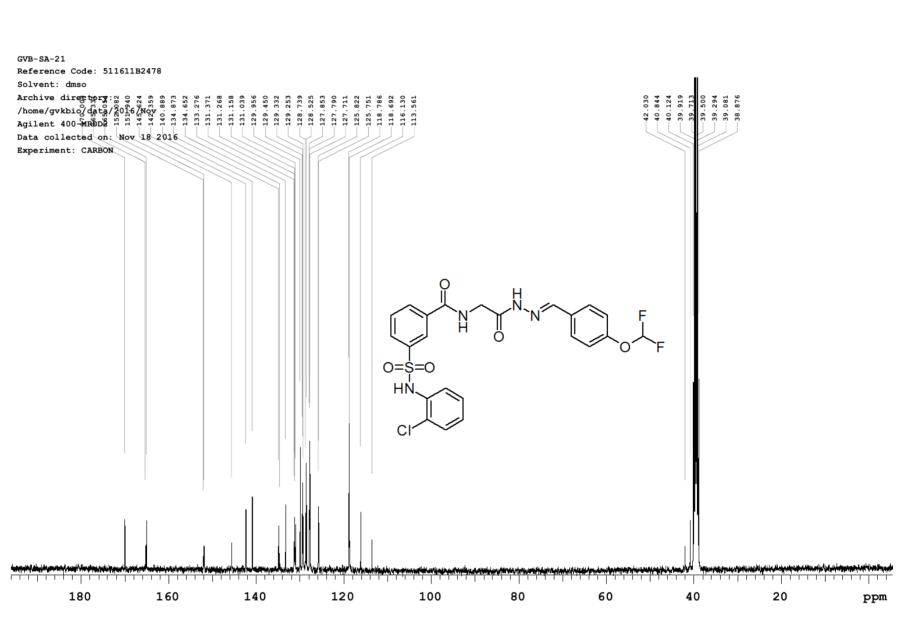


-1.5Minimum: Maximum: 5.0 10000.0 50.0 Calc. Mass Mass mDa PPM DBE i-FIT Formula -0.6 487.0837 487.0843 -1.214.5 102982.1 C22 H20 N4 O5 S C1

HRMS of Compound-1h



 $^1\mbox{H}$ NMR of $\mbox{\bf Compound-1i}$ in DMSO-d $_6$



 $^{13}\mbox{CNMR}$ of $\mbox{Compound-1i}$ in DMSO-d $_6$

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Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

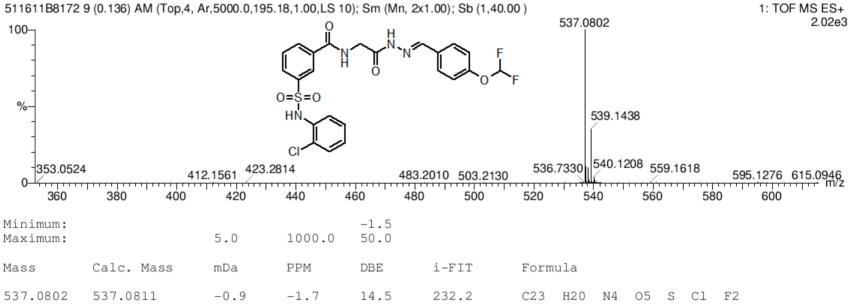
Monoisotopic Mass, Even Electron Ions

335 formula(e) evaluated with 2 results within limits (up to 1 best isotopic matches for each mass)

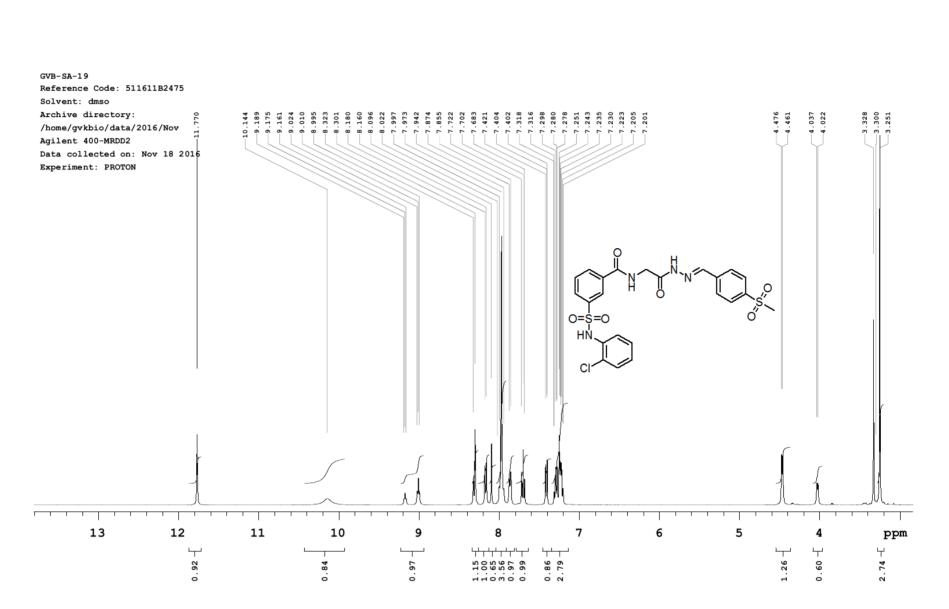
Elements Used:

C: 0-24 H: 0-20 N: 0-4 O: 0-5 S: 0-1 CI: 0-1 F: 0-2

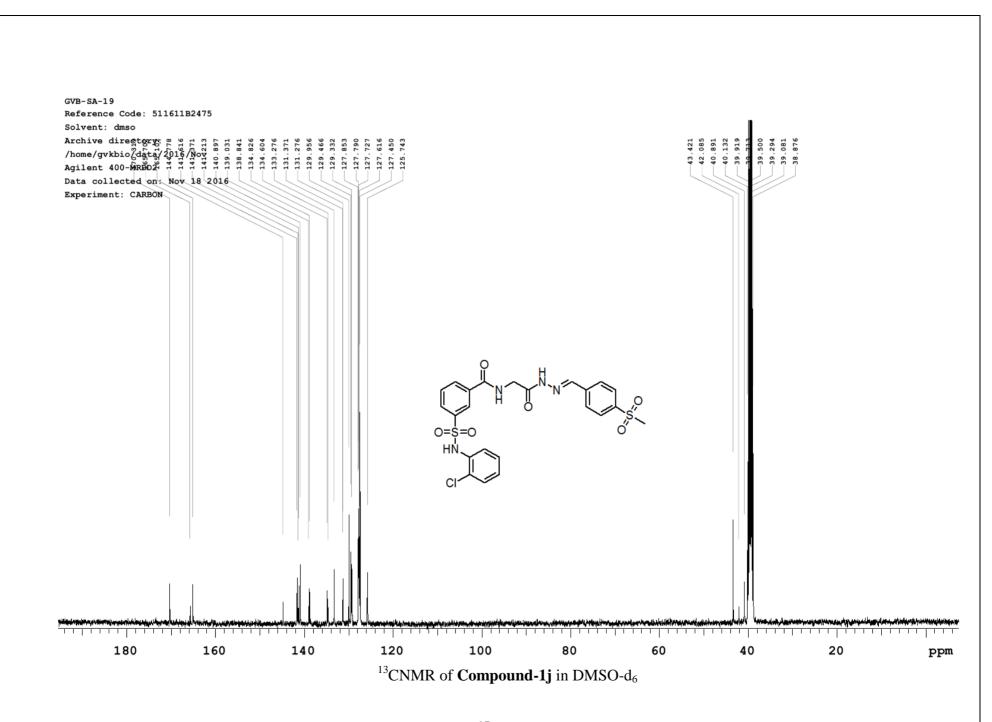
GVB-SA21



HRMS of Compound-1i



 $^{1}\text{H NMR of }\text{Compound-1j}\text{ in DMSO-d}_{6}$



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Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

203 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass)

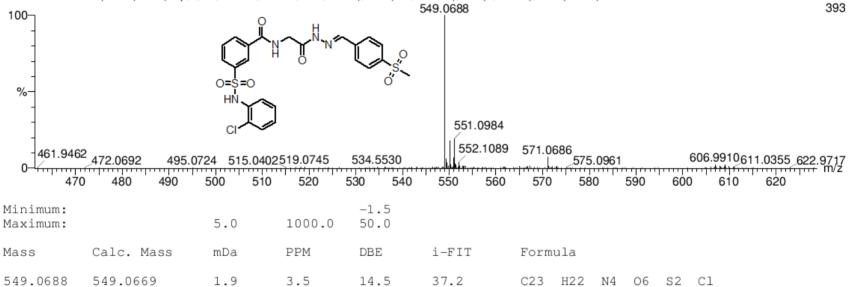
Elements Used:

C: 0-23 H: 0-22 N: 0-4 O: 0-6 S: 0-2 CI: 0-1

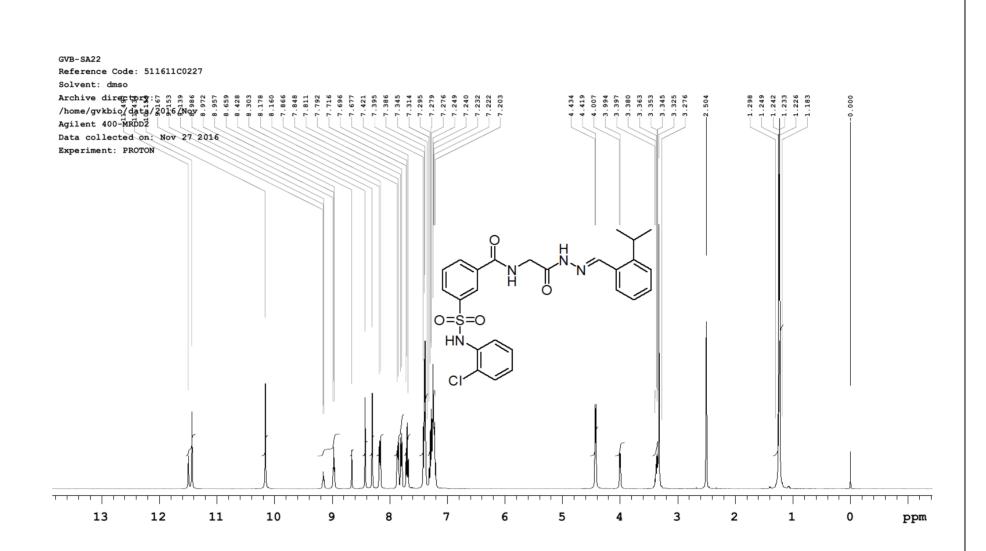
GVB-SA19

511611B8169 56 (0.800) AM (Top,4, Ar,5000.0,195.15,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00); Cm (28:56)

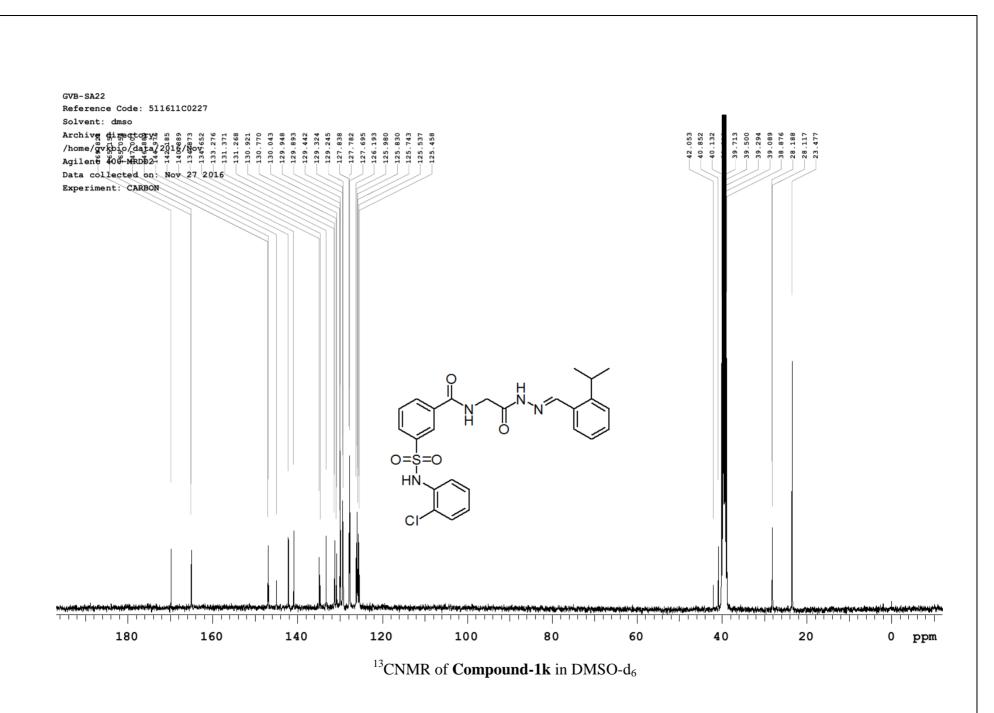
1: TOF MS ES+



HRMS of Compound-1j



 $^1\mbox{H}$ NMR of $\mbox{\bf Compound-1k}$ in DMSO-d $_6$



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Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

93 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass)

Elements Used:

C: 0-25 H: 0-26 N: 0-4 O: 0-4 S: 0-1 CI: 0-1

GVB-SA22

511611B8551 8 (0.127) AM (Top,4, Ar,5000.0,195.16,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00) 1: TOF MS ES+ 4.75e3 100-% 514.1898 516.1650 571.1795 591.0963 241.2198273.1116 351.0653 388.1931 459.1898 653.1914 726.1837 M/z 350 375 425 725 300 450 500 550 575 600 650 Minimum: -1.5Maximum: 5.0 1000.0 50.0 Mass Calc. Mass mDa PPM DBE i-FIT Formula 1.6 513.1379 513.1363 3.1 14.5 525.4 C25 H26 N4 O4 S C1

HRMS of Compound-1k

GVB-SA-20

Reference Code: 511611B2476

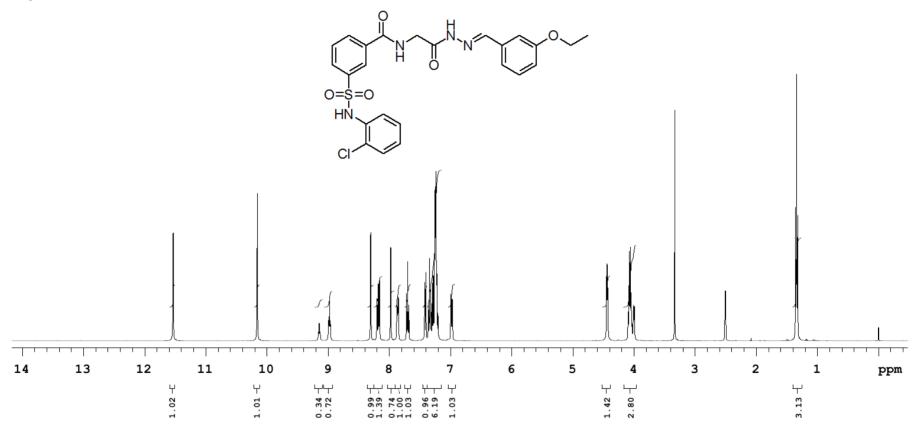
Solvent: dmso
Archive directory:

/home/gvkbio/data/2016/Nov

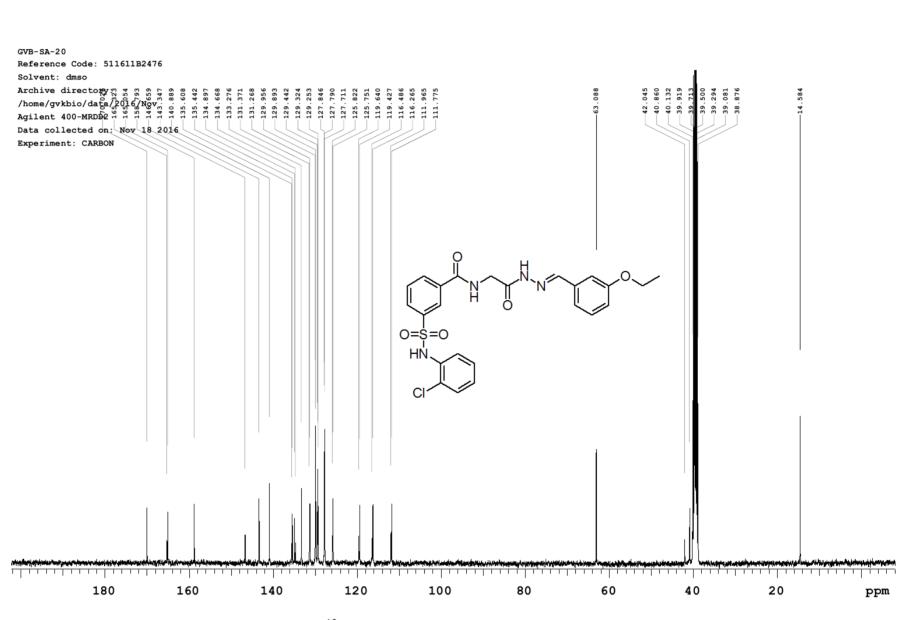
Agilent 400-MRDD2

Data collected on: Nov 18 2016

Experiment: PROTON



 $^{1}\text{H NMR}$ of **Compound-11** in DMSO-d₆



 $^{13}\mbox{CNMR}$ of $\mbox{Compound-11}$ in DMSO-d $_6$

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Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

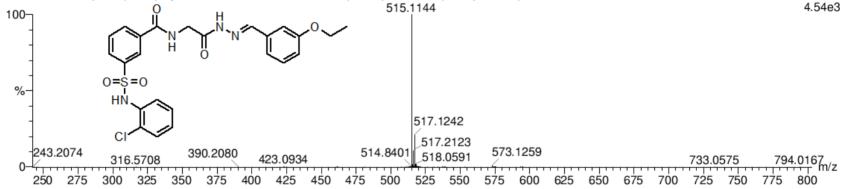
Monoisotopic Mass, Even Electron Ions

161 formula(e) evaluated with 7 results within limits (up to 1 best isotopic matches for each mass) Elements Used:

C: 0-24 H: 0-24 N: 0-4 O: 0-5 S: 0-2 CI: 0-1 GVB-SA20

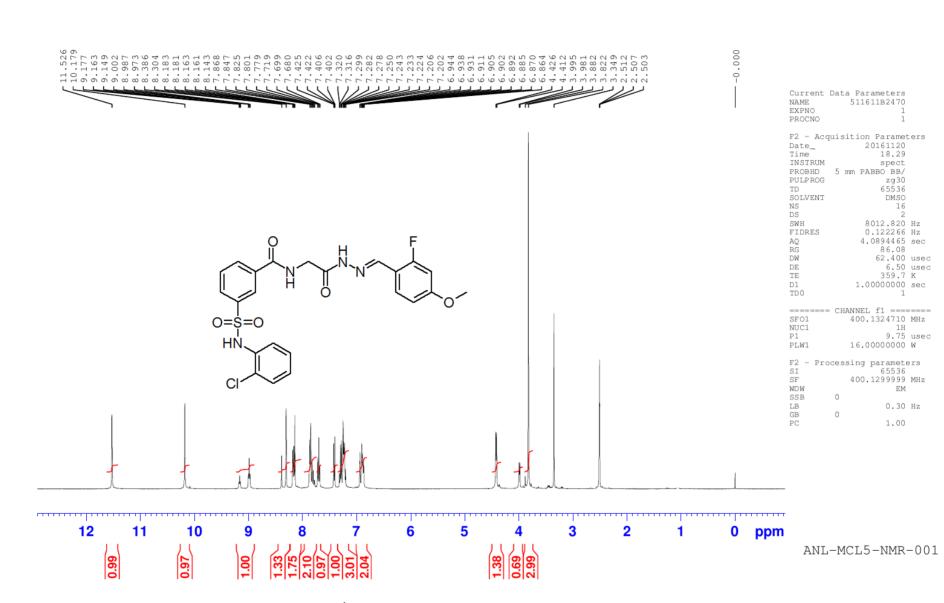
511611B8171 8 (0.126) AM (Top,4, Ar,5000.0,195.16,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00)

1: TOF MS ES+

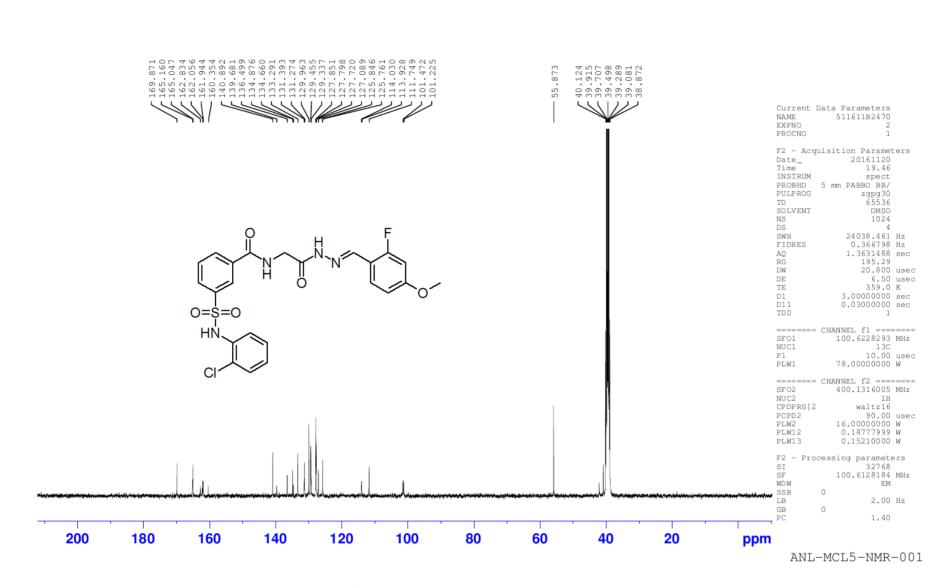


Minimum: -1.5Maximum: 5.0 1000.0 50.0 Mass Calc. Mass mDa PPM DBE i-FIT Formula 515.1144 515.1156 -1.2-2.3 14.5 481.0 C24 H24 N4 O5 S C1

HRMS of Compound-11



¹H NMR of **Compound-1m** in DMSO-d₆



¹³CNMR of **Compound-1m** in DMSO-d₆

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Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

-0.2

-0.4

Selected filters: None

Monoisotopic Mass, Even Electron Ions

519.0905

230 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass)

Elements Used:

C: 0-23 H: 0-21 N: 0-4 O: 0-5 F: 0-1 S: 0-1 CI: 0-1

GVB-SA16

519.0903

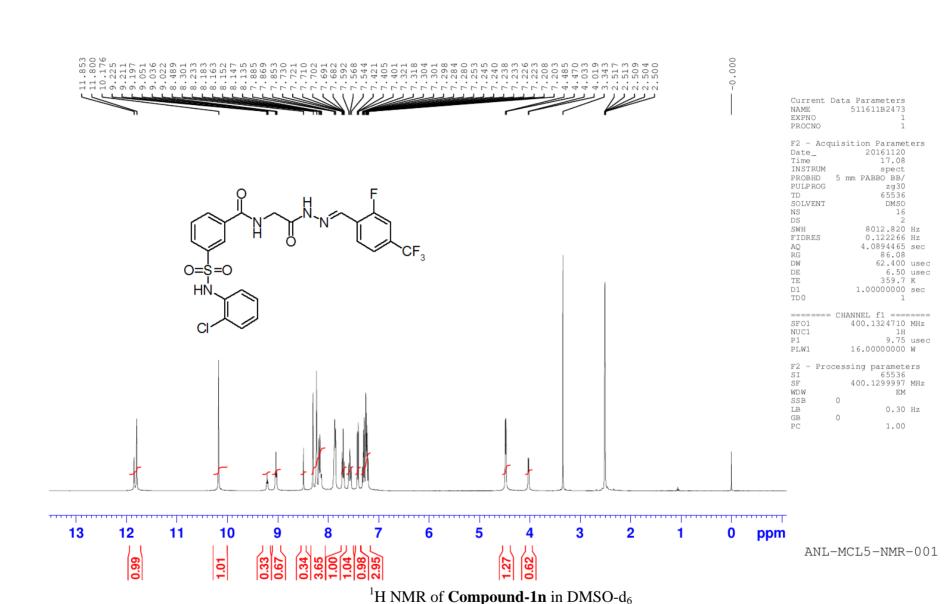
511611B8166 14 (0.186) AM (Top,4, Ar,5000.0,195.17,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00) 1: TOF MS ES+ 519.0903 4.20e3 100-521.1299 518.2645 522.0795 465.2796 577.2268 540 560 580 600 620 640 660 680 700 720 740 760 780 500 520 460 480 -1.5Minimum: Maximum: 5.0 1000.0 50.0 Mass Calc. Mass mDa PPM DBE i-FIT Formula

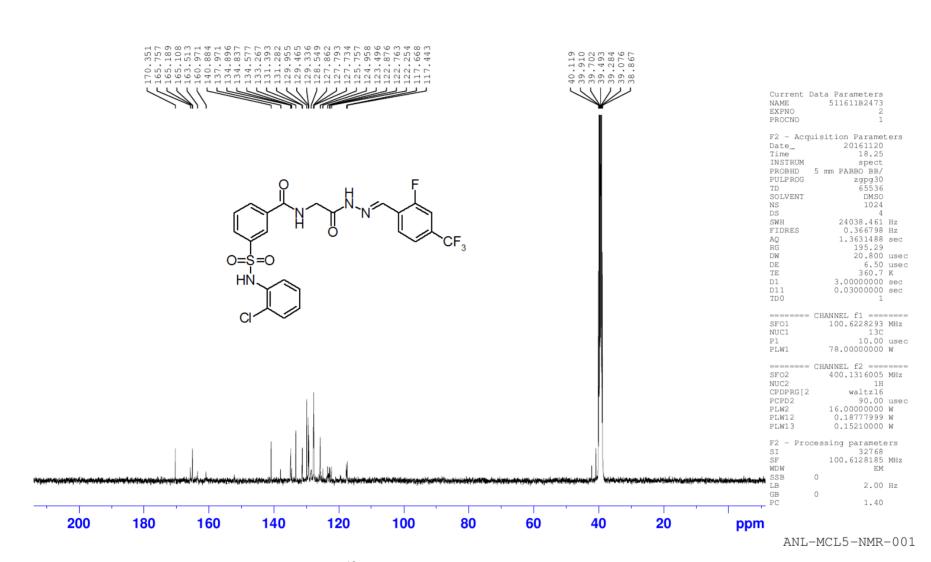
174.9

C23 H21 N4 O5 F S C1

HRMS of Compound-1m

14.5





 13 CNMR of **Compound-1n** in DMSO-d₆

Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

489 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass) Elements Used:

C: 0-23 H: 0-18 N: 0-4 O: 0-4 F: 0-4 S: 0-1 CI: 0-1

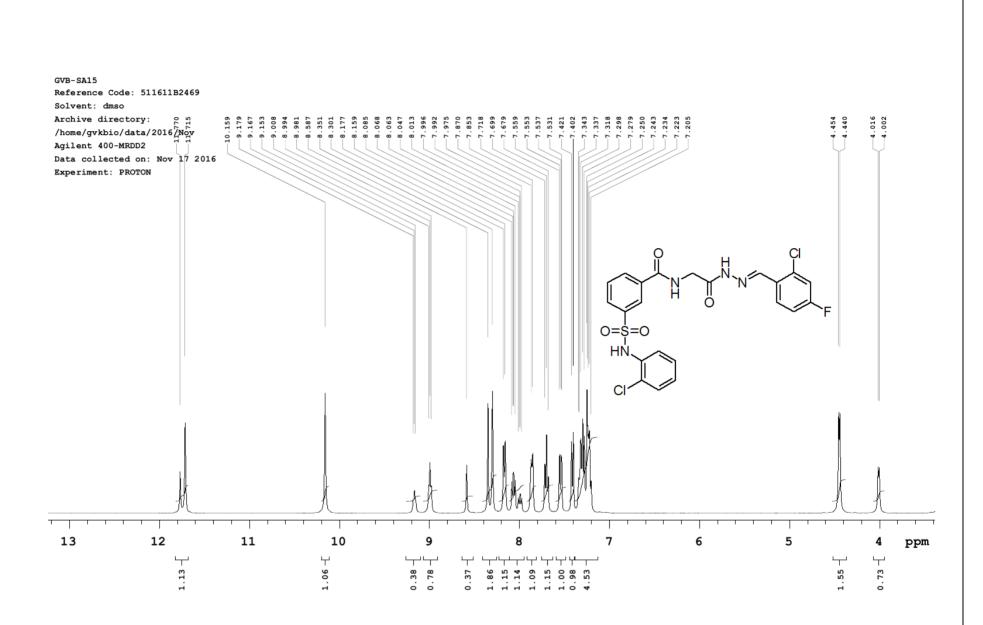
GVB-SA17

511611B8167 16 (0.238) AM (Top,4, Ar,5000.0,195.20,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00)

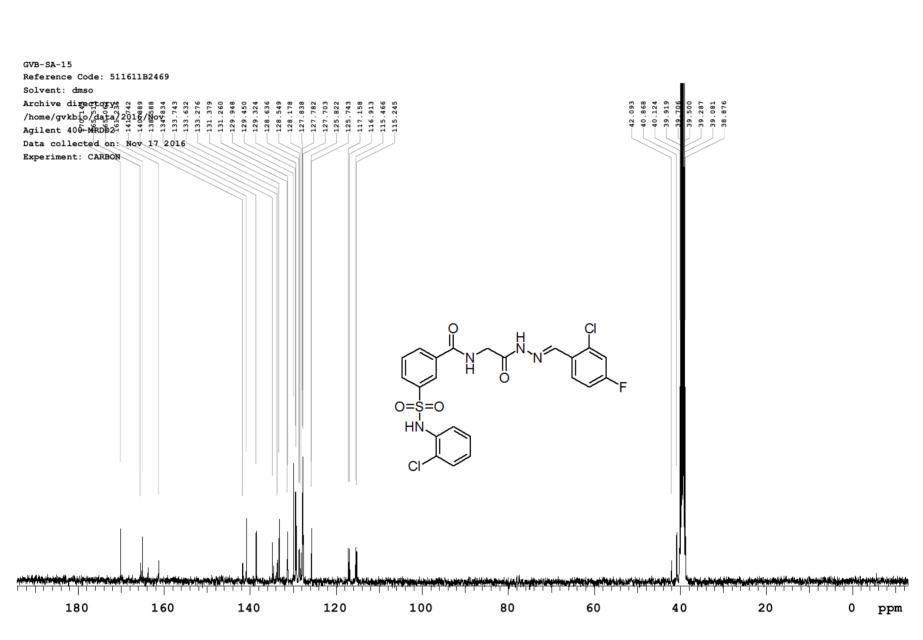
1: TOF MS ES+ 1.67e3 100-557.1711 559.0898 559.2841 351.0585 555.9849 635.0689 392.0527 432.0691 503.1351 615.1018 440 460 480 520 540 560 580 640 660 680 360 380 400 420 500 600 620

Minimum: Maximum:		5.0	1000.0	-1.5 50.0								
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Formula						
557.0684	557.0673	1.1	2.0	14.5	128.2	C23	H18	N4	04	F4	S	Cl

HRMS of Compound-1n



¹H NMR of **Compound-1o** in DMSO-d₆



 13 CNMR of **Compound-1o** in DMSO- d_6

Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

290 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass)

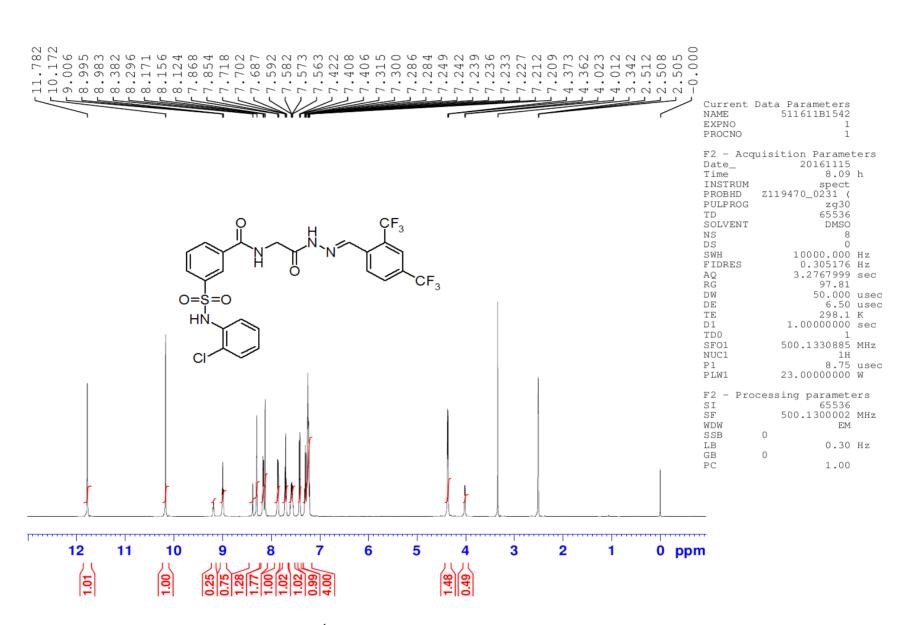
Elements Used:

C: 0-22 H: 0-18 N: 0-4 O: 0-4 F: 0-1 S: 0-1 CI: 0-2

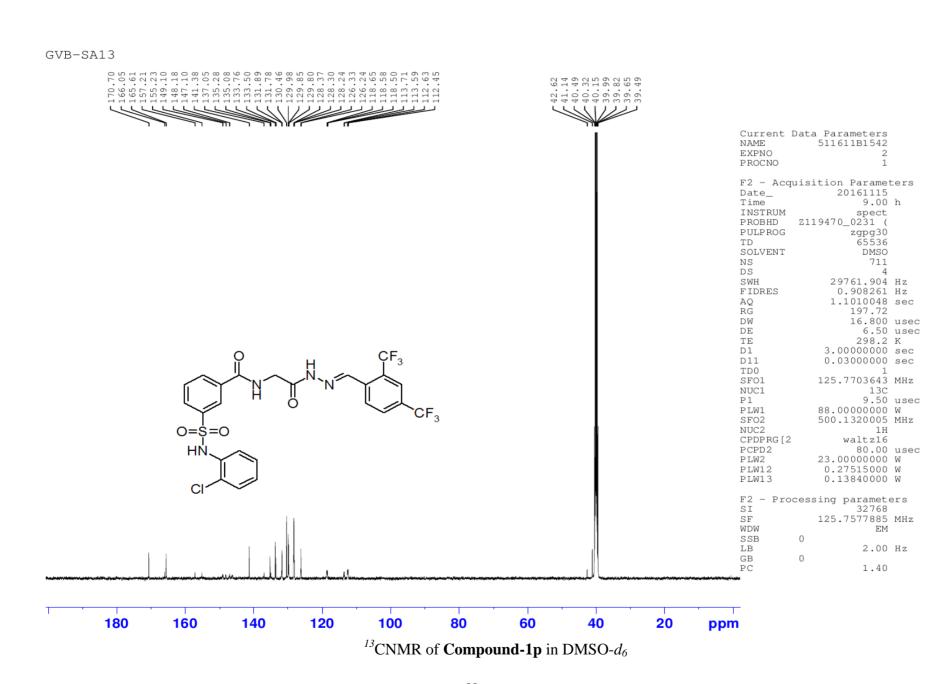
GVB-SA15

511611B8164 17 (0.251) AM (Top,4, Ar,5000.0,195.15,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00) 1: TOF MS ES+ 523.0408 1.37e3 100-525.0217 527.0397 477.1814 522.5323 545.0145 583.0936 601.0676 640.9149 m/z 480 490 500 510 520 530 540 550 560 570 580 590 600 610 620 630 640 Minimum: -1.55.0 Maximum: 1000.0 50.0 Calc. Mass PPM DBE i-FIT Formula Mass mDa 523.0408 523.0410 -0.2 -0.414.5 120.3 C22 H18 N4 O4 F S C12

HRMS of Compound-10



¹H NMR of **Compound-1p** in DMSO-d₆



Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

-1.9

-3.1

Selected filters: None

Monoisotopic Mass, Even Electron Ions

607.0641

689 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass)

Elements Used:

C: 0-24 H: 0-18 N: 0-4 O: 0-4 F: 0-6 S: 0-1 CI: 0-1

GVB-SA13

607.0622

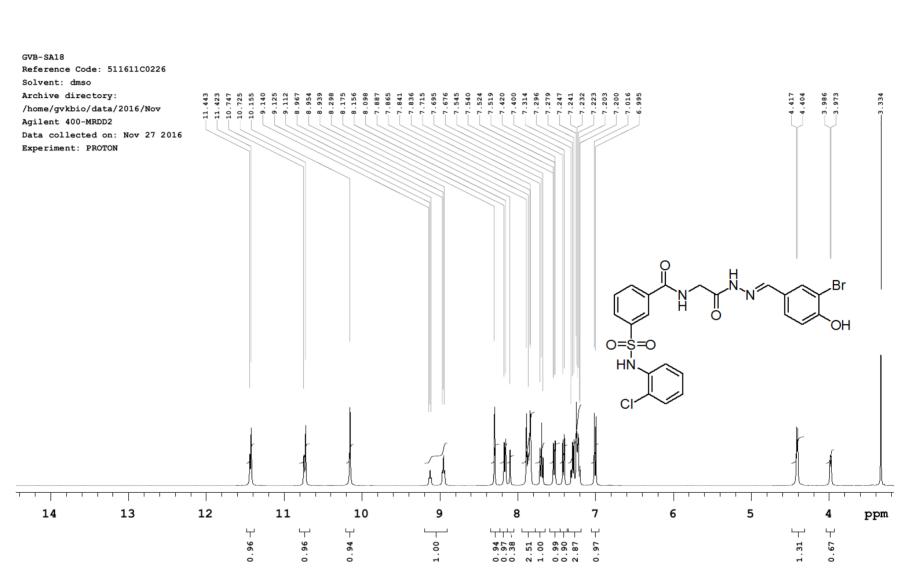
511611B8159 16 (0.238) AM (Top,4, Ar,5000.0,195.17,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00) 1: TOF MS ES+ 607.0622 795 100-607.2051 609.0174 606.7883 609.2559 554.2897 665.1365 624.3243 590 595 600 605 610 615 620 625 630 635 640 645 650 655 660 665 565 570 575 580 585 -1.5Minimum: 5.0 1000.0 50.0 Maximum: Mass Calc. Mass mDa PPM DBE i-FIT Formula

85.5

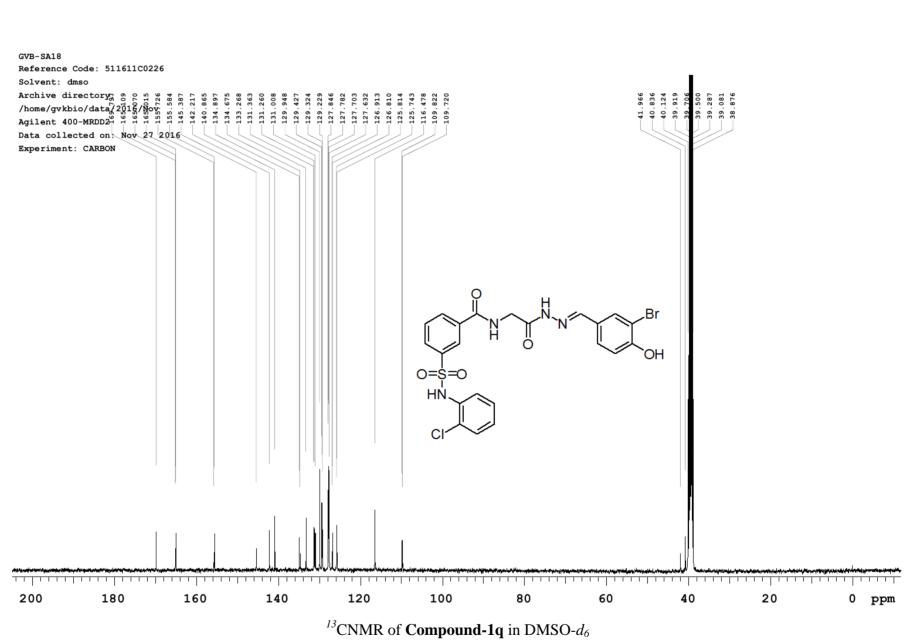
C24 H18 N4 O4 F6 S C1

HRMS of Compound-1p

14.5



 1 H NMR of **Compound-1q** in DMSO-d₆



Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

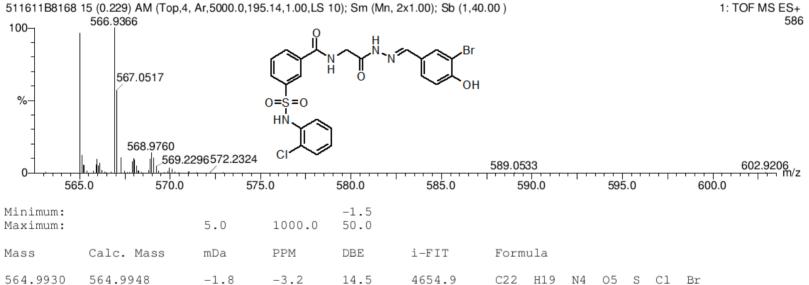
Monoisotopic Mass, Even Electron Ions

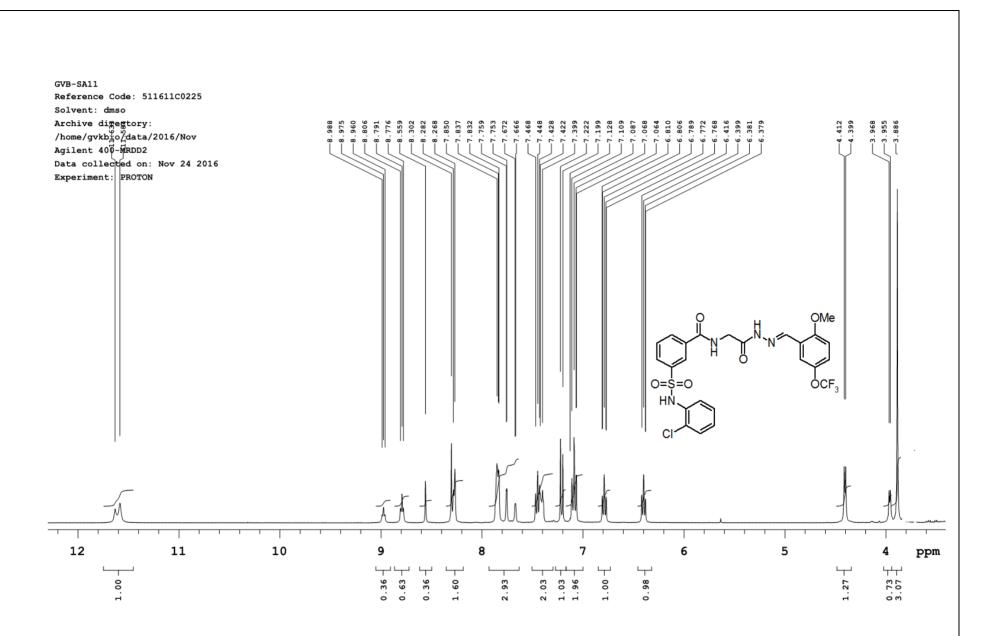
233 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass)

Elements Used:

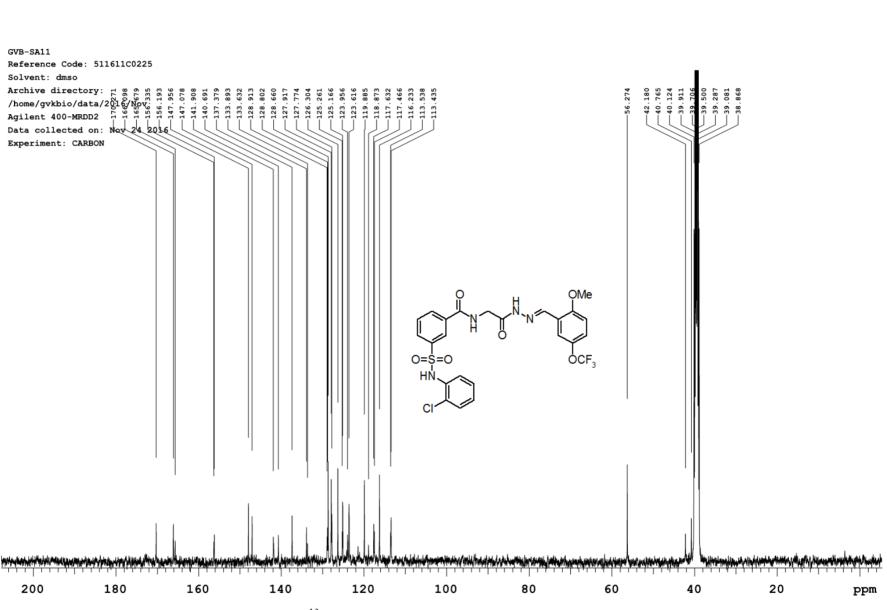
C: 0-22 H: 0-19 N: 0-4 O: 0-5 S: 0-1 CI: 0-1 Br: 0-1

GVB-SA18





¹H NMR of **Compound-1r** in DMSO-d₆



 13 CNMR of **Compound-1r** in DMSO- d_6

Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

549 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass)

Elements Used:

C: 0-24 H: 0-21 N: 0-4 O: 0-6 F: 0-3 S: 0-1 CI: 0-1

GVB-SA11

511611B8155 9 (0.118) AM (Top,4, Ar,5000.0,195.18,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00) 1: TOF MS ES+ 585.0807 1.06e3 QMe 100-587.0821 607.0095 609.1075 628.9656 651.0776 672.8693691.2649 493.2615 584.1223 500 520 540 560 580 600 620 640 660 700 720 740 760 Minimum: -1.55.0 1000.0 Maximum: 50.0 Calc. Mass PPM DBE i-FIT Mass mDa Formula 585.0807 585.0822 -1.5-2.6 14.5 54.1 C24 H21 N4 O6 F3 S C1

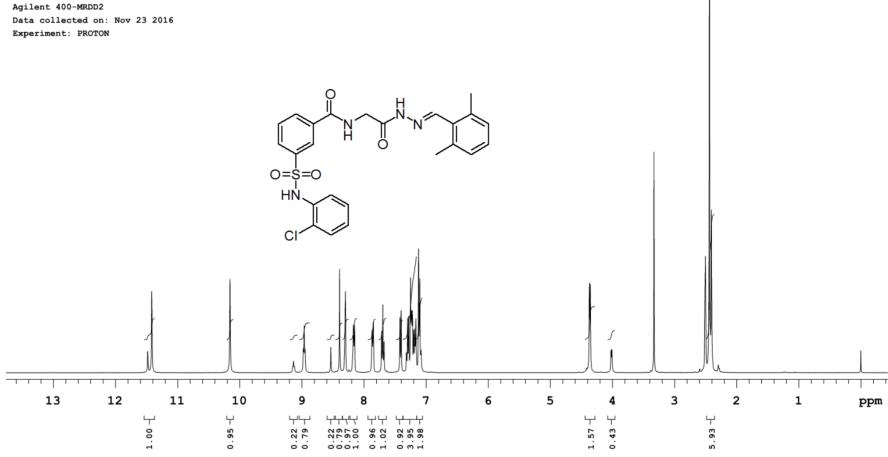
HRMS of Compound-1r

GVB-SA23

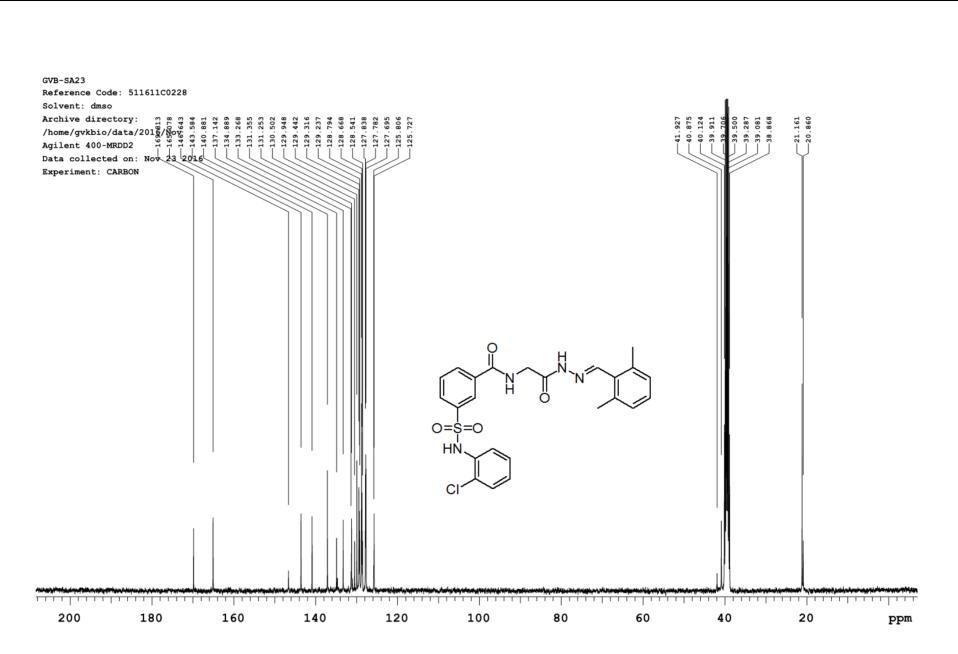
Reference Code: 511611C0228

Solvent: dmso Archive directory:

/home/gvkbio/data/2016/Nov



 $^{1}\text{H NMR of }\text{Compound-1s}\text{ in DMSO-d}_{6}$



 13 CNMR of **Compound-1s** in DMSO- d_6

Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

87 formula(e) evaluated with 2 results within limits (up to 1 best isotopic matches for each mass)

Elements Used:

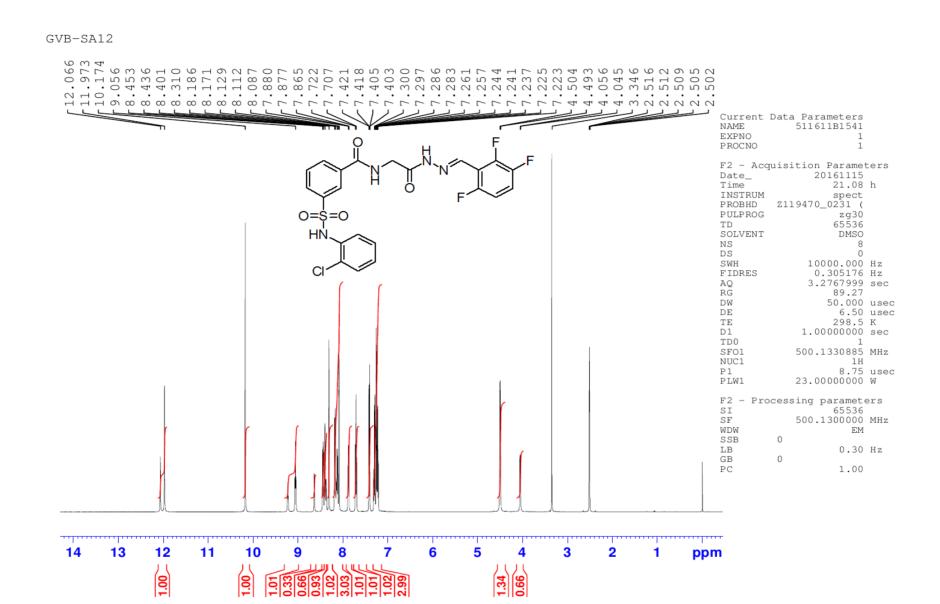
C: 0-25 H: 0-26 N: 0-4 O: 0-4 S: 0-1 CI: 0-1

GVB-SA23

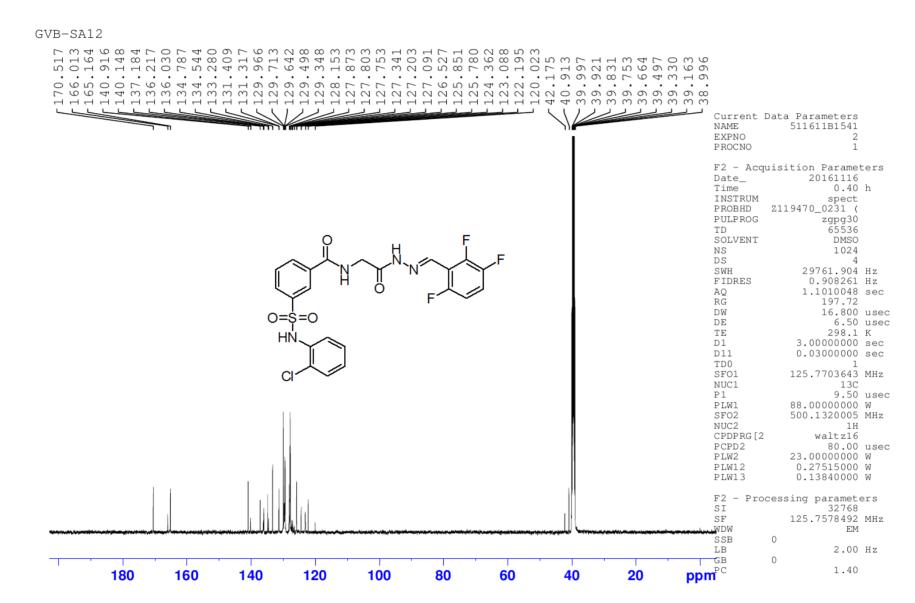
511611B8552 19 (0.268) AM (Top,4, Ar,5000.0,195.18,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00) 1: TOF MS ES+ 3.45e3 100-501.1338 501.2421 374.0804 498.4103 521.0458 538.9415 0-կուդրությունուրությունությու 410 430 440 450 470 490 510 520

Minimum: -1.55.0 Maximum: 1000.0 50.0 i-FIT Calc. Mass PPM Formula Mass mDa DBE 499.1229 499.1207 2.2 4.4 14.5 440.4 C24 H24 N4 O4 S C1

HRMS of Compound-1s



¹H NMR of **Compound-1t** in DMSO-d₆



 13 CNMR of **Compound-1t** in DMSO- d_6

Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

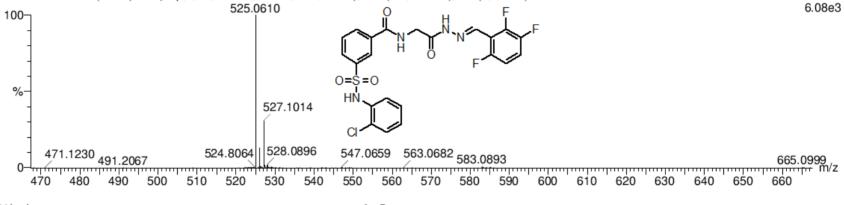
Monoisotopic Mass, Even Electron Ions

389 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass) Elements Used:

C: 0-22 H: 0-17 N: 0-4 O: 0-4 F: 0-3 S: 0-1 CI: 0-1 GVB-SA12

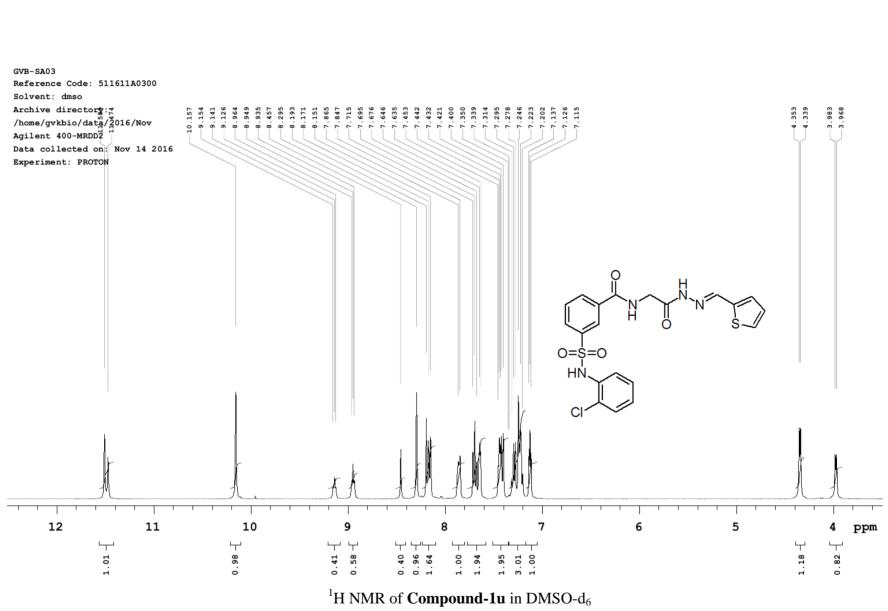
511611B8157 12 (0.164) AM (Top,4, Ar,5000.0,195.16,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00)

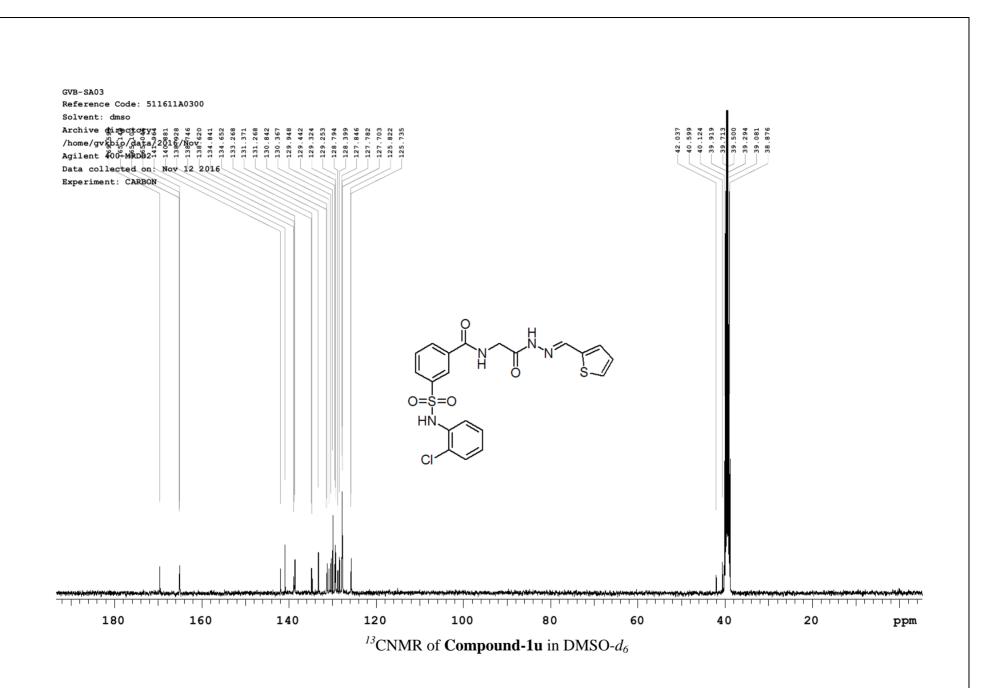
1: TOF MS ES+



Minimum: -1.5Maximum: 5.0 1000.0 50.0 Calc. Mass PPM DBE i-FIT Mass mDa Formula 525.0610 525.0611 -0.1 -0.2 14.5 349.2 C22 H17 N4 O4 F3 S C1

HRMS of Compound-1t





Page 1

Single Mass Analysis

Tolerance = 10000.0 PPM / DBE: min = -1.5, max = 50.0

-1.7

-3.6

Selected filters: None

Monoisotopic Mass, Even Electron Ions

477.0458

143 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

C: 0-20 H: 0-18 N: 0-4 O: 0-4 S: 0-2 CI: 0-1

GVB-SA03

477.0441

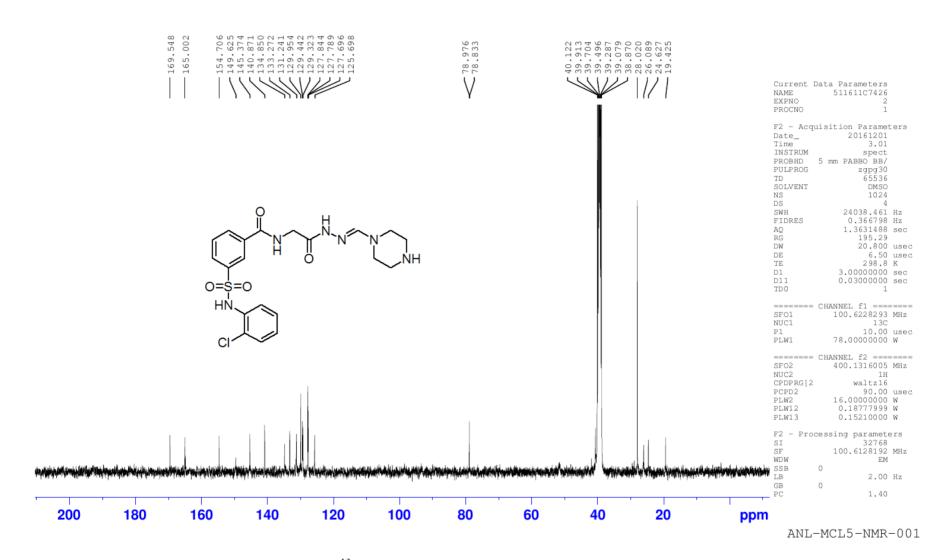
511611B4317 25 (0.362) AM (Top,4, Ar,5000.0,195.15,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00) 1: TOF MS ES+ 1.29e3 100-479.0207 479.1582 351.0823 364.0349 498.9683 481.1180 423.1699 0-հուրադանադադադադադադադադադադանընդուրաըստրաբարարար 400 410 420 450 470 490 460 Minimum: -1.5Maximum: 5.0 10000.0 50.0 Mass Calc. Mass mDa PPM DBE i-FIT Formula

78957.8

C20 H18 N4 O4 S2 C1

HRMS of Compound-1u

13.5



¹³CNMR of **Compound-1v** in DMSO- d_6

Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

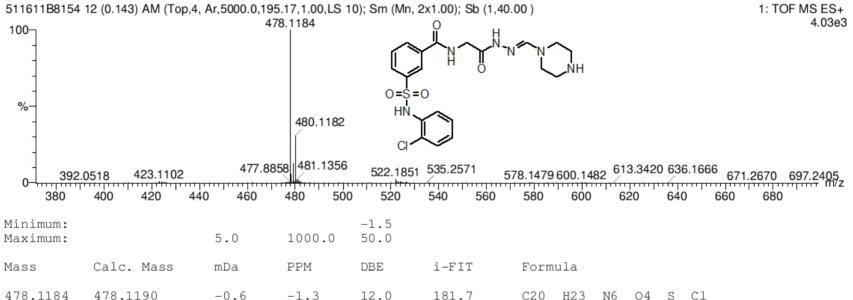
Monoisotopic Mass, Odd and Even Electron lons

133 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass)

Elements Used:

C: 0-20 H: 0-23 N: 0-6 O: 0-4 S: 0-1 CI: 0-1

GVB-SA10



HRMS of Compound-1v

GVB-SA14

Reference Code: 511611B2466

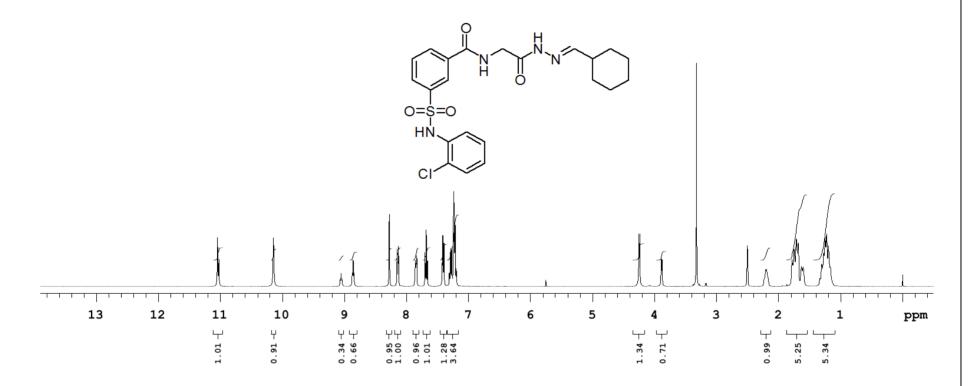
Solvent: dmso
Archive directory:

/home/gvkbio/data/2016/Nov

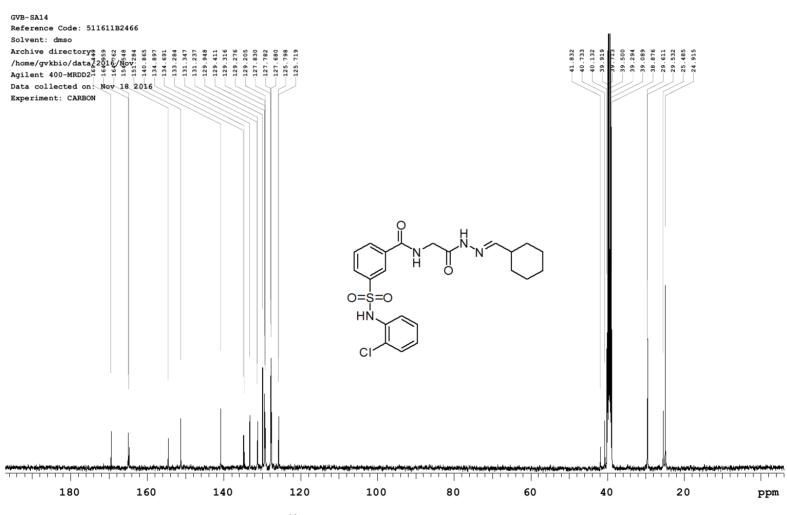
Agilent 400-MRDD2

Data collected on: Nov 18 2016

Experiment: PROTON



 $^{1}\text{H NMR of }\text{Compound-1}\text{w}\text{ in DMSO-d}_{6}$



 13 CNMR of **Compound-1w** in DMSO- d_6

Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

Calc. Mass

477.1363

94 formula(e) evaluated with 1 results within limits (up to 1 best isotopic matches for each mass)

PPM

2.3

mDa

1.1

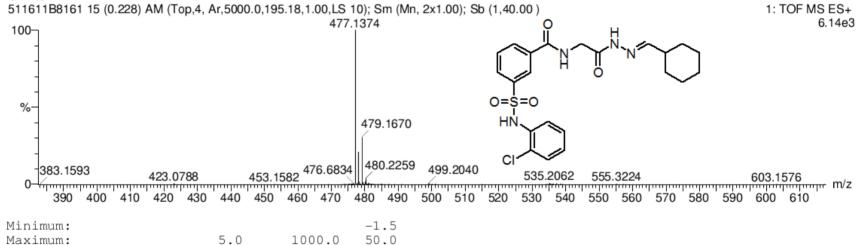
Elements Used:

C: 0-22 H: 0-26 N: 0-4 O: 0-4 S: 0-1 CI: 0-1

GVB-SA14

Mass

477.1374



i-FIT

132.3

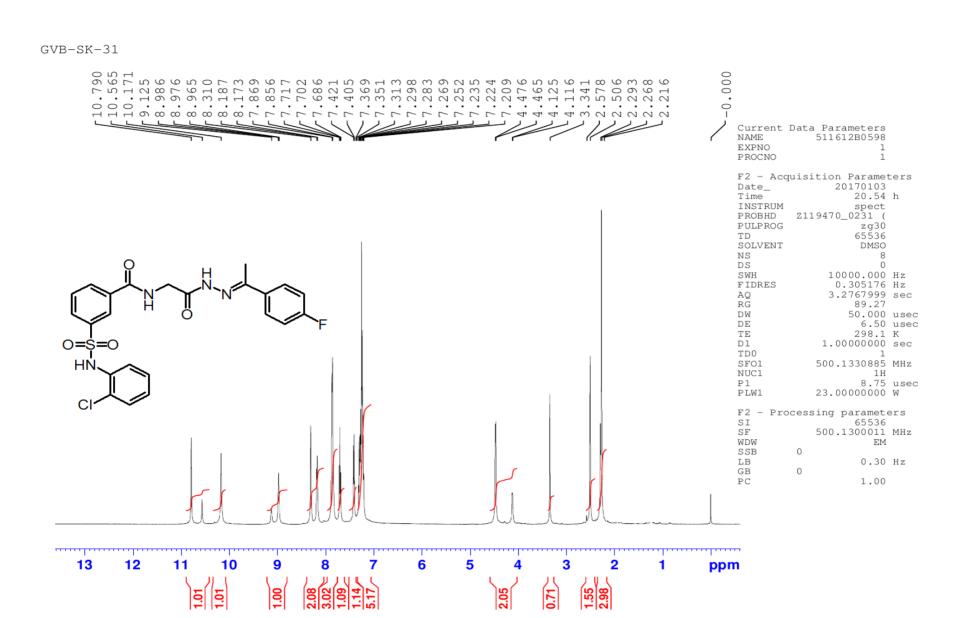
Formula

C22 H26 N4 O4 S C1

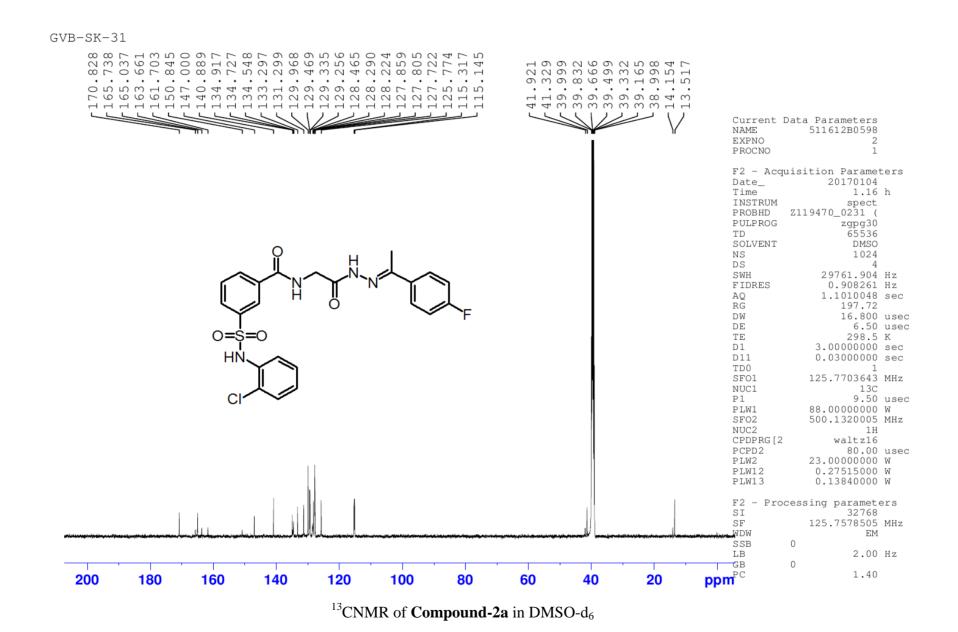
HRMS of Compound-1w

DBE

11.5



¹H NMR of **Compound-2a** in DMSO-d₆



Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

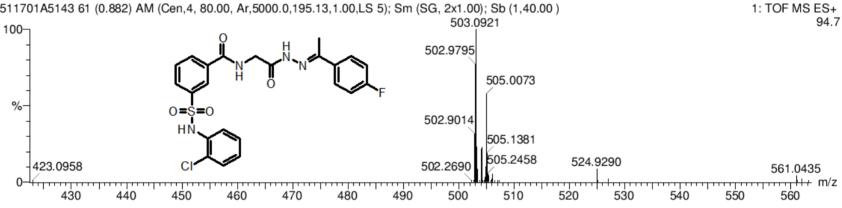
190 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

C: 0-23 H: 0-21 N: 0-4 O: 0-4 F: 0-1 S: 0-1 CI: 0-1

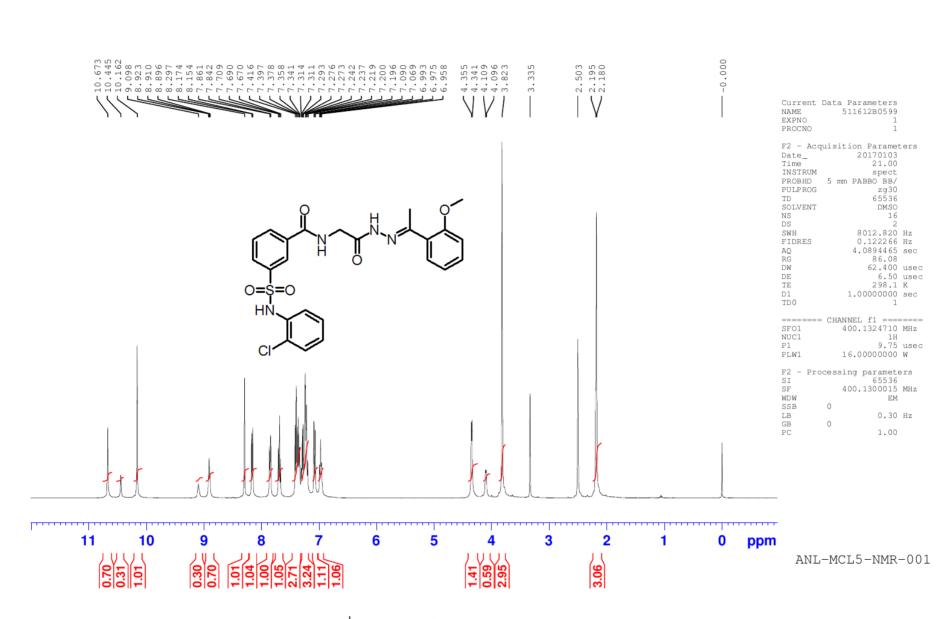
SAMPLE CODE:COMPOUND-31

511701A5143 61 (0.882) AM (Cen,4, 80.00, Ar,5000.0,195.13,1.00,LS 5); Sm (SG, 2x1.00); Sb (1,40.00)

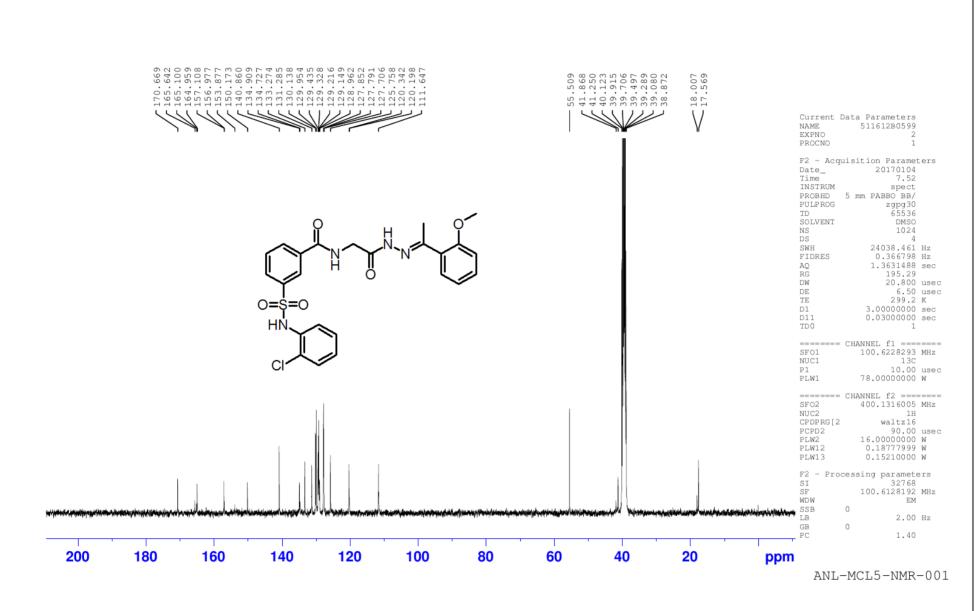


Minimum: Maximum:		5.0	1000.0	-1.5 50.0								
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Formula						
503.0921	503.0956	-3.5	-7.0	14.5	7.9	C23	H21	N4	04	F	S	Cl

HRMS of Compound-2a



¹H NMR of **Compound-2b** in DMSO-d₆



¹³CNMR of **Compound-2b** in DMSO-d₆

Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

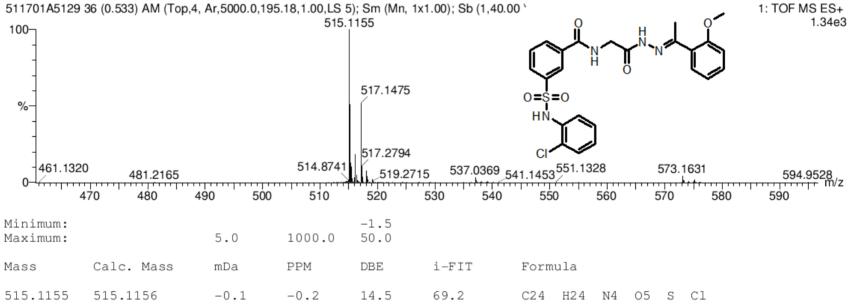
Monoisotopic Mass, Even Electron Ions

199 formula(e) evaluated with 69 results within limits (up to 1 closest results for each mass)

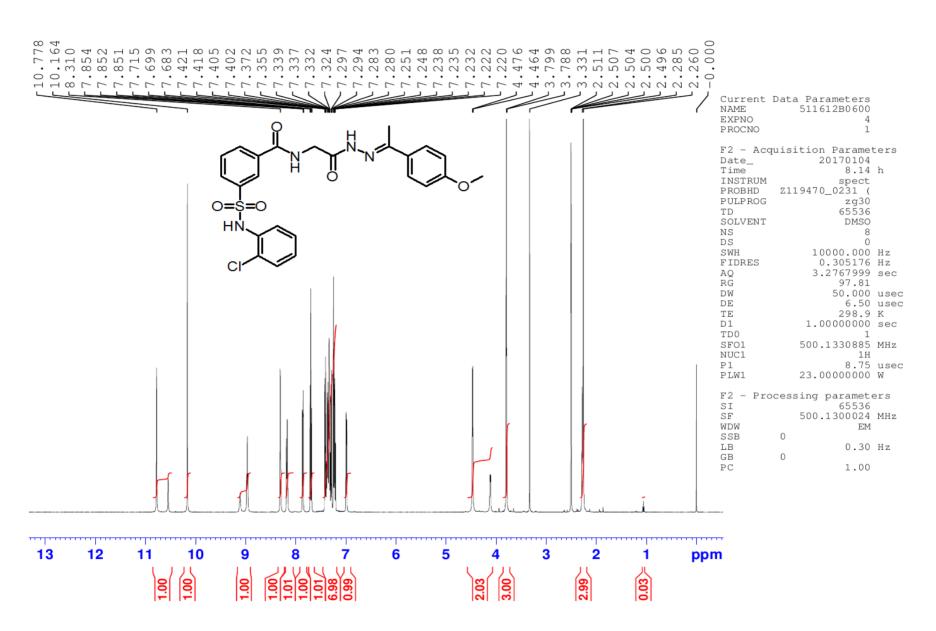
Elements Used:

C: 0-29 H: 0-25 N: 0-4 O: 0-5 F: 0-1 S: 0-1 CI: 0-1

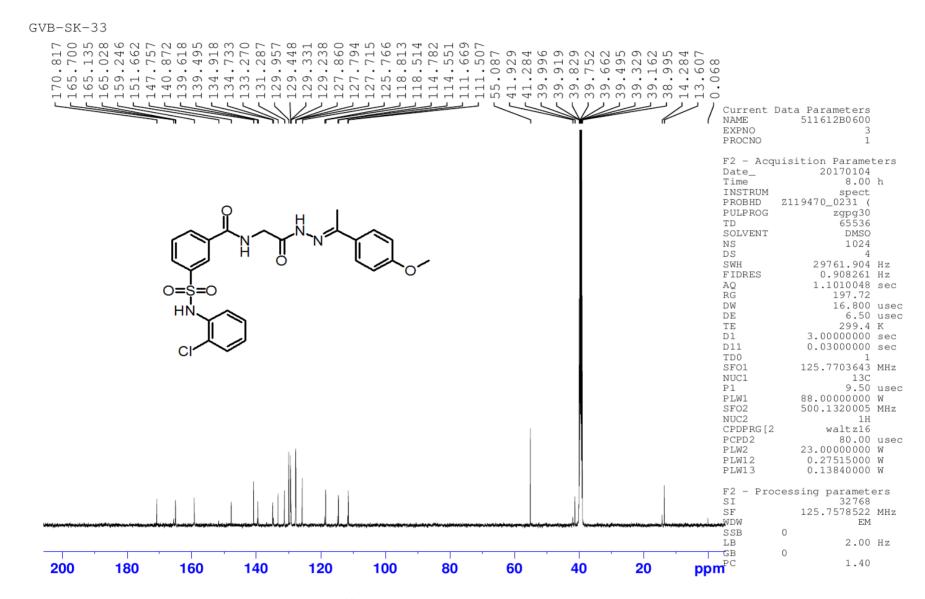
SAMPLE CODE:COMPOUND-32



HRMS of Compound-2b



¹H NMR of **Compound-2c** in DMSO-d₆



¹³CNMR of **Compound-2c** in DMSO-d₆

Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

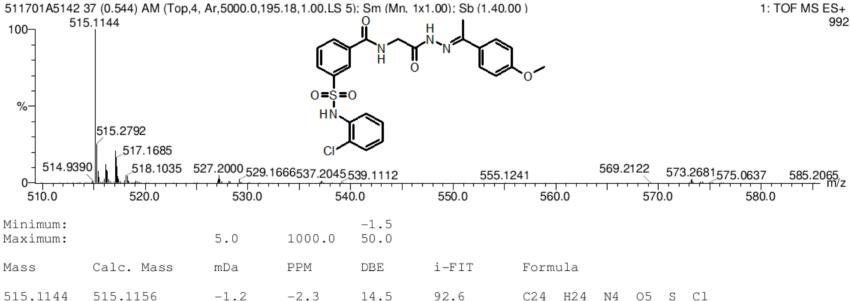
Monoisotopic Mass, Even Electron Ions

113 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

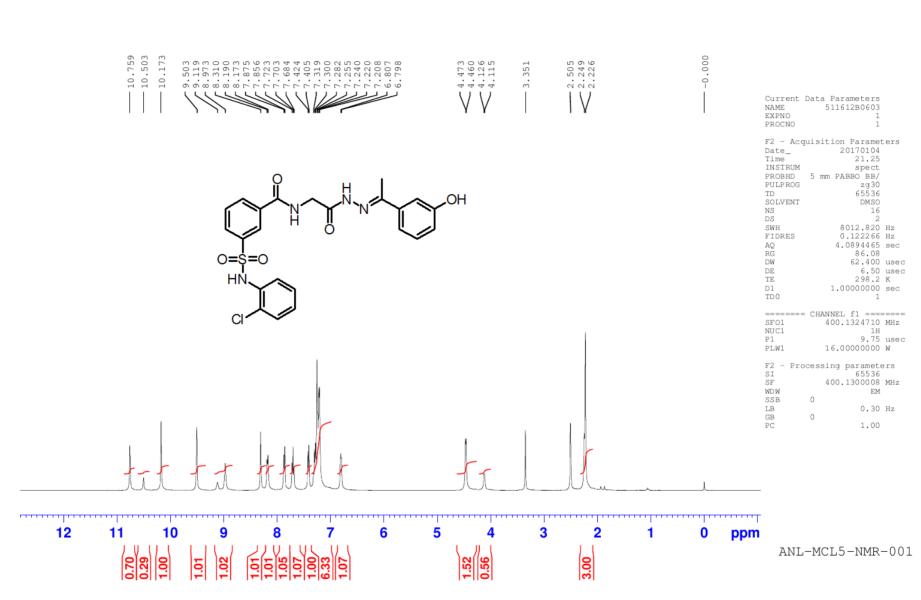
Elements Used:

C: 0-24 H: 0-24 N: 0-4 O: 0-5 S: 0-1 CI: 0-1

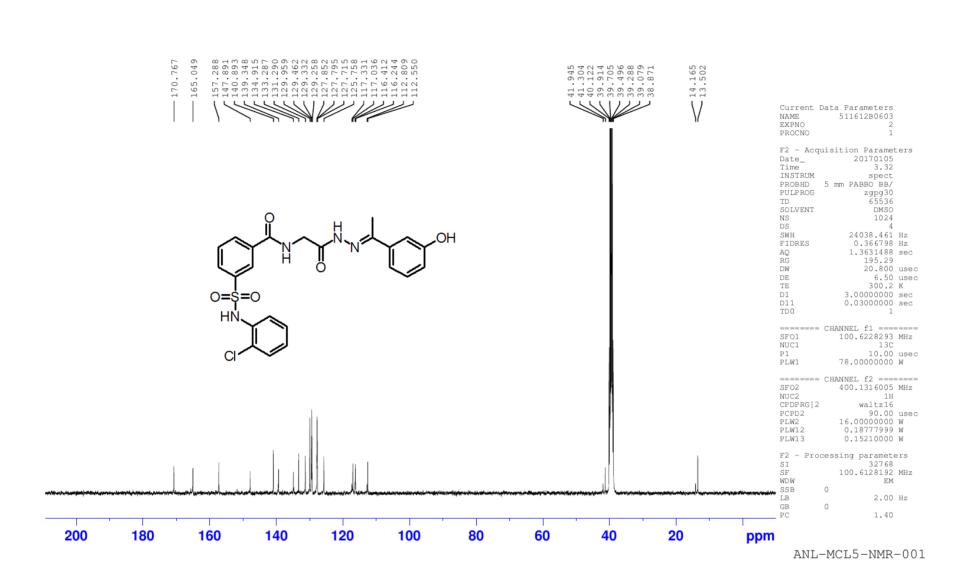
SAMPLE CODE:COMPOUND-33



HRMS of Compound-2c



¹H NMR of **Compound-2d** in DMSO-d₆



¹³CNMR of **Compound-2d** in DMSO-d₆

Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

221 formula(e) evaluated with 4 results within limits (up to 1 closest results for each mass)

Elements Used:

C: 0-23 H: 0-22 N: 0-4 O: 0-5 F: 0-1 S: 0-1 CI: 0-1

SAMPLE CODE: COMPOUND-36

511701A5132 18 (0.259) AM (Top,4, Ar,5000.0,195.16,1.00,LS 5); Sm (Mn, 1x1.00); Sb (1,40.00) 1: TOF MS ES+ 6.42e3 501.1035 100-0=S=0 % 503.1077 504.1275 500.8979 294.0909 350.9822 423.1324 447.1730 523.0488 559.1100 619.1860 340 360 520 300 320 380 400 420 440 460 480 500 540 580 600 Minimum: -1.55.0 Maximum: 1000.0 50.0 PPM DBE i-FIT Mass Calc. Mass mDa Formula 501.1035 501.0999 3.6 7.2 14.5 91.8 C23 H22 N4 O5 S C1

HRMS of Compound-2d

Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

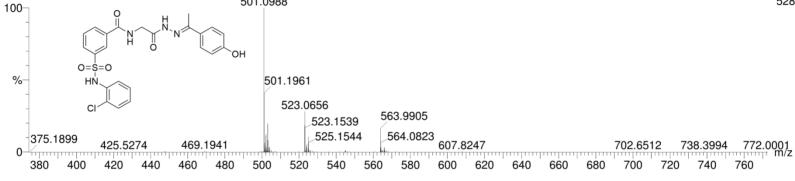
113 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

C: 0-23 H: 0-22 N: 0-4 O: 0-5 S: 0-1 CI: 0-1

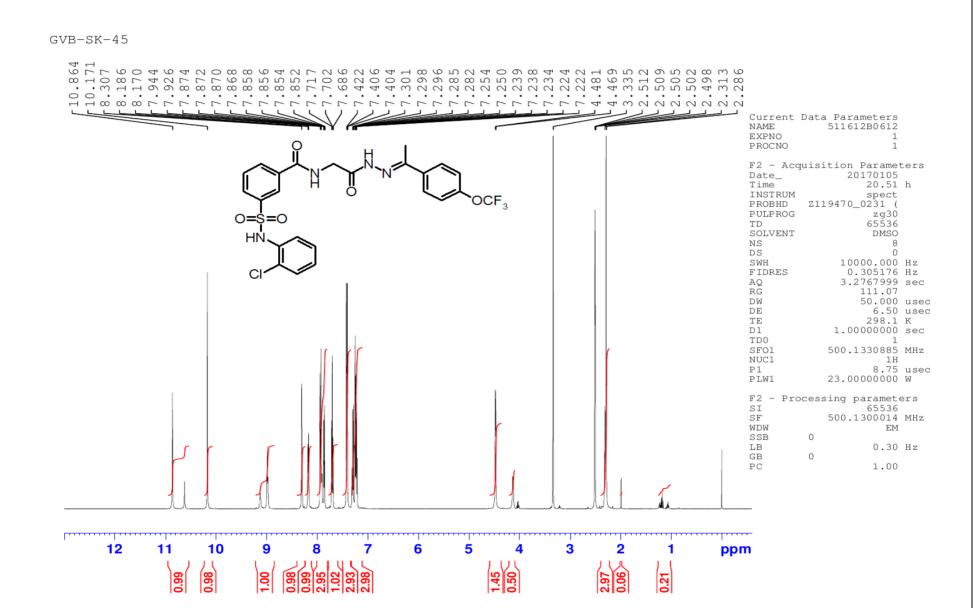
GVB-SK44-01

511702A3696 11 (0.157) AM (Top,4, Ar,5000.0,195.49,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00) 1: TOF MS ES+ 528

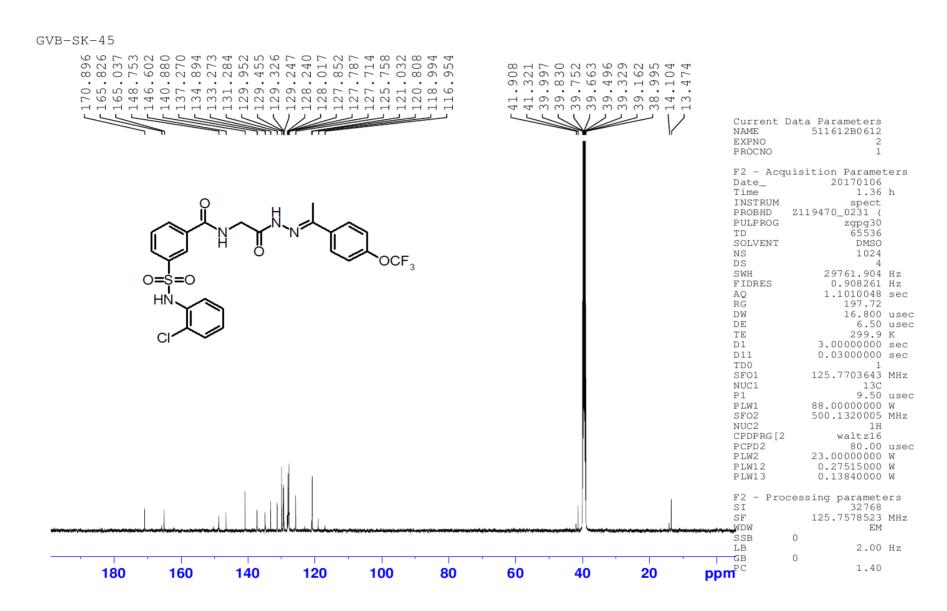


Minimum: -1.55.0 1000.0 Maximum: 50.0 Calc. Mass mDa PPM DBE i-FIT Formula Mass -1.1-2.250.9 501.0988 501.0999 14.5 C23 H22 N4 O5 S C1

HRMS of Compound-2e



¹H NMR of **Compound-2f** in DMSO-d₆



¹³CNMR of **Compound-2f** in DMSO-d₆

Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

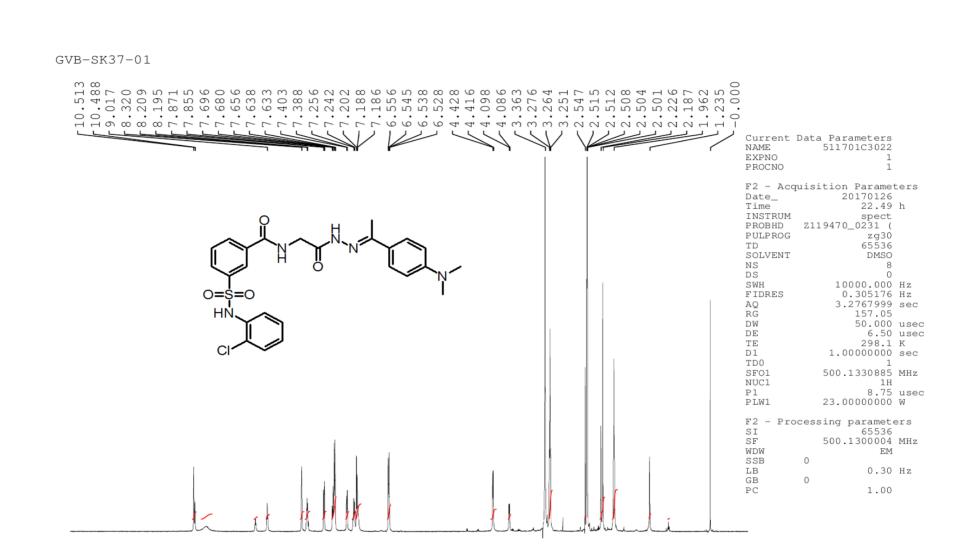
469 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

C: 0-24 H: 0-30 N: 0-4 O: 0-5 S: 0-1 CI: 0-1 F: 0-3

SAMPLE CODE:COMPOUND-45 511701A5130 56 (0.797) AM (Top,4, Ar,5000.0,195.15,1.00,LS 5); Sm (Mn, 1x1.00); Sb (1,40.00) 1: TOF MS ES+ 893 569.0854 100-571.0362 515.0394 591.0215 568.6010 353.1430 503.1387 392.0139 627.0258 440 360 380 400 420 460 480 500 520 540 560 580 600 620 Minimum: -1.55.0 1000.0 50.0 Maximum: Calc. Mass mDa PPM DBE i-FIT Formula Mass 569.0854 569.0873 -1.9-3.3 14.5 38.7 C24 H21 N4 O5 S C1 F3

HRMS of Compound-2f

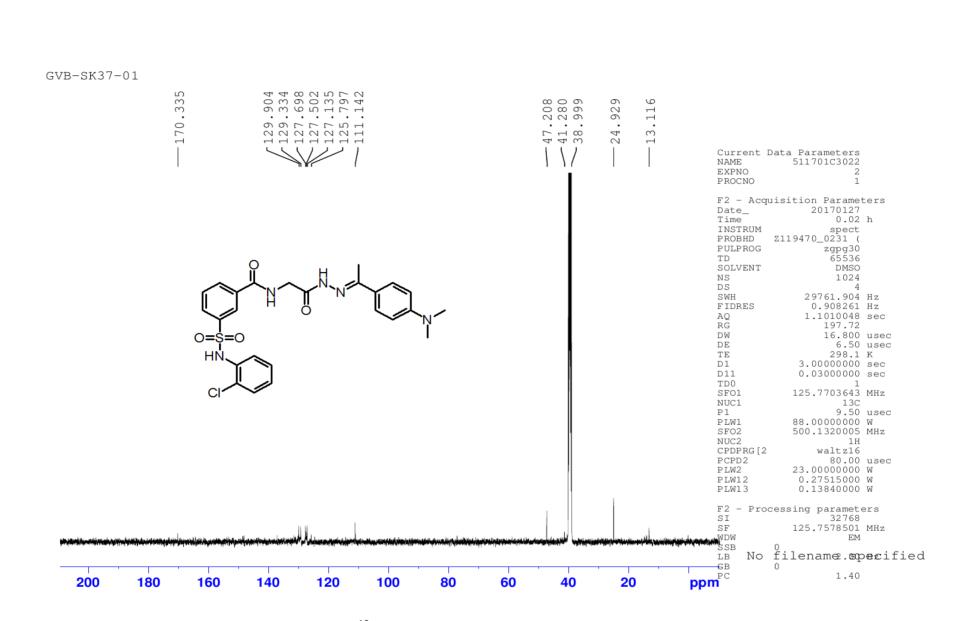


¹H NMR of **Compound-2g** in DMSO-d₆

ppm

12

11



¹³CNMR of **Compound-2g** in DMSO-d₆

Page 1

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

113 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

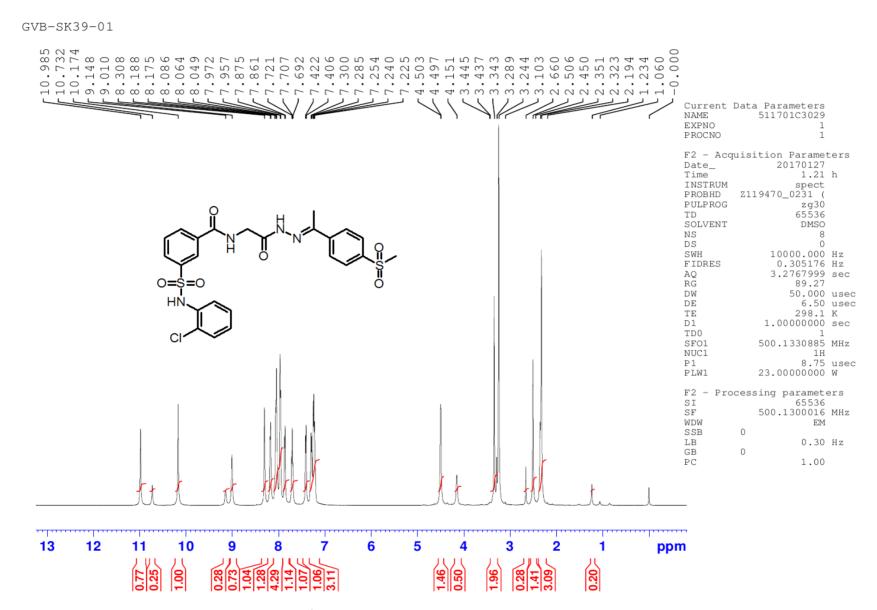
C: 0-25 H: 0-27 N: 0-5 O: 0-4 S: 0-1 CI: 0-1

GVB-SK37

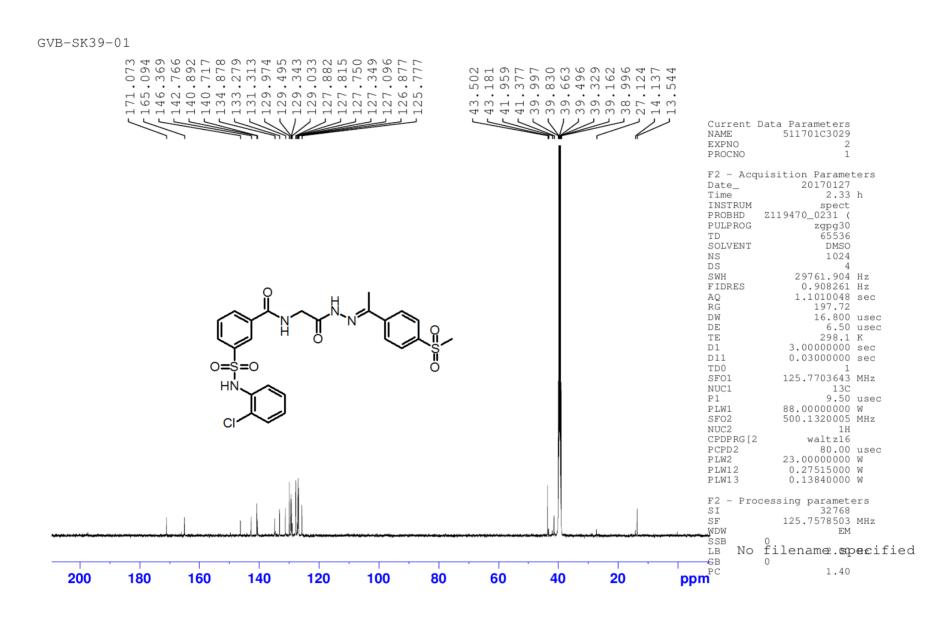
511702A2311 12 (0.165) AM (Top,4, Ar,5000.0,195.50,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00) 1: TOF MS ES+ 528.1440 3.42e3 100-% 530.1541 195.5202 527.7668 531.1215 568.1376 178.5628 236.5289 712.8326 m/z 294.3227 351.3608 418.5677 325 350 375 400 425 450 475 500 525 550 575 600 625 650 675 700 225 250 275 300 Minimum: -1.5

Maximum: 5.0 1000.0 50.0 i-FIT Mass Calc. Mass mDa PPM DBE Formula -3.2-6.1528.1440 528.1472 14.5 196.0 C25 H27 N5 O4 S C1

HRMS of Compound-2g



¹H NMR of **Compound-2h** in DMSO-d₆



¹³CNMR of **Compound-2h** in DMSO-d₆

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Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

563.0826

203 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

0.2

0.1

Elements Used:

C: 0-24 H: 0-24 N: 0-4 O: 0-6 S: 0-2 CI: 0-1

GVB-SK39-01

563.0827

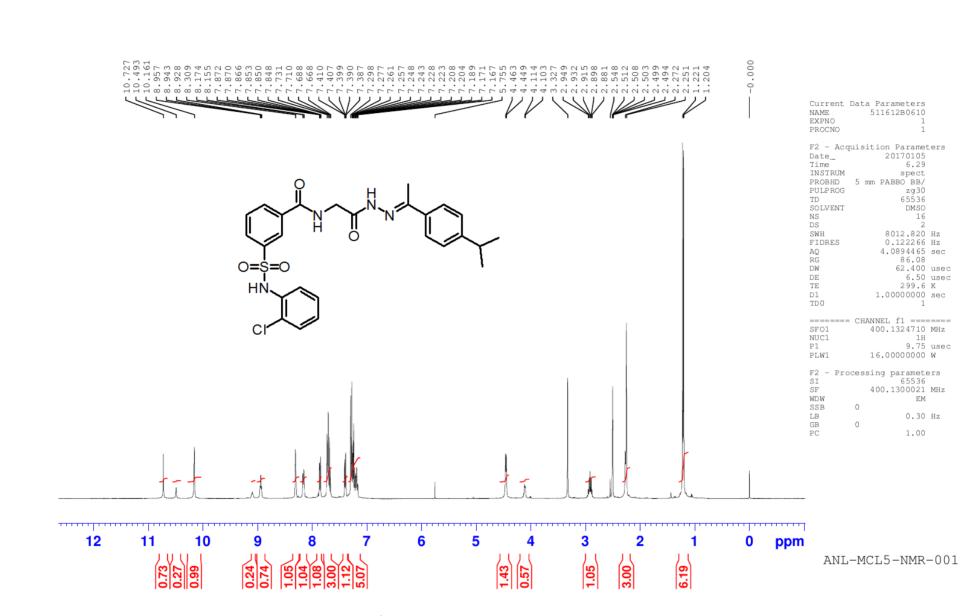
511702A2304 9 (0.137) AM (Top,4, Ar,5000.0,195.52,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00) 1: TOF MS ES+ 563.0827 315 100-563.1514 % 563.2201 565.1122 565.3533 562,7390 392.2685 528.2061 759.0891 587.0115 436.2755 400 500 520 540 580 460 480 560 620 640 680 700 720 440 600 660 740 -1.5Minimum: 5.0 1000.0 50.0 Maximum: Mass Calc. Mass mDa PPM DBE i-FIT Formula

HRMS of Compound-2h

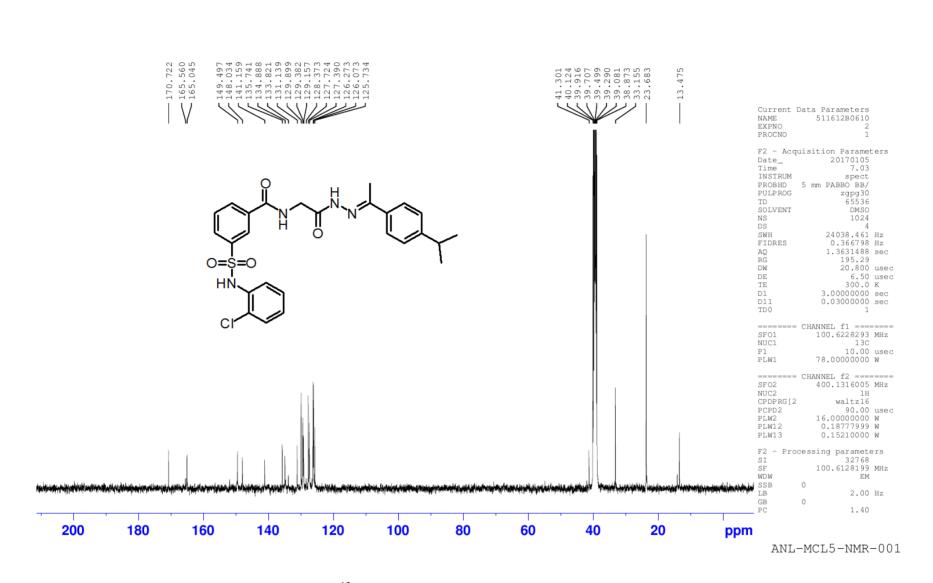
26.5

C24 H24 N4 O6 S2 Cl

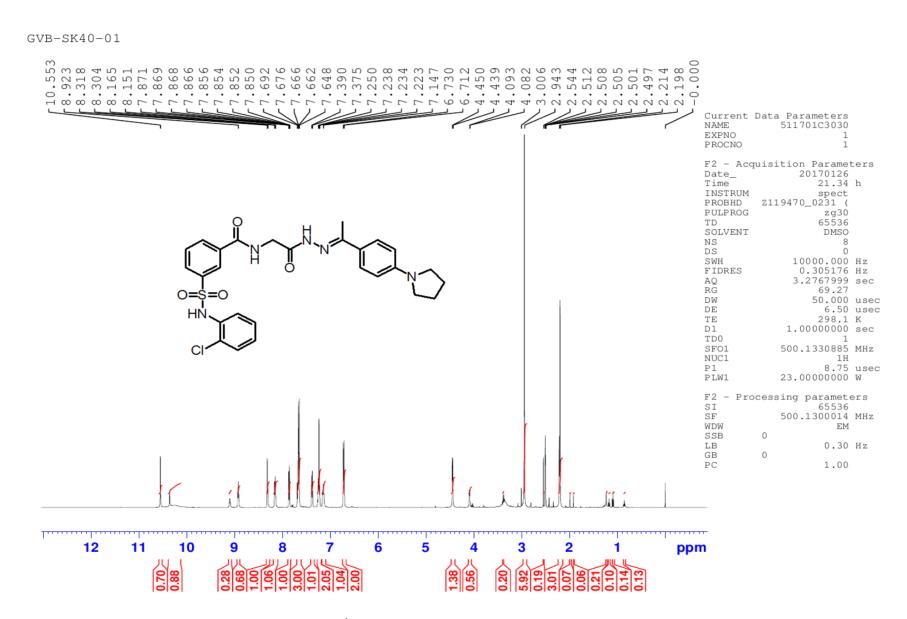
14.5



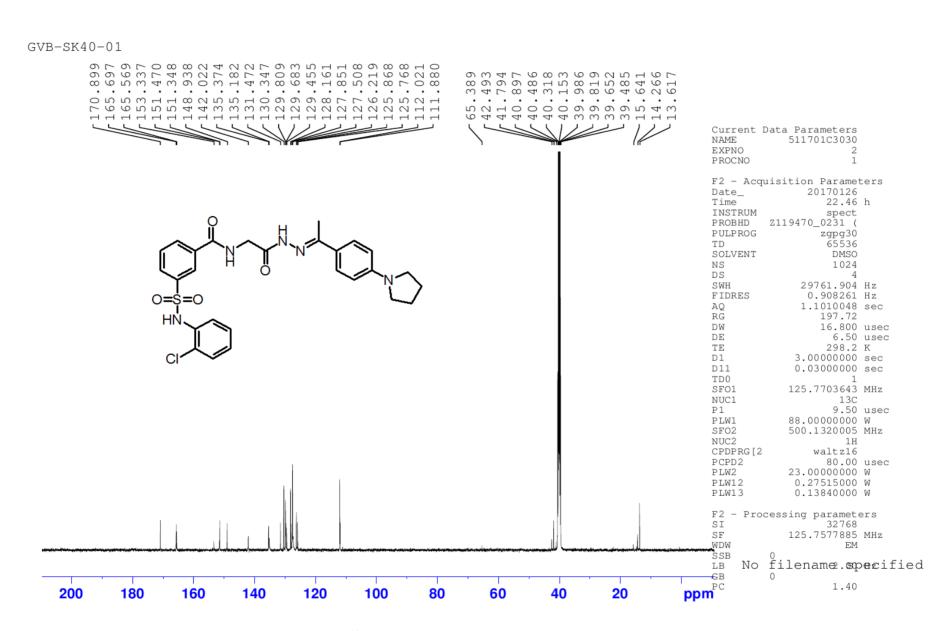
¹H NMR of **Compound-2i** in DMSO-d₆



¹³CNMR of **Compound-2i** in DMSO-d₆



¹H NMR of **Compound-2j** in DMSO-d₆



¹³CNMR of **Compound-2j** in DMSO-d₆

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Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

113 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

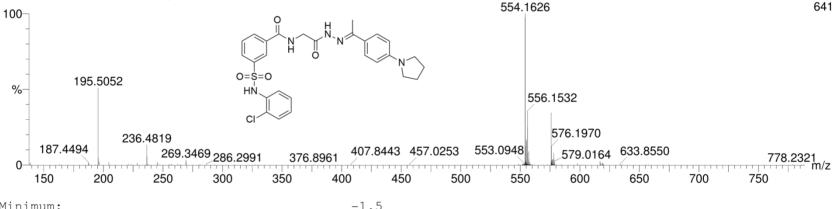
Elements Used:

C: 0-27 H: 0-29 N: 0-5 O: 0-4 S: 0-1 CI: 0-1

GVB-SK40

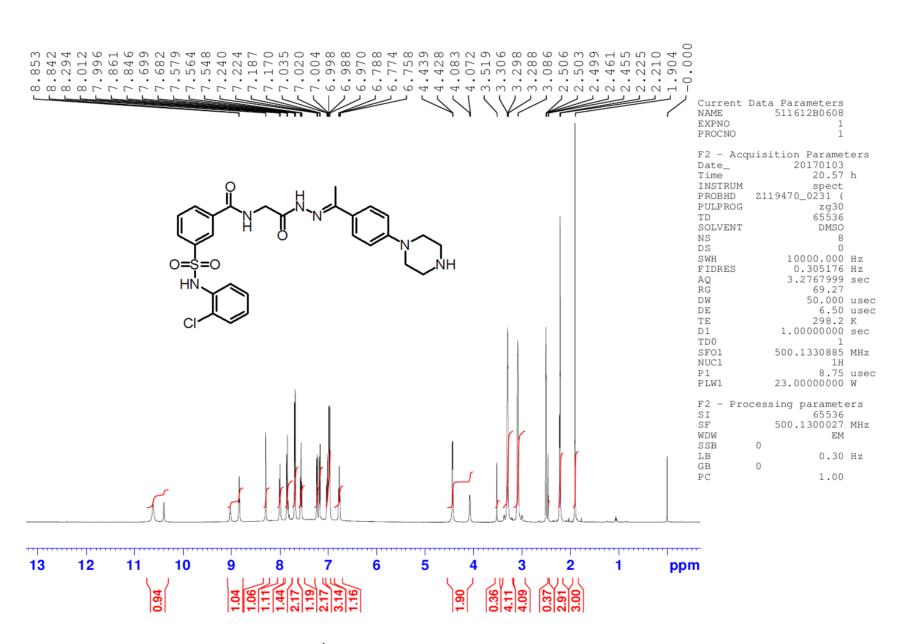
511702A2314 9 (0.136) AM (Top,4, Ar,5000.0,195.51,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00)

1: TOF MS ES+

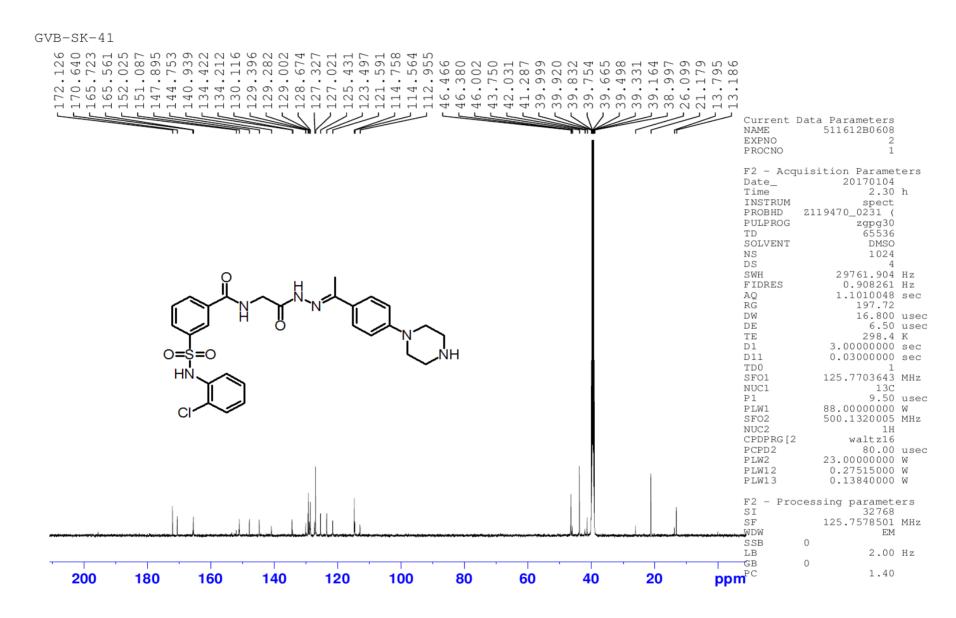


Minimum: -1.5Maximum: 5.0 1000.0 50.0 Calc. Mass Mass mDa PPM DBE i-FIT Formula 554.1626 554.1629 -0.3-0.515.5 31.4 C27 H29 N5 O4 S C1

HRMS of Compound-2J



¹H NMR of **Compound-2k** in DMSO-d₆



¹³CNMR of **Compound-2k** in DMSO-d₆

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Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

133 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

539.1884

540

Elements Used:

487.2667

500

490

100-

C: 0-27 H: 0-30 N: 0-6 O: 0-4 S: 0-1 CI: 0-1

515.2338

520

510

SAMPLE CODE:COMPOUND-41

511701A5134 32 (0.462) AM (Top,4, Ar,5000.0,195.17,1.00,LS 5); Sm (Mn, 2x1.00); Sb (1,40.00)

530

1: TOF MS ES+
4.19e3

630

640

Minimum: Maximum:		5.0	1000.0	-1.5 50.0		
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Formula
569.1760	569.1738	2.2	3.9	15.5	30.6	C27 H30 N6 O4

550

HRMS of Compound-2K

568.8530

560

571.1616

572.2020

580

591.2084

590

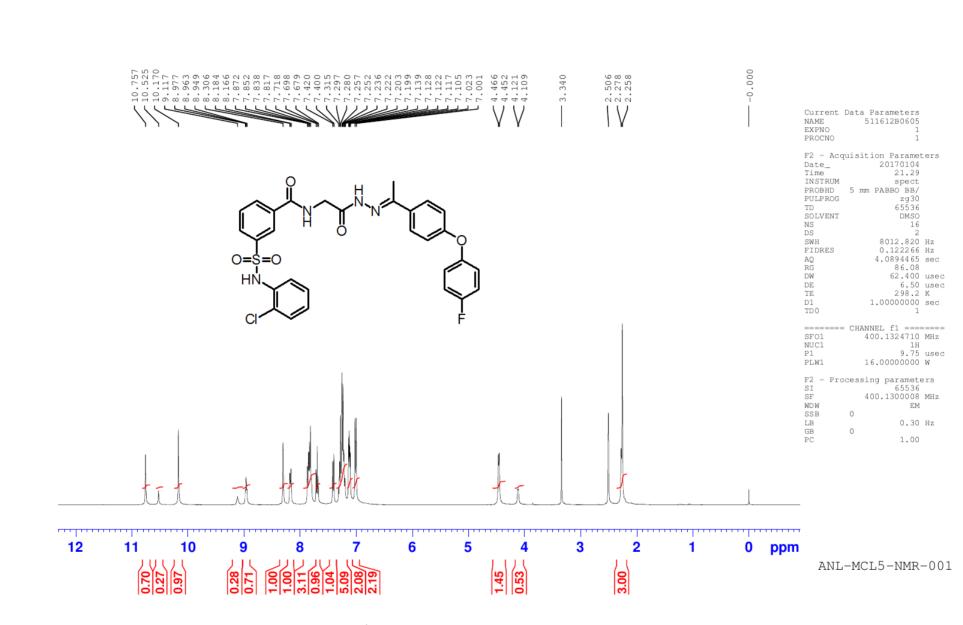
609.0690

610

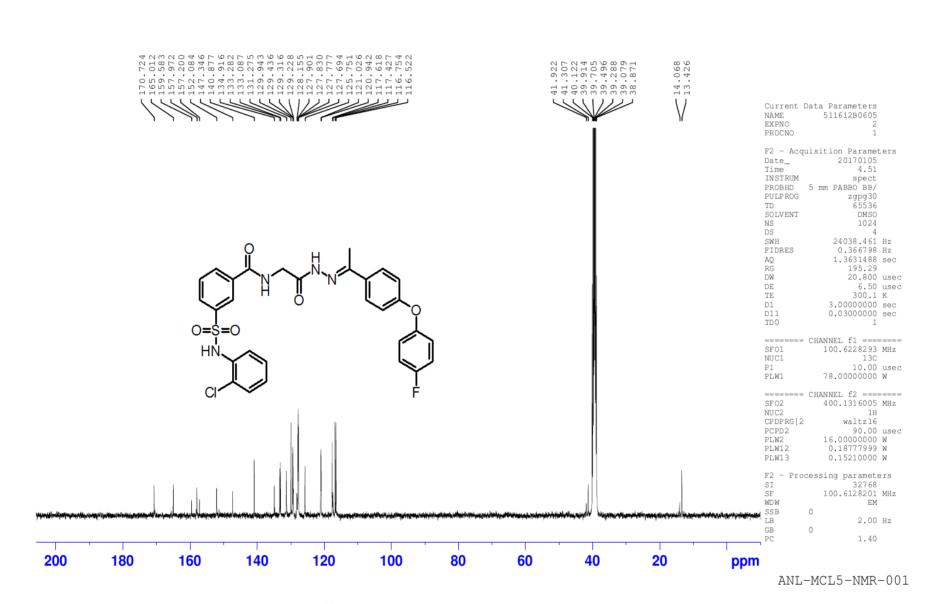
620

S Cl

600



¹H NMR of **Compound-2m** in DMSO-d₆



¹³CNMR of **Compound-2m** in DMSO-d₆

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Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

226 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

C: 0-29 H: 0-25 N: 0-4 O: 0-5 F: 0-1 S: 0-1 CI: 0-1

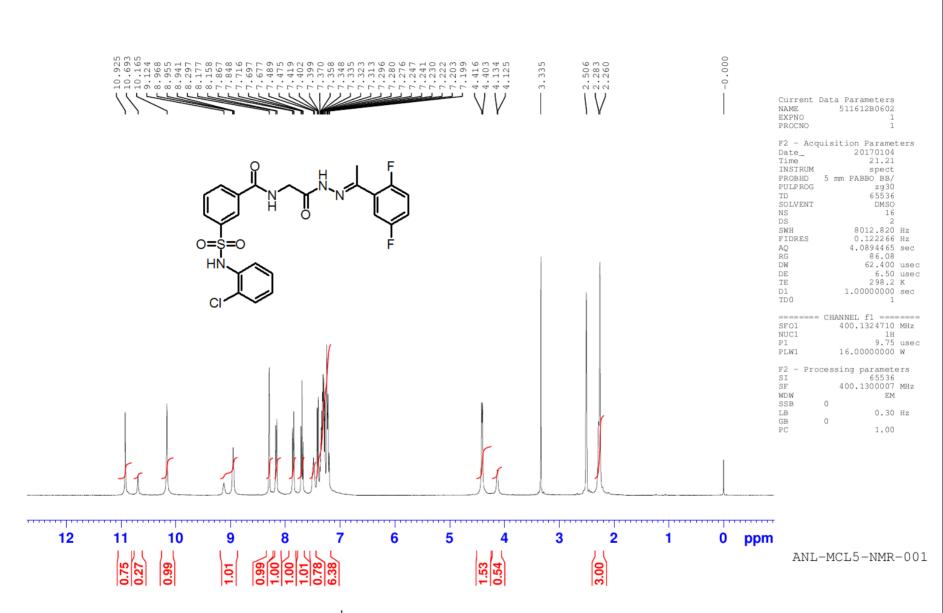
SAMPLE CODE:COMPOUND-38

511701A5149 28 (0.392) AM (Top,4, Ar,5000.0,195.13,1.00,LS 5); Sm (Mn, 1x1.00); Sb (1,40.00)

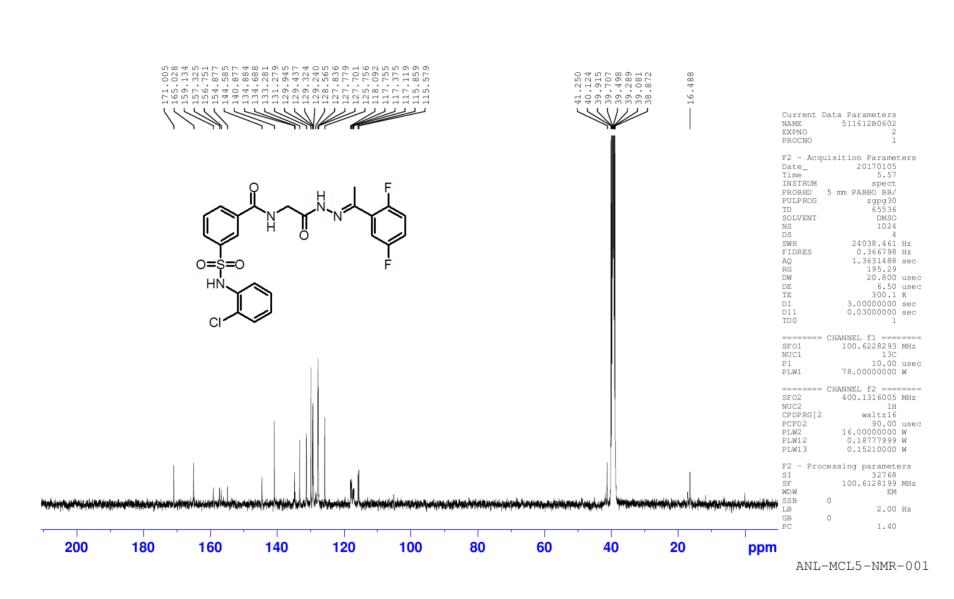
1: TOF MS ES+ 595.1225 1.39e3 100-597.0584 597.2001 594.8160 501.1286 .598.1930 527.0728 541.2516 569.0087 618.2709 653.1298 m/z 510 530 540 550 560 570 580 590 600 610 620 630 640 660 -1.55.0 1000.0 50.0

Minimum: Maximum: Calc. Mass PPM DBE i-FIT Mass mDa Formula 595.1225 595.1218 0.7 1.2 18.5 78.0 C29 H25 N4 O5 F S C1

HRMS of Compound-2m



¹H NMR of **Compound-2n** in DMSO-d₆



¹³CNMR of **Compound-2n** in DMSO-d₆

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Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

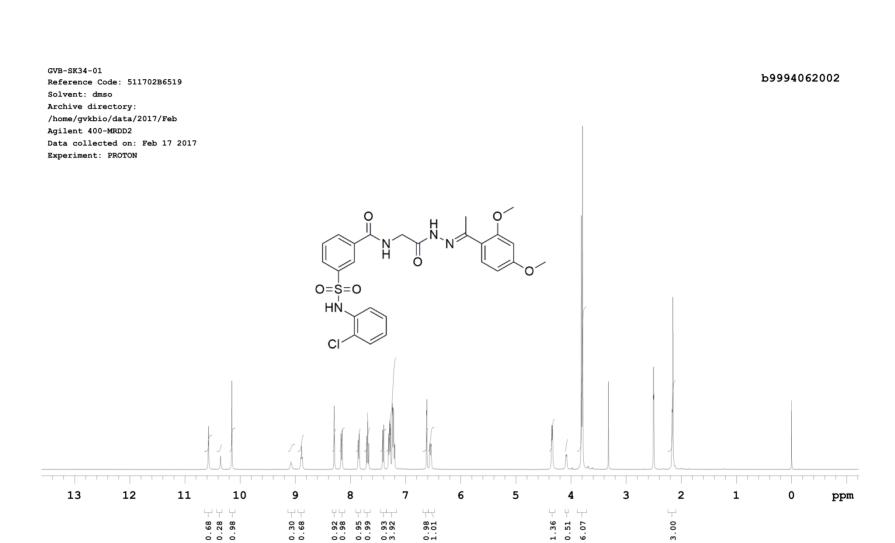
337 formula(e) evaluated with 3 results within limits (up to 1 closest results for each mass)

Elements Used:

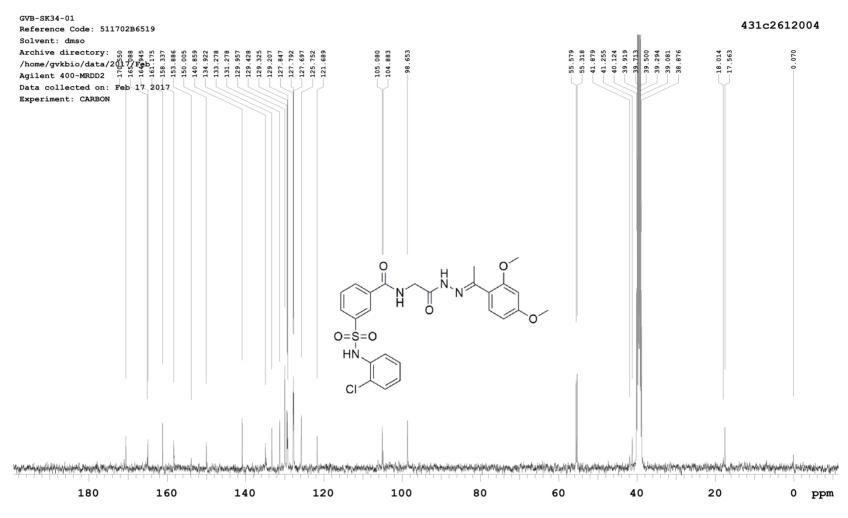
C: 0-23 H: 0-20 N: 0-4 O: 0-5 S: 0-1 CI: 0-1 F: 0-2

SAMPLE CODE:COMPOUND-35 511701A5137 15 (0.229) AM (Top,4, Ar,5000.0,195.16,1.00,LS 5); Sm (Mn, 1x1.00); Sb (1,40.00) 1: TOF MS ES+ 521.0829 6.02e3 100-523.1265 523.2260 543.1298 323.2168 351.0203 503.1191 520.6967 57<u>9.1435</u> m/z 396.1026 425.0851 464.1913 580 360 380 400 420 440 460 480 500 540 560 Minimum: -1.55.0 Maximum: 1000.0 50.0 Calc. Mass mDa PPM DBE i-FIT Formula Mass 521.0829 -3.3 -6.3 14.5 579.7 521.0862 C23 H20 N4 O4 S C1 F2

HRMS of Compound-2n



 $^{1}\text{H NMR of }$ Compound-20 in DMSO-d₆



Plotname: 511702B6519_CARBON_01.REC_plot01

 $^{13}\mbox{CNMR}$ of $\mbox{Compound-2o}$ in DMSO-d $_{6}$

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1: TOF MS ES+

Single Mass Analysis

Tolerance = 1000.0 PPM / DBE: min = -1.5, max = 50.0

Selected filters: None

Monoisotopic Mass, Even Electron Ions

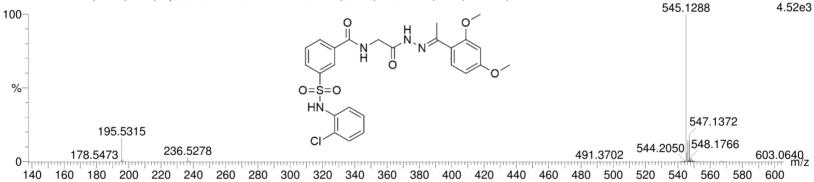
133 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

C: 0-25 H: 0-26 N: 0-4 O: 0-6 S: 0-1 CI: 0-1

GVB-SK34-01

511702A3706 13 (0.175) AM (Top,4, Ar,5000.0,195.54,1.00,LS 10); Sm (Mn, 2x1.00); Sb (1,40.00)



Minimum: Maximum:		5.0	1000.0	-1.5 50.0							
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Formula					
545.1288	545.1262	2.6	4.8	14.5	427.1	C25	H26	N4	06	S	Cl

HRMS of Compound-20