

Synthesis and Antibacterial Evaluation of Quinoline-Sulfonamide Hybrid Compounds: A Promising Strategy Against Bacterial Resistance

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Table S1: Minimum inhibitory concentration (MIC) of compound QS-3 and standard drug CIP in µg/mL.

Compound	<i>E. coli</i>	<i>E. faecalis</i>	<i>P. aeruginosa</i>	<i>K. pneumoniae</i>	<i>Salmonella typhi</i>
QS-3	128	128	64	256	512
CIP	0.5	0.5	0.5	0.5	0.5

Table S2: Zone of inhibition (mm) against environmental Multidrug-resistant bacterial isolates.

S. No.	Isolate Name	QS-3	AMP	CIP
1.	EJH01	-	-	15
2.	EJH02	-	30	26
3.	EJH04	-	-	-
4.	EJH05	-	20	25
5.	EJH07	-	25	22
6.	EJH11	-	21	26
7.	EJH12	-	-	25
8.	EJH13	-	-	20
9.	EJH14	-	-	22
10.	EJH15	-	20	17
11.	EJH18	-	17	25
12.	EJH19	-	17	21
13.	EJH20	-	20	19
14.	EJH21	-	18	26
15.	EJH22	-	32	14
16.	EJH23	-	20	24
17.	EJH24	-	23	28
18.	EJH25	--	22	21
19.	A200	-	-	18
20.	AA 201	-	-	16

21.	AA 202	8	-	-
22.	AA 209	-	30	21
23.	AA 216	-	-	24
24.	AA 221	-	-	21
25.	AA 224	-	-	11
26.	AA 237	-	26	25
27.	AA 240	-	23	25
28.	AA 243	-	20	19
29.	AA 245	-	18	22
30.	AA 248	-	25	27
31.	AA 261	-	25	29
32.	AA 269	-	-	-
33.	AA 273	-	15	14
34.	AA 276	-	22	23
35.	AA 290	8	-	-

Table S3. Drug ability or Lipinski's 'rule-of-five' data of compounds QS1-QS12.

Compounds	Molecular Wt.	No. of HBA	No. of HBD	Mol log P	Molar Refractivity (MR)	Lipinski violation
QS-1	460.94	4	2	3.13	128.71	0
QS-2	427.88	4	2	3.59	113.37	0
QS-3	439.91	4	2	2.88	119.9	0
QS-4	459.95	3	2	3.88	130.92	0
QS-5	451.97	3	2	3.85	127.99	0

QS-6	485.98	3	2	4.25	138.85	1
QS-7	460.94	4	2	3.13	128.71	0
QS-8	427.88	4	2	3.59	113.37	0
QS-9	439.91	4	2	2.88	119.9	0
QS-10	459.95	3	2	3.88	130.92	0
QS-11	451.97	3	2	3.85	127.99	0
QS-12	485.98	3	2	4.25	138.85	1

NMR spectrum of compound QS1-QS12

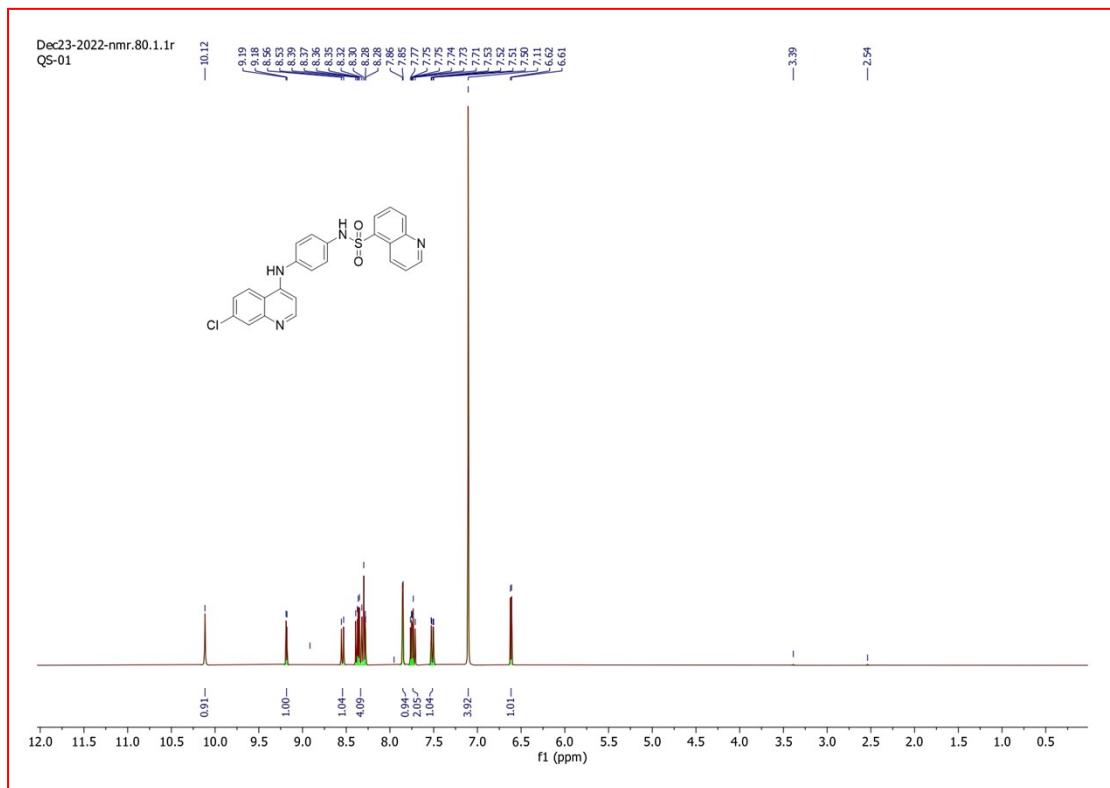


Figure.- S1. ^1H NMR Spectrum of compound (QS-1)

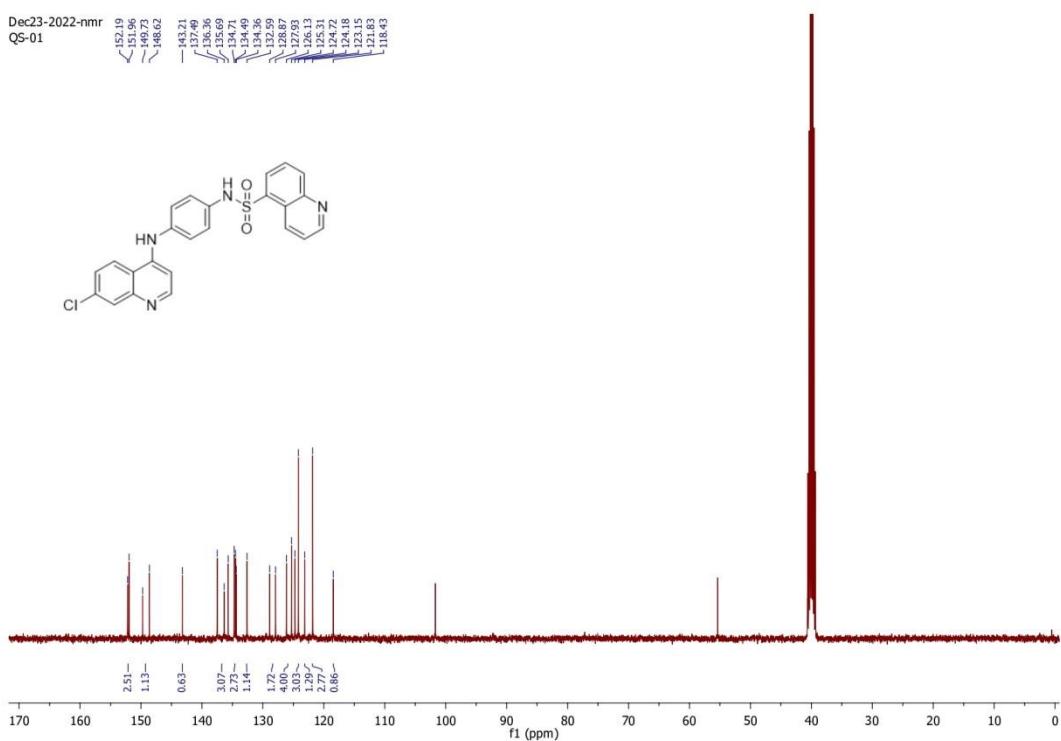


Figure.- S2. ^{13}C NMR Spectrum of compound (QS-1)

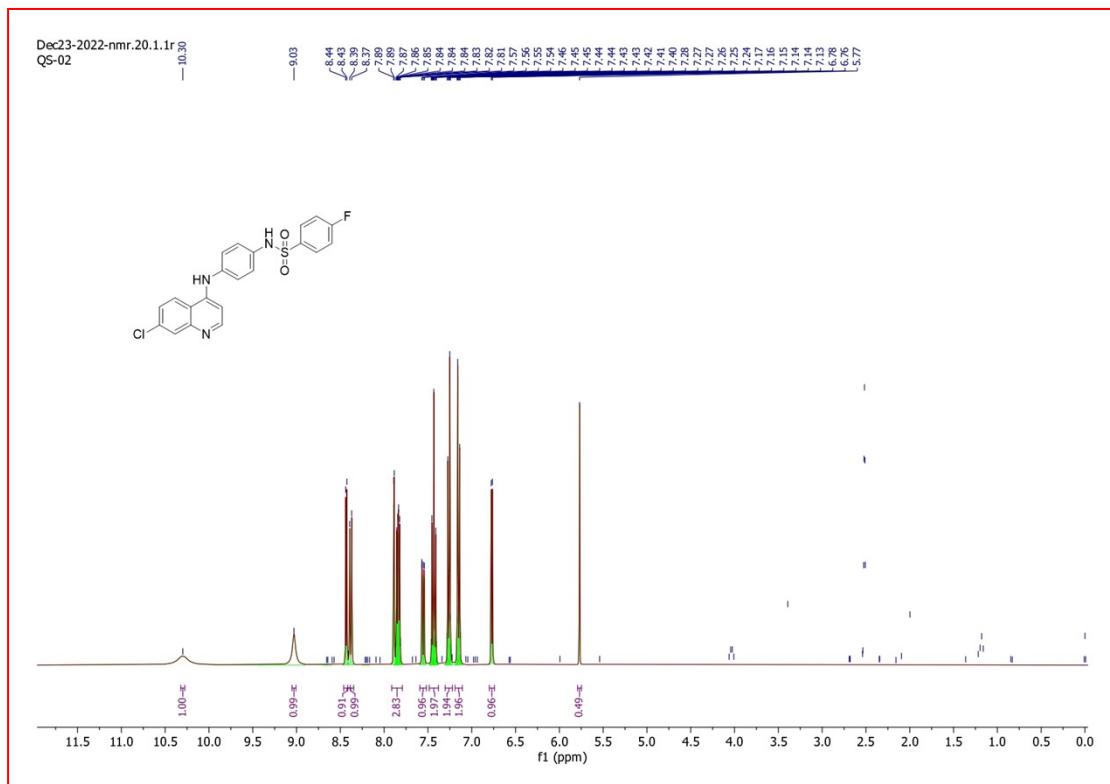


Figure-S3. ^1H NMR Spectrum of compound (QS-2)

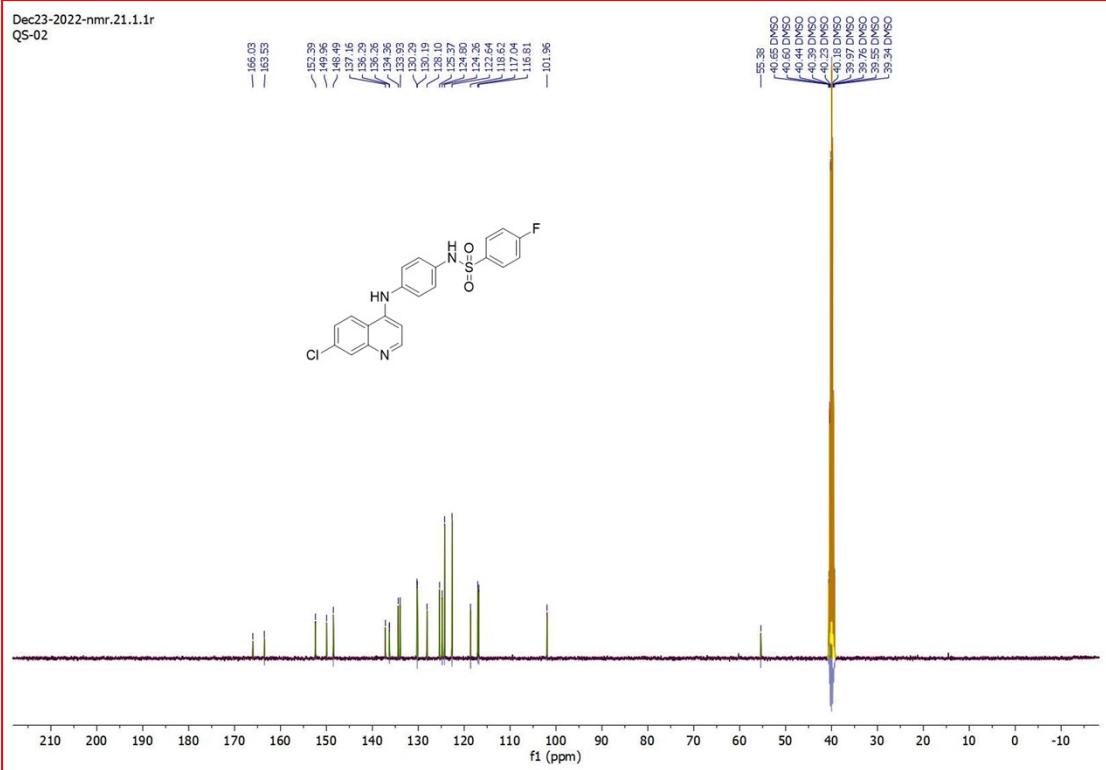


Figure-S4. ^{13}C NMR Spectrum of compound (QS-2)

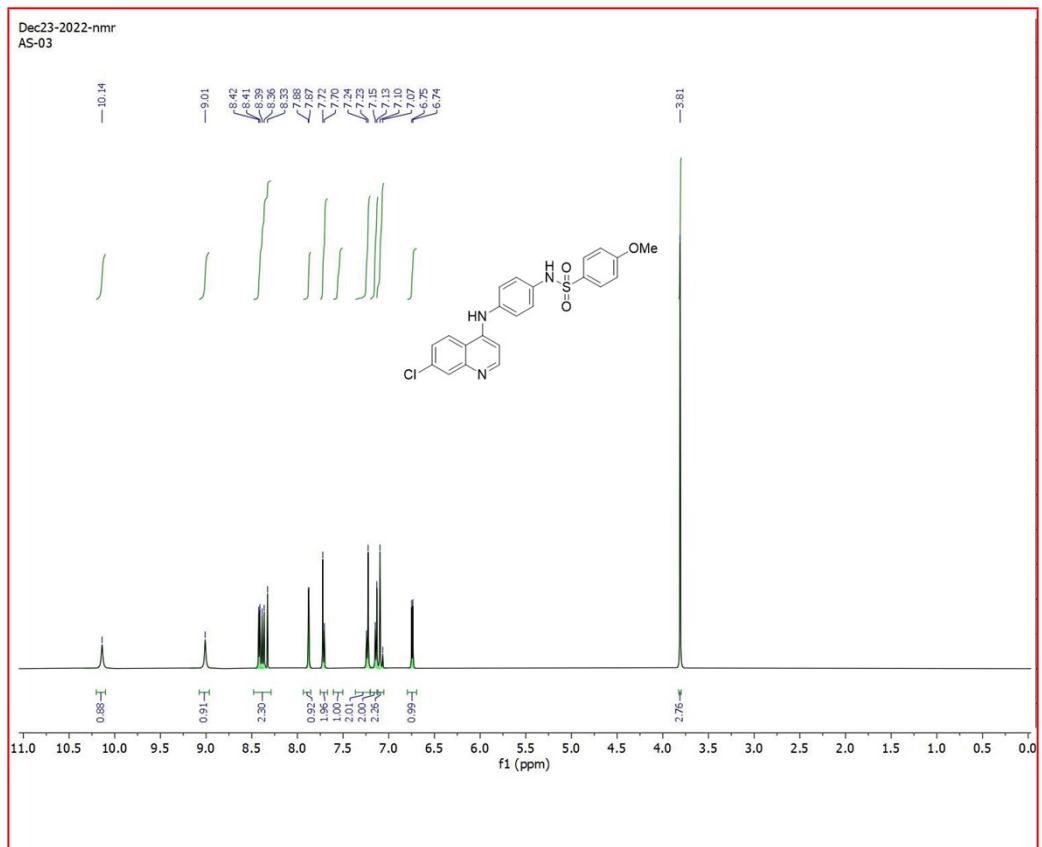


Figure-S5. ^1H NMR Spectrum of compound (QS-3)

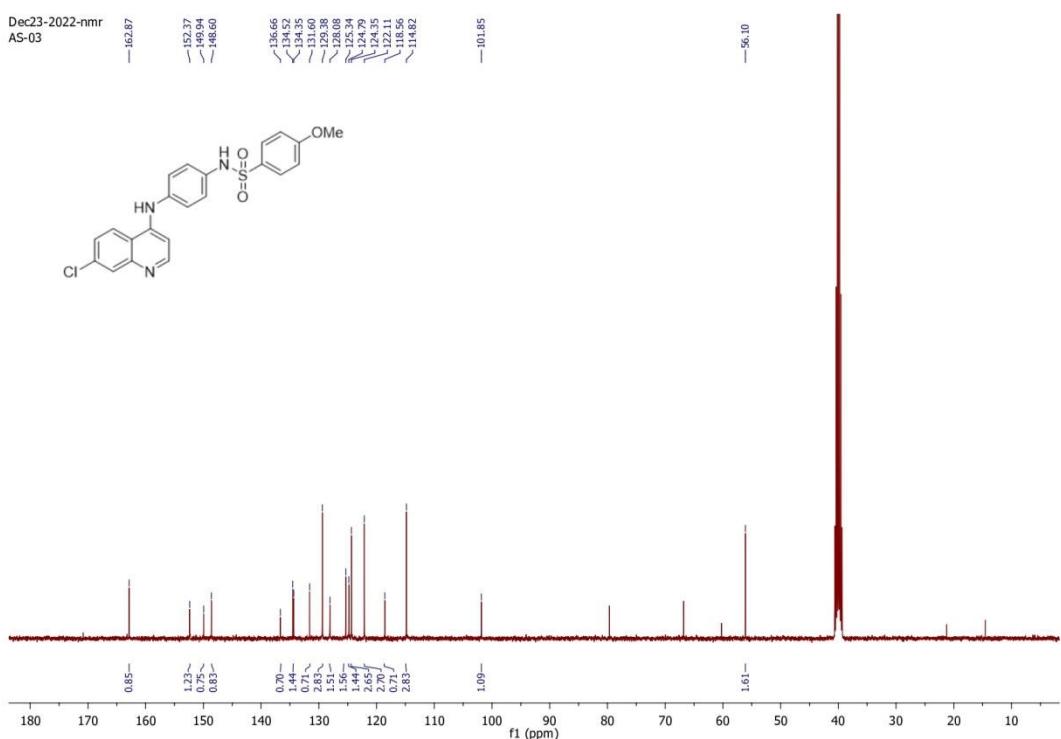


Figure-S6. ^{13}C NMR Spectrum of compound (QS-3)

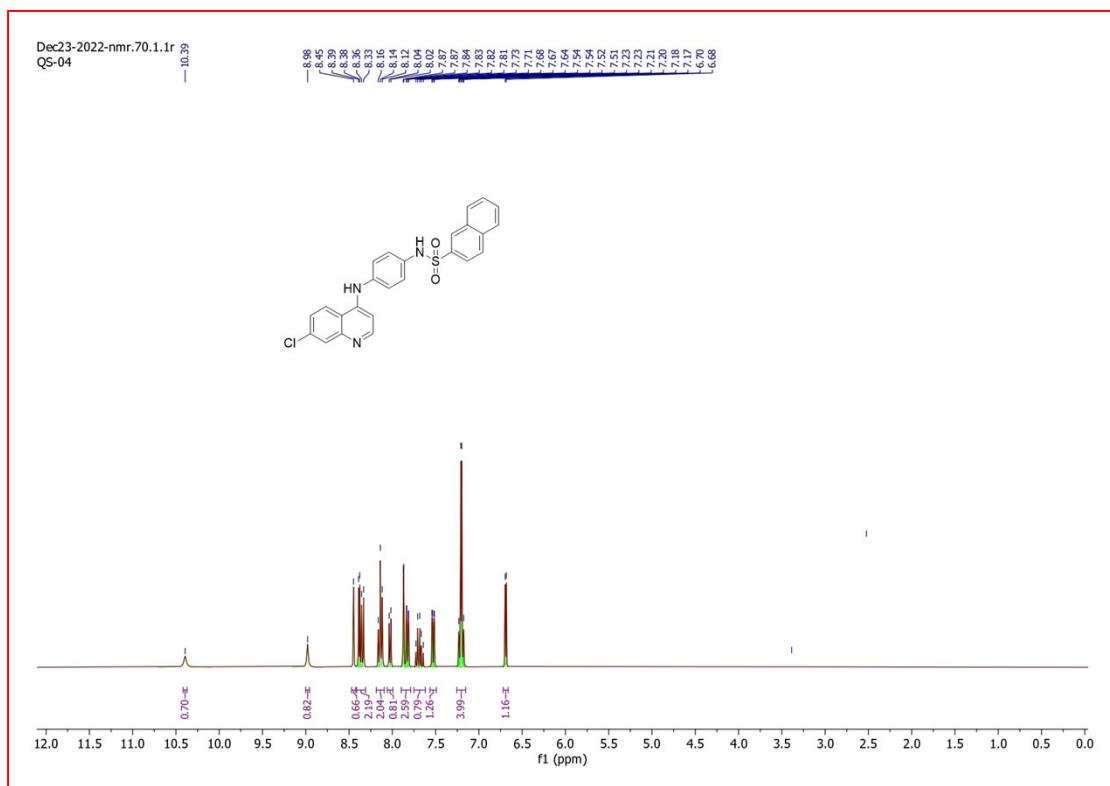


Figure.- S7. ^1H NMR Spectrum of compound (QS-4)

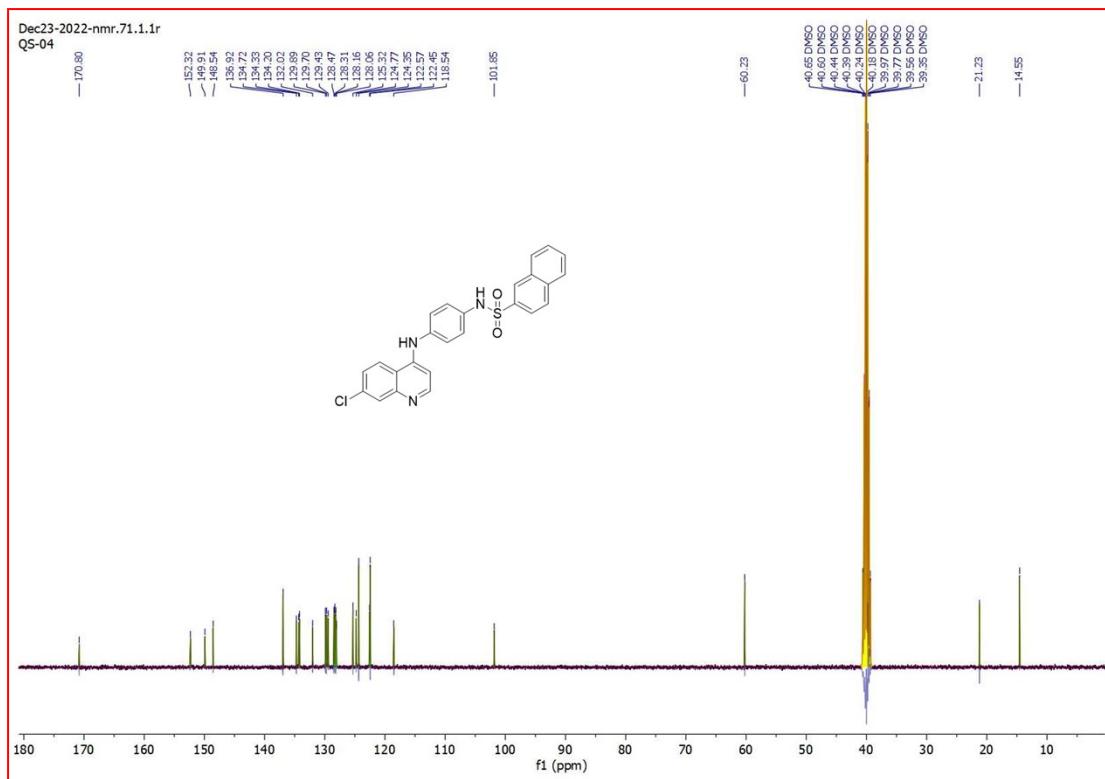


Figure.- S8. ^{13}C NMR Spectrum of compound (QS-4)

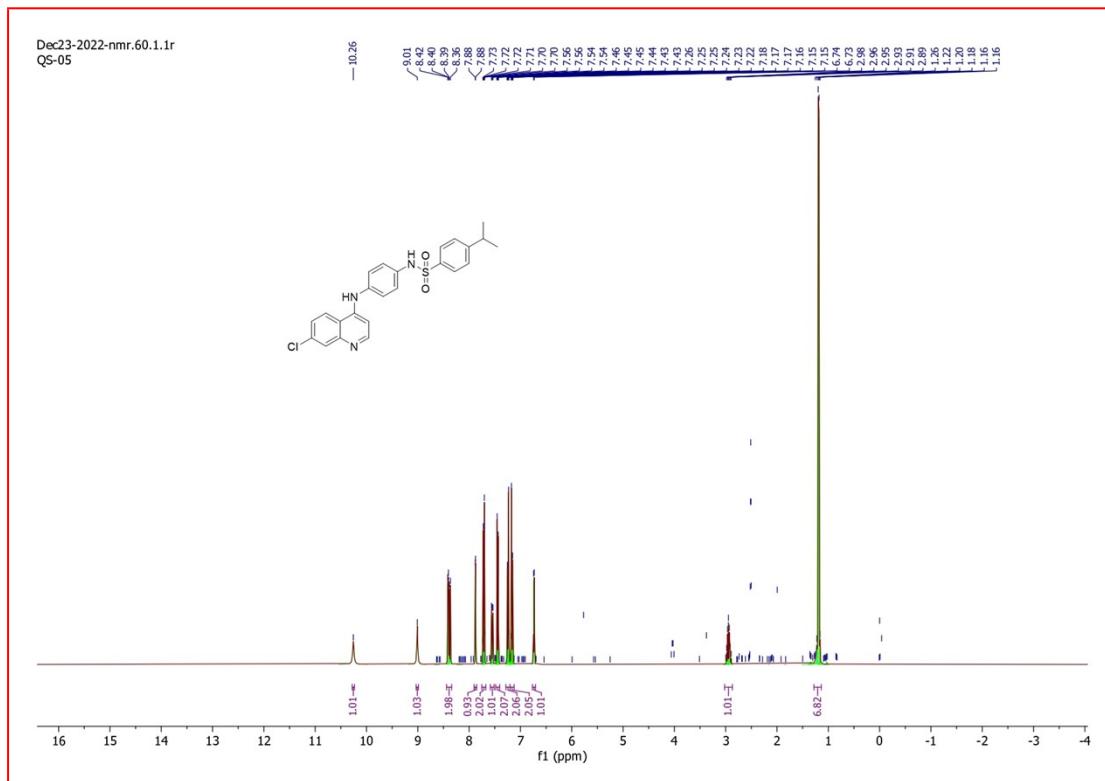


Figure.- S9. ^1H NMR Spectrum of compound (QS-5)

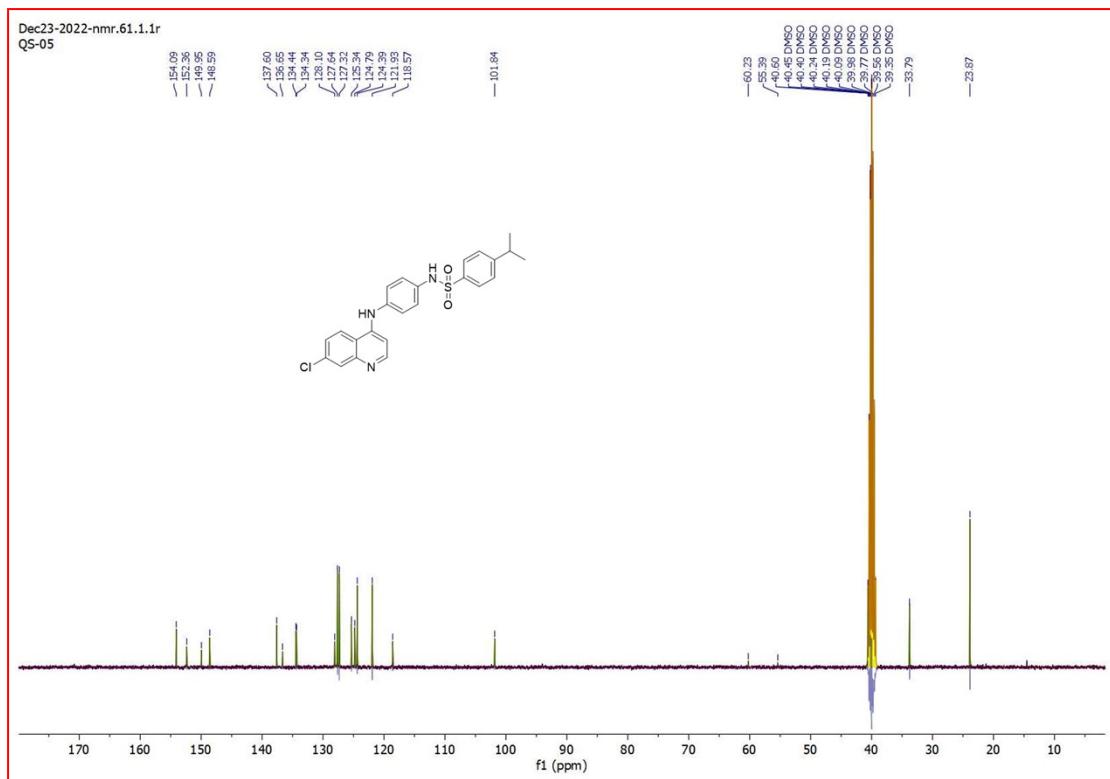


Figure.- S10. ¹³C NMR Spectrum of compound (QS-5)

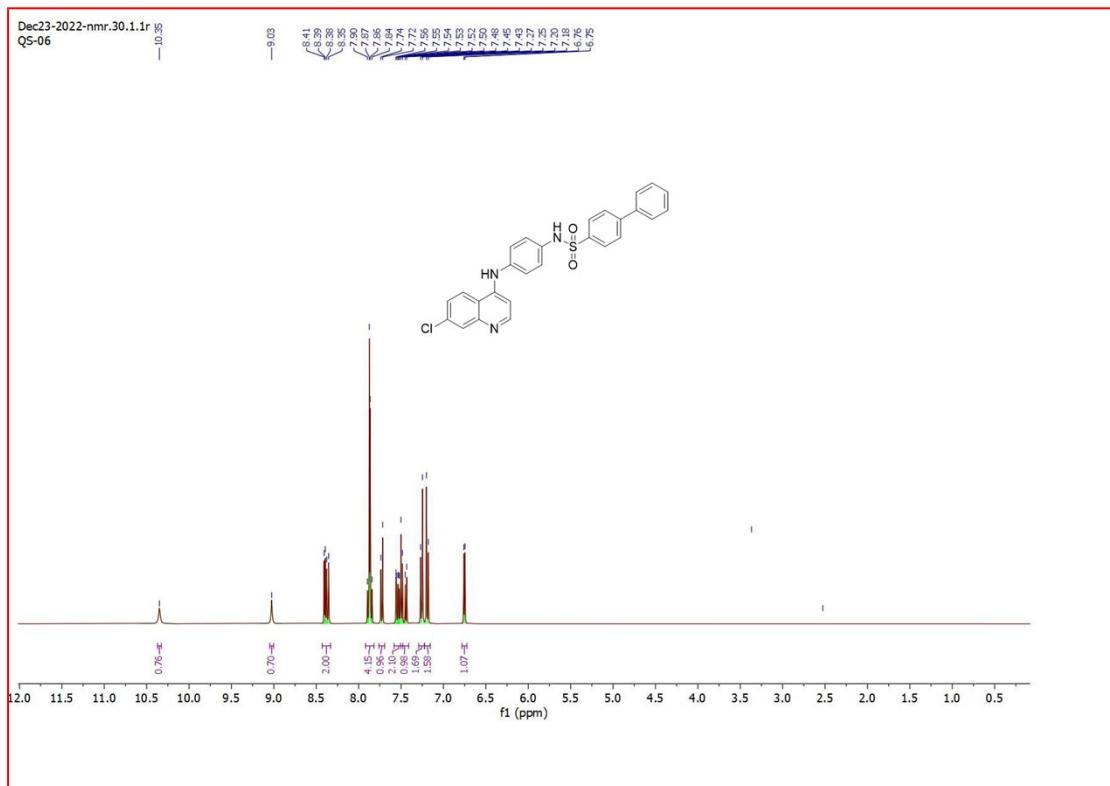


Figure.- S11. ¹H NMR Spectrum of compound (QS-6)

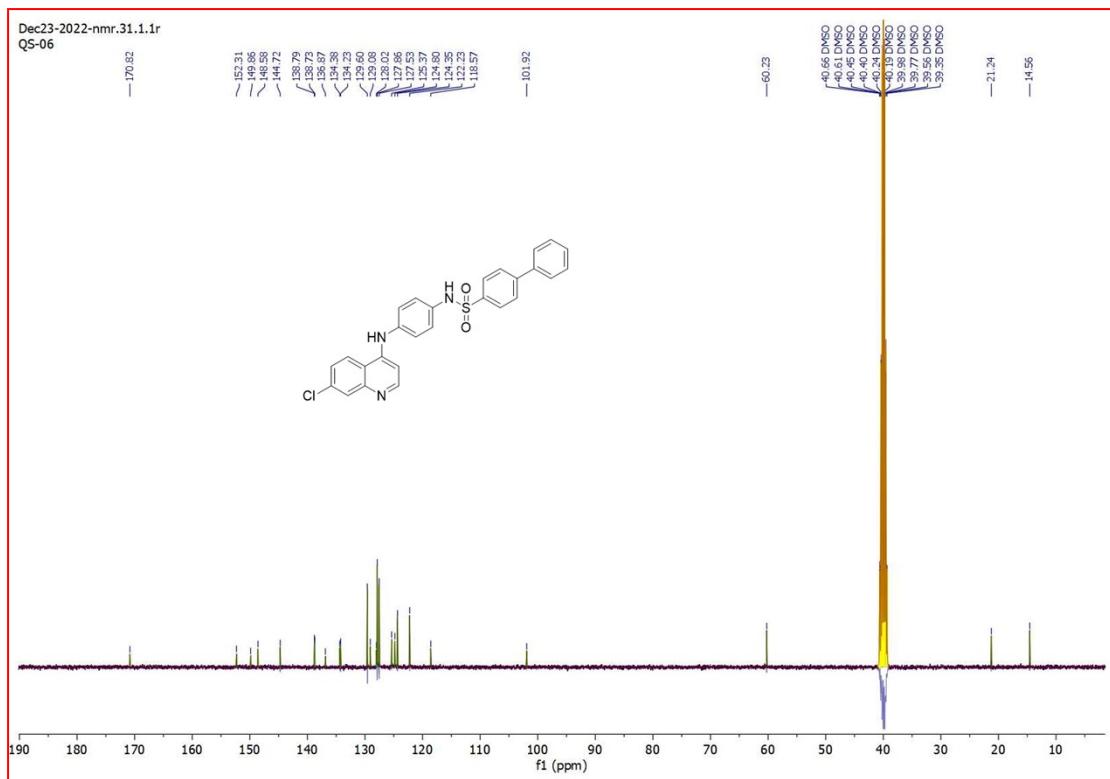


Figure.- S12. ^{13}C NMR Spectrum of compound (QS-6)

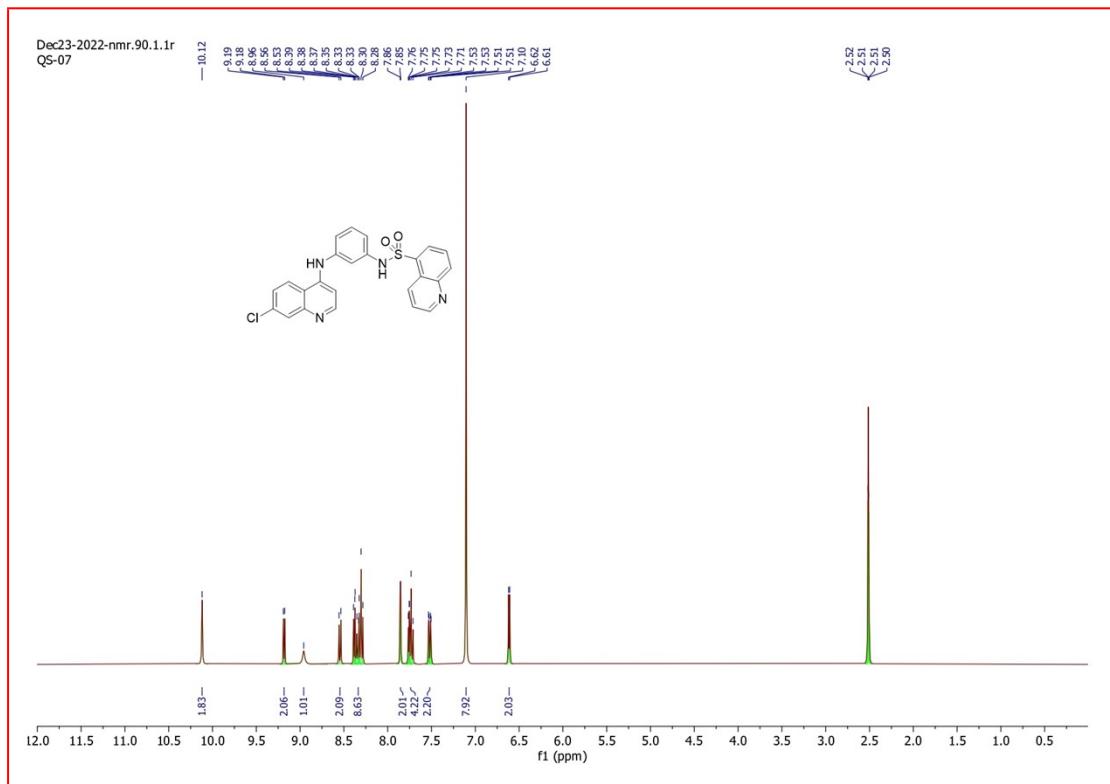


Figure.- S13. ^1H NMR Spectrum of compound (QS-7)

Dec23-2022-nmr.91.1.1r
QS-07

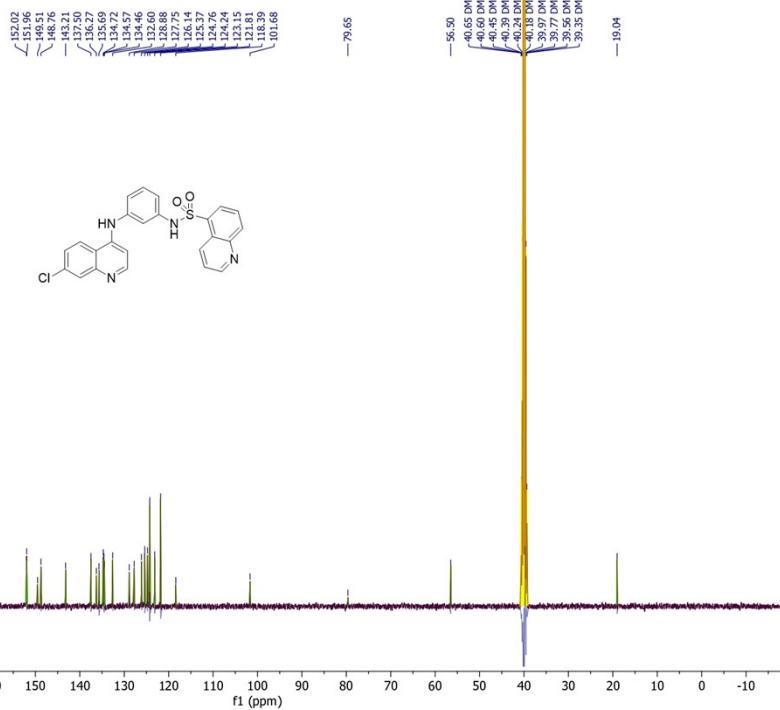


Figure.- S14. ¹³C NMR Spectrum of compound (QS-7)

Dec23-2022-nmr
QS-08

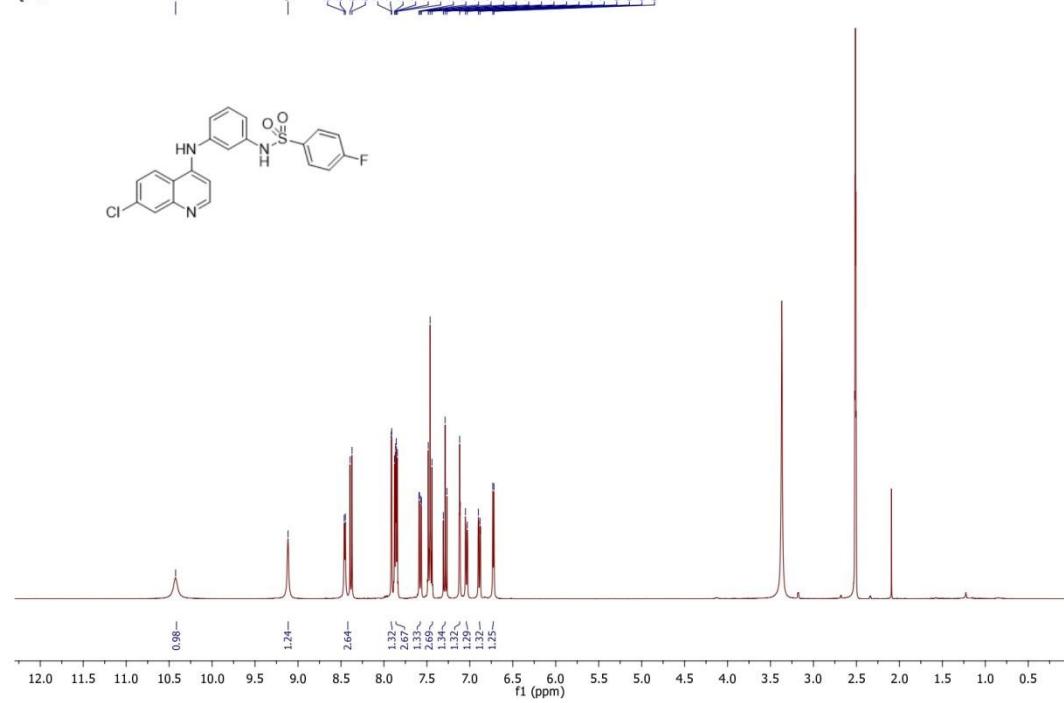


Figure.- S15. ¹H NMR Spectrum of compound (QS-8)

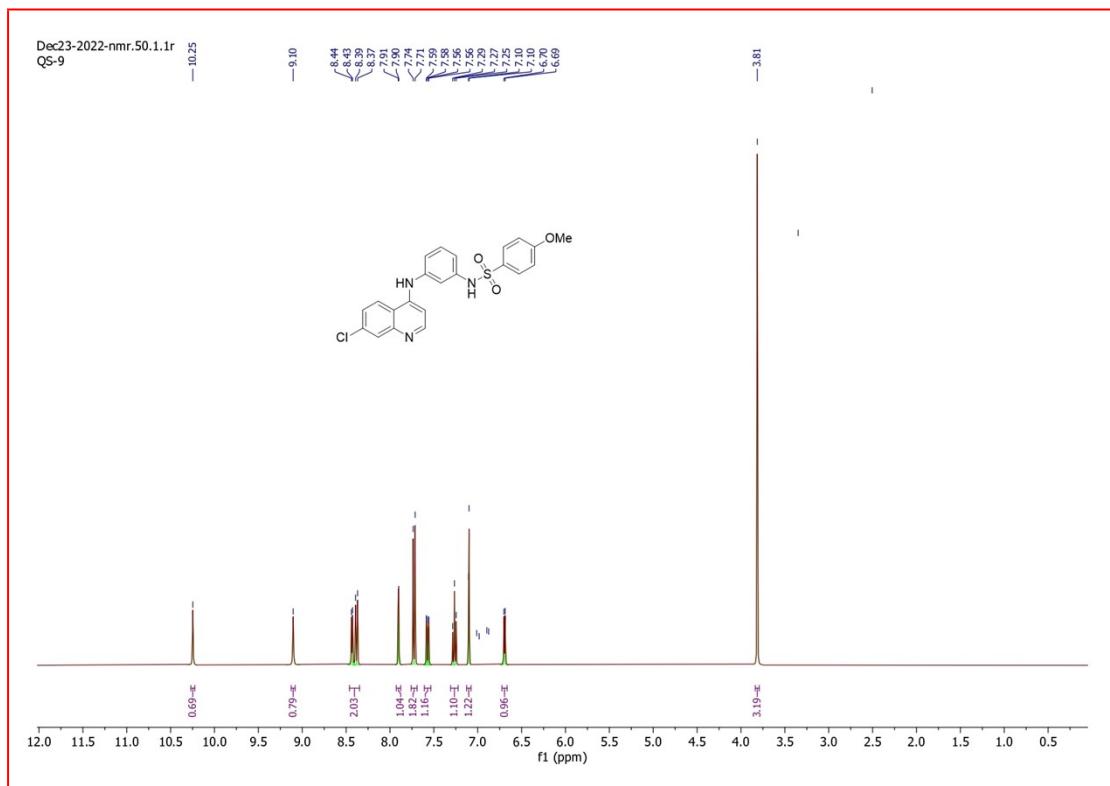
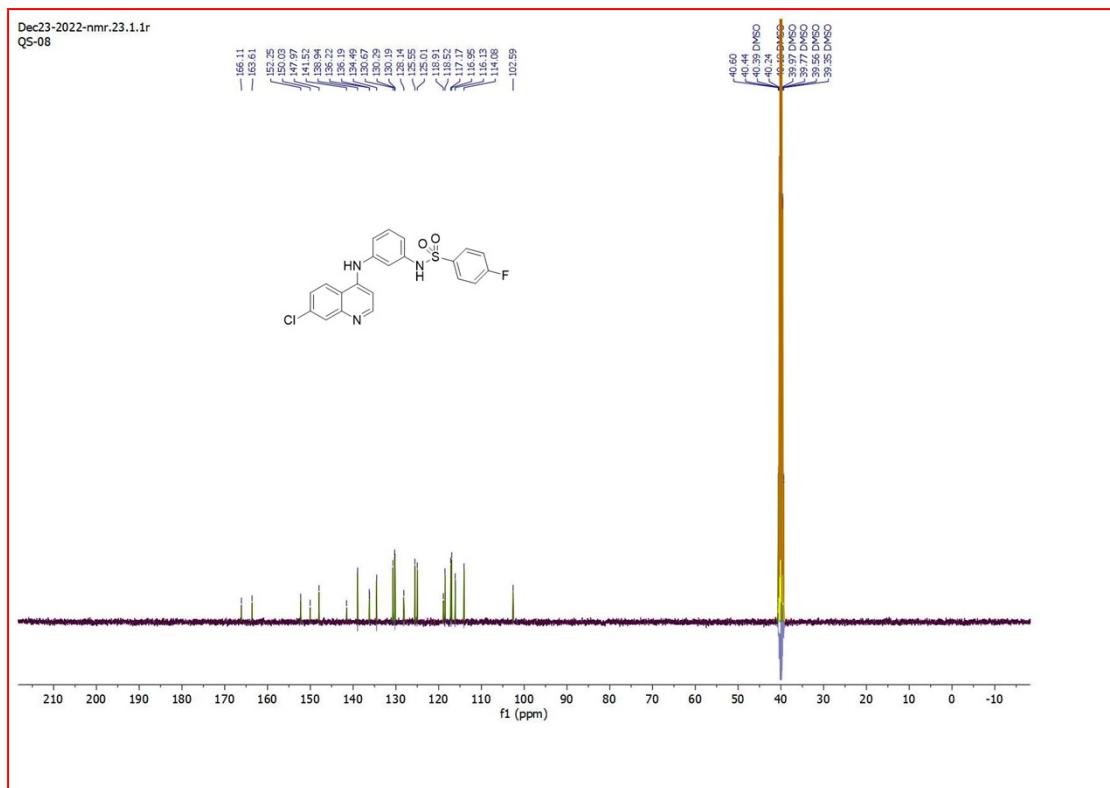


Figure.- S17. ^1H NMR Spectrum of compound (QS-9)

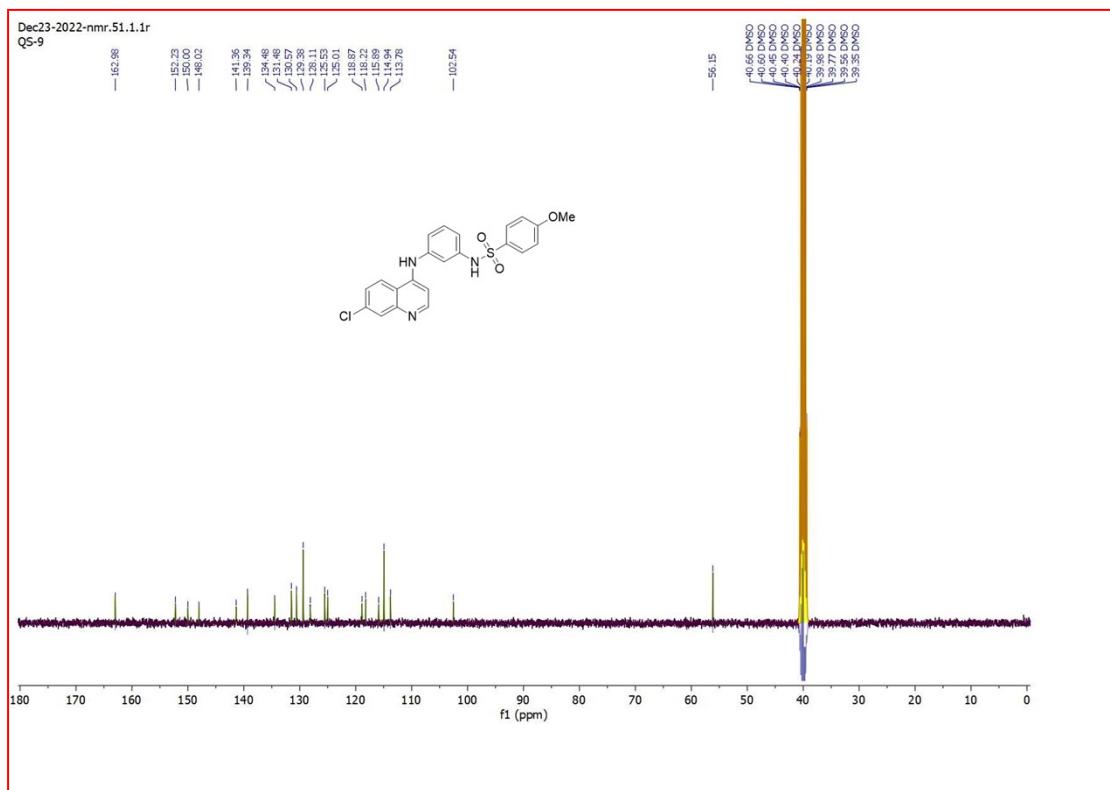


Figure.- S18. ^{13}C NMR Spectrum of compound (QS-9)

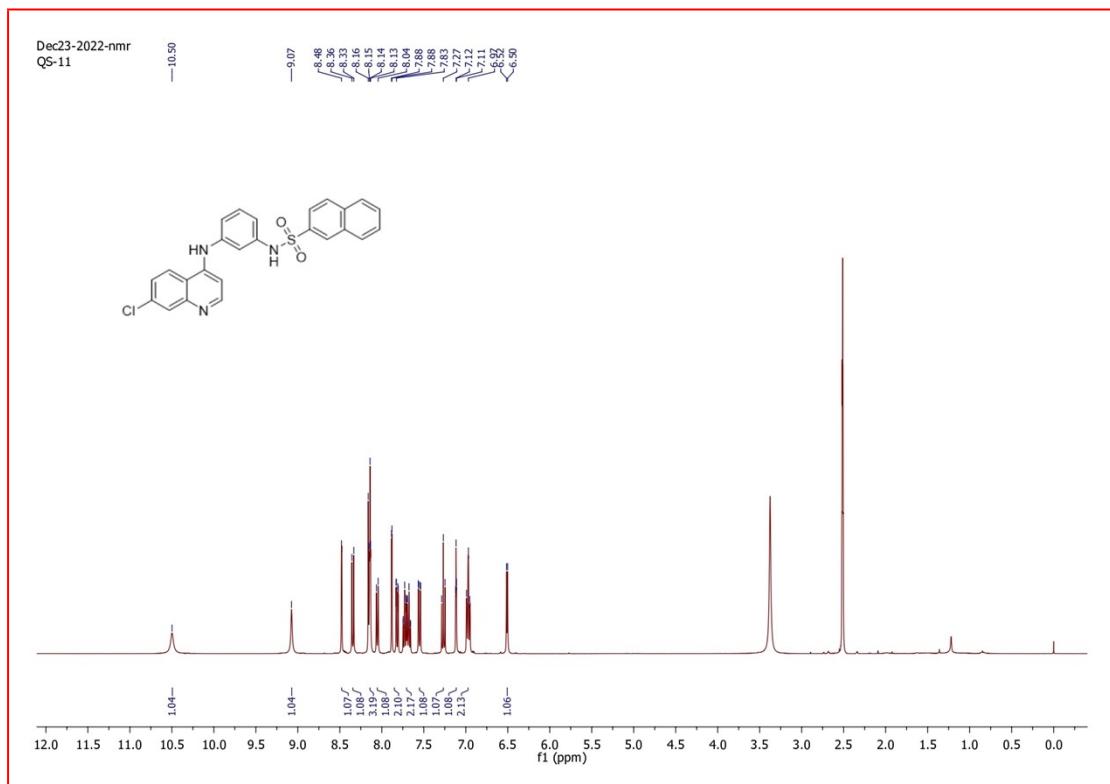


Figure.- S19. ^1H NMR Spectrum of compound (QS-10)

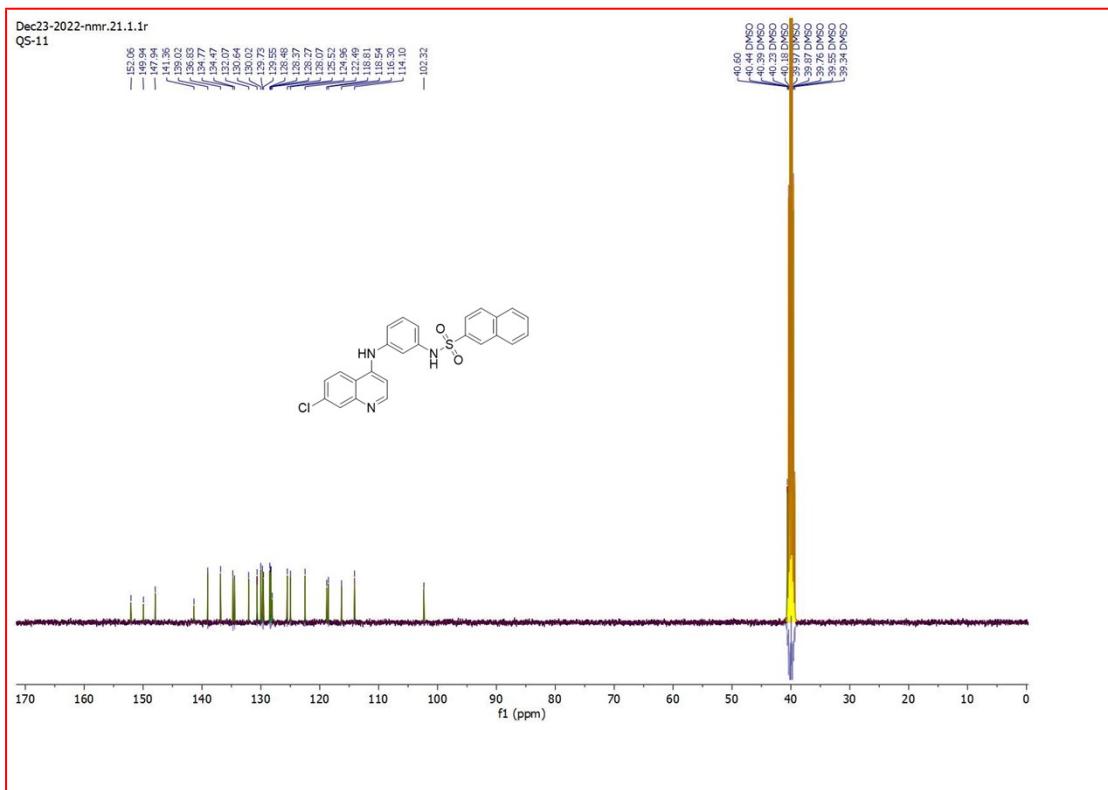


Figure.- S20. ^{13}C NMR Spectrum of compound (QS-10)

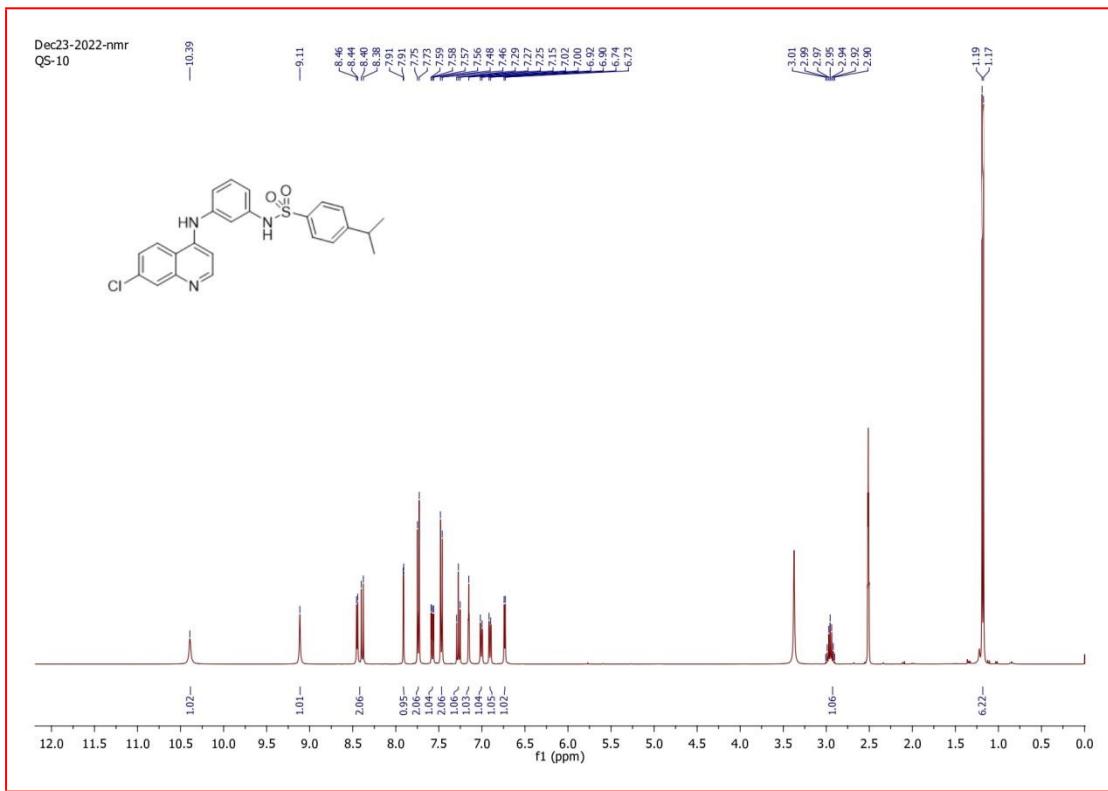


Figure.- S21. ^1H NMR Spectrum of compound (QS-11)

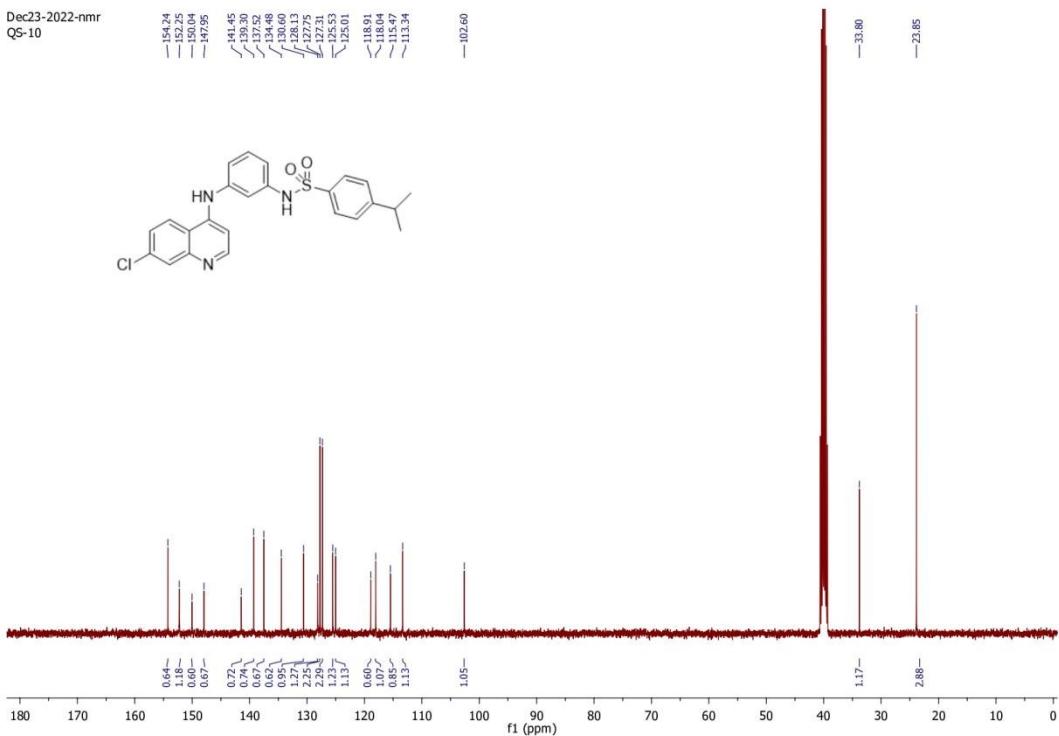


Figure.- S22. ^{13}C NMR Spectrum of compound (QS-11)

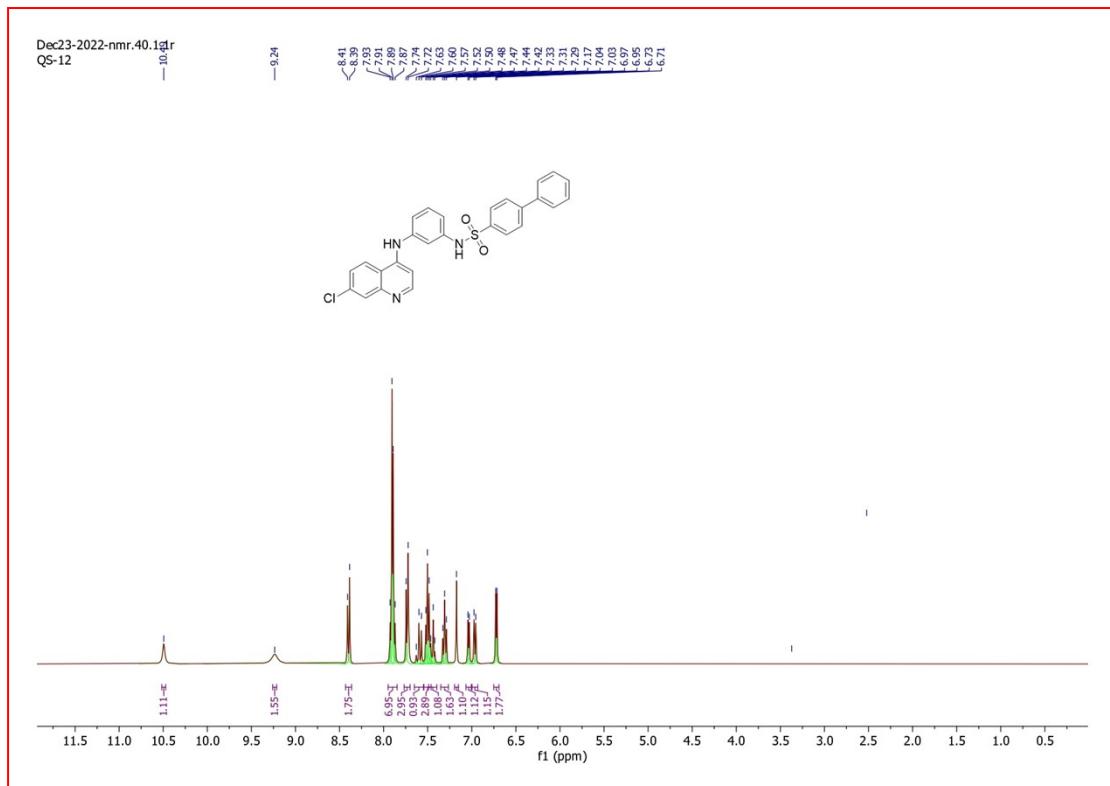


Figure.- S23. ^1H NMR Spectrum of compound (QS-12)

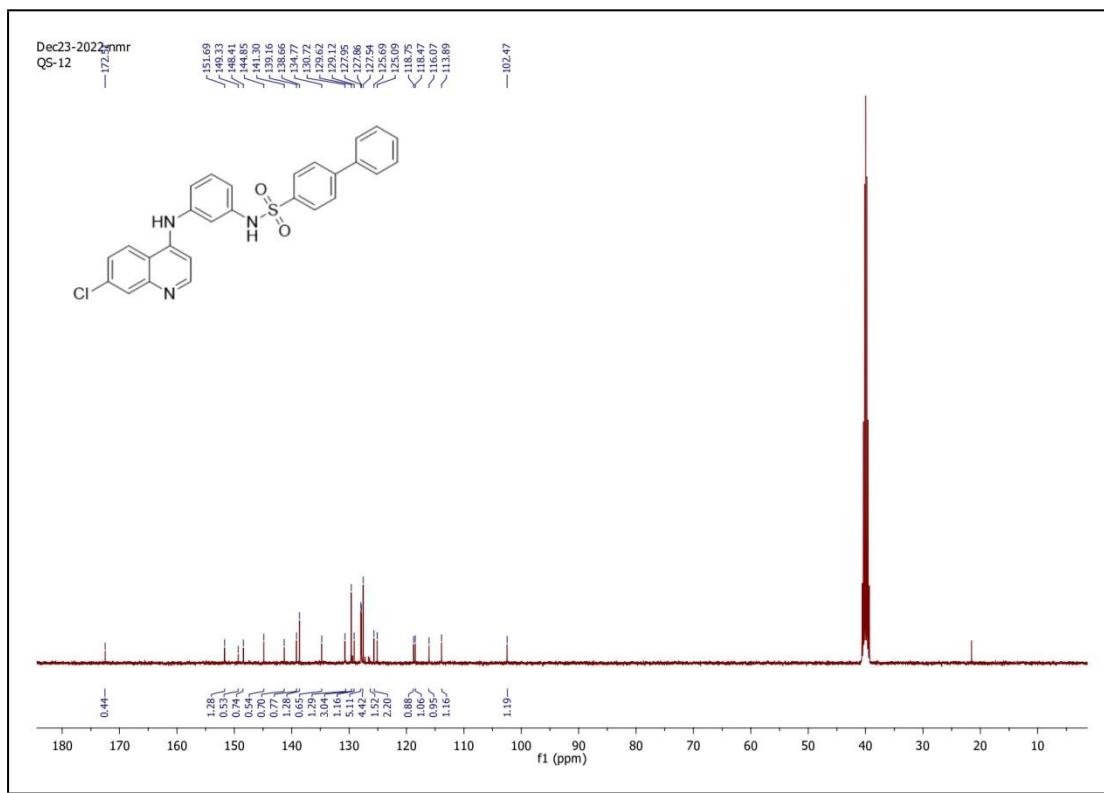


Figure - S24. ^{13}C NMR Spectrum of compound (QS-12)

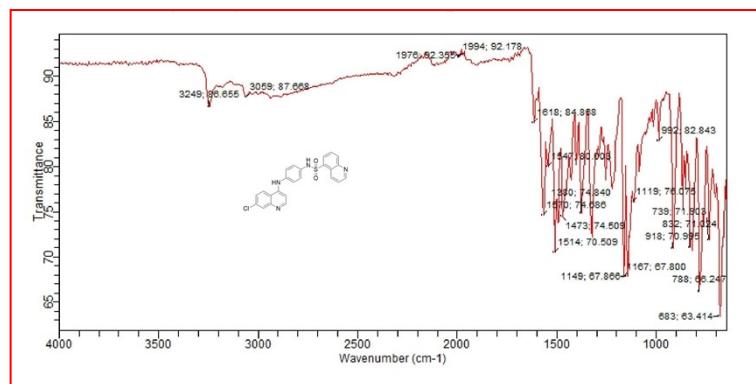


Figure - S25. FTIR Spectrum of compound (QS-1)

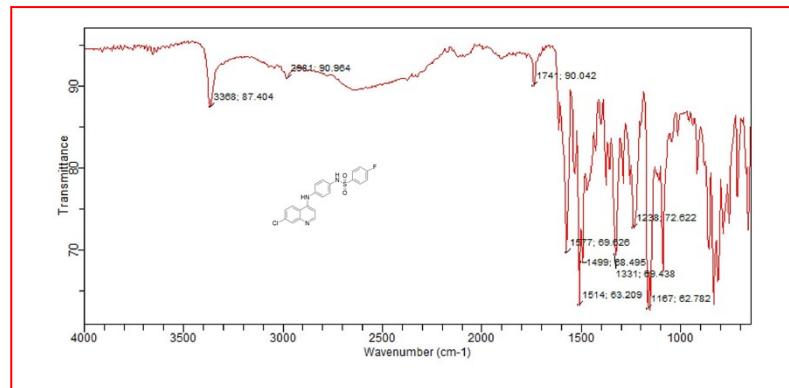


Figure - S26. FTIR Spectrum of compound (QS-2)

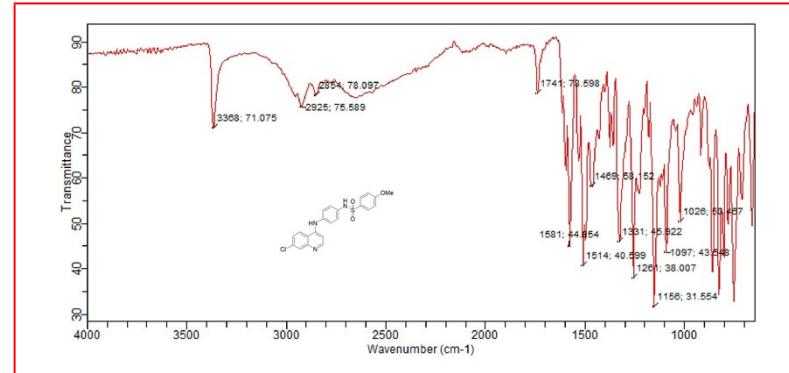


Figure - S27. FTIR Spectrum of compound (QS-3)

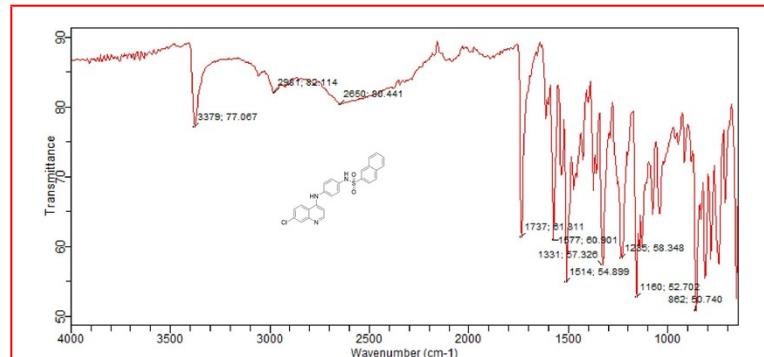


Figure - S28. FTIR Spectrum of compound (QS-4)

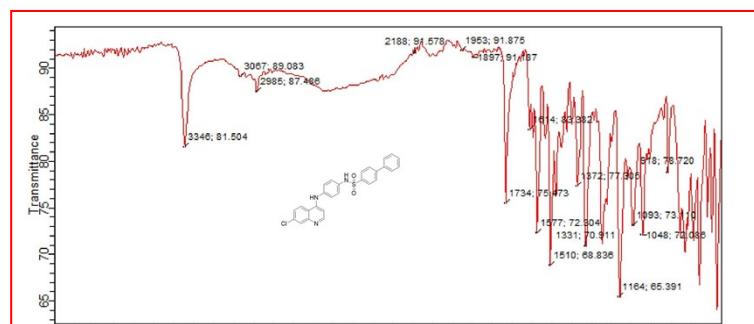


Figure - S29. FTIR Spectrum of compound (QS-5)

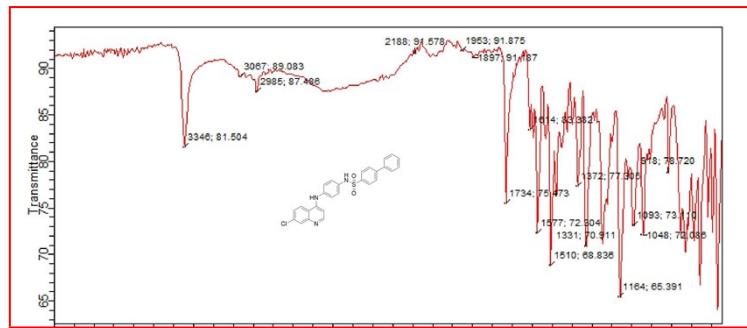


Figure - S30. FTIR Spectrum of compound (QS-6)

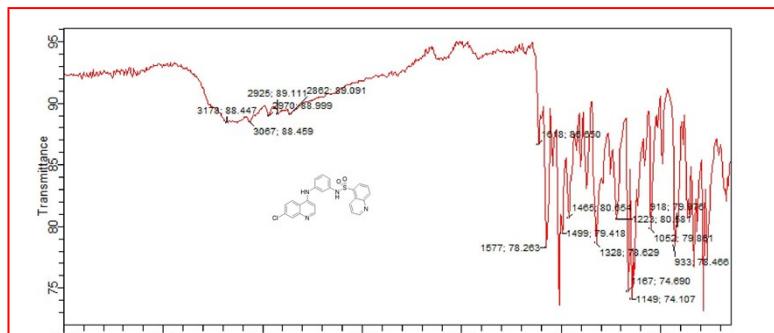


Figure - S31. FTIR Spectrum of compound (QS-7)

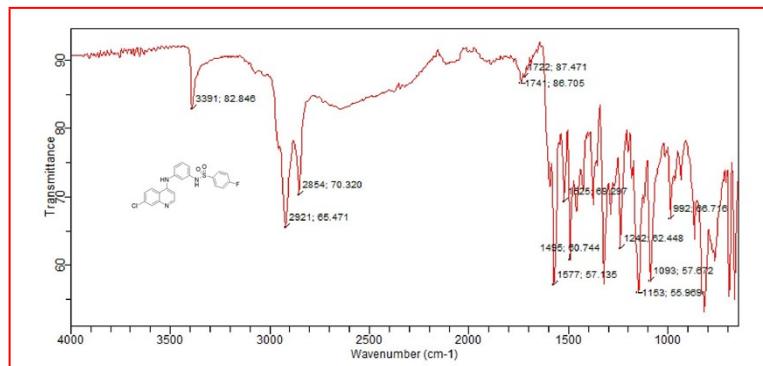


Figure - S32. FTIR Spectrum of compound (QS-8)

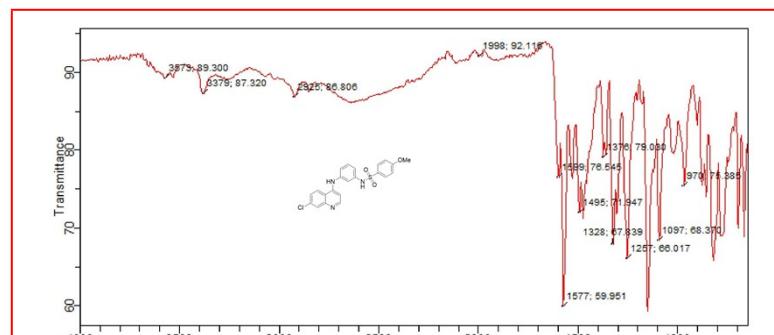


Figure - S33. FTIR Spectrum of compound (QS-9)

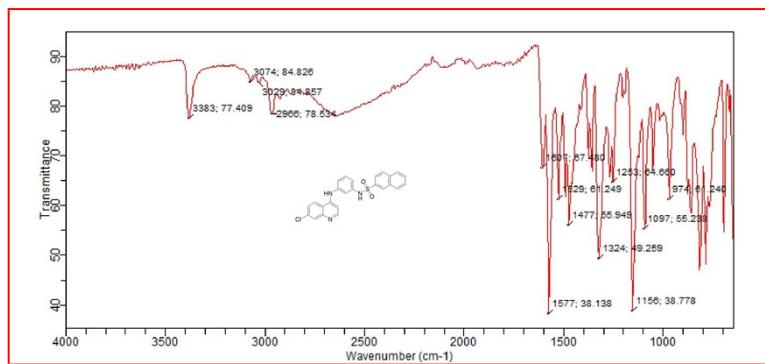


Figure - S34. FTIR Spectrum of compound (QS-10)

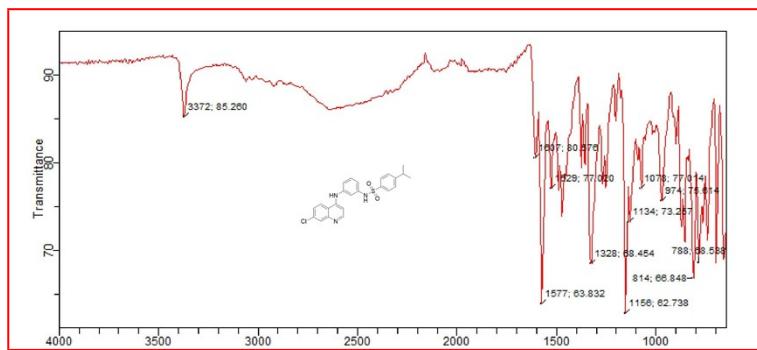


Figure - S35. FTIR Spectrum of compound (QS-11)

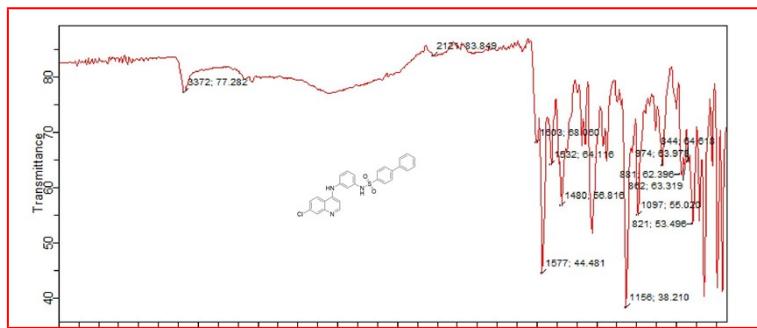


Figure - S36. FTIR Spectrum of compound (QS-12)

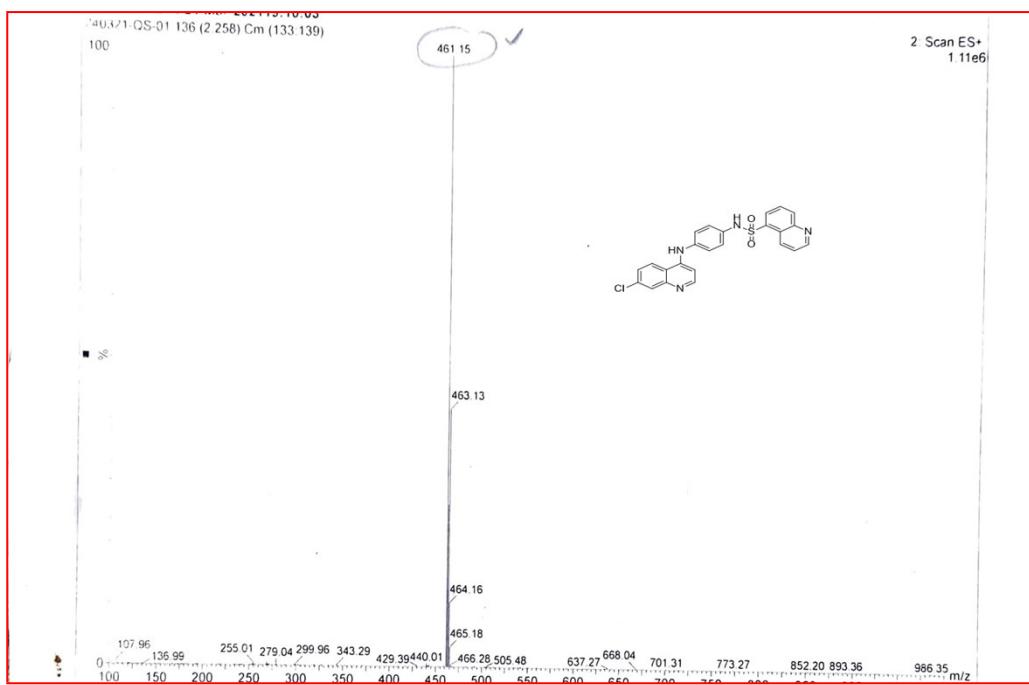


Figure - S37. Mass spectrum of compound (QS-1)

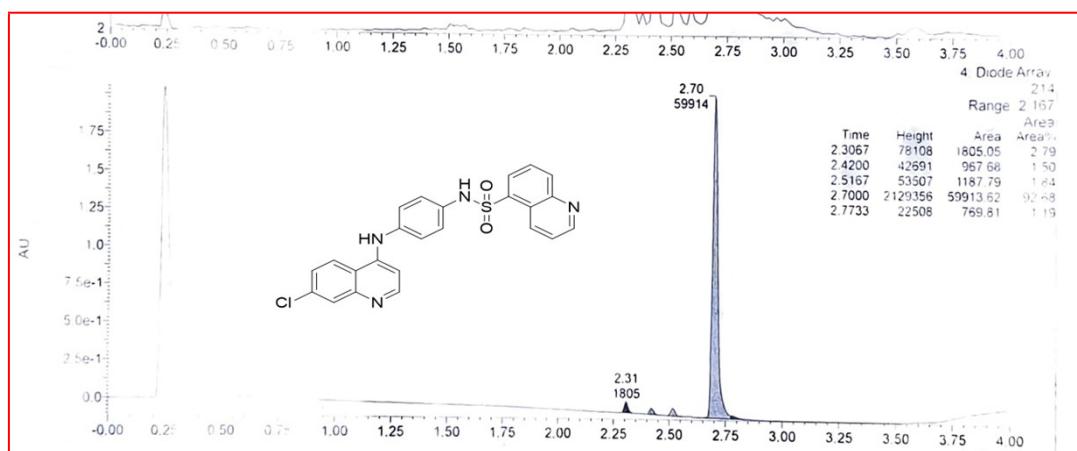


Figure - S38. Purity of compound (QS-1)

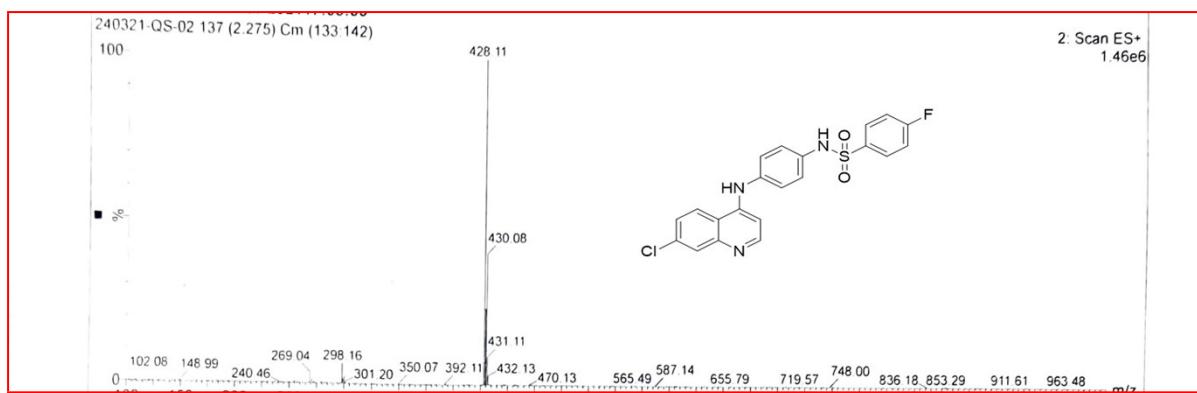


Figure - S39. Mass spectrum of compound (QS-2)

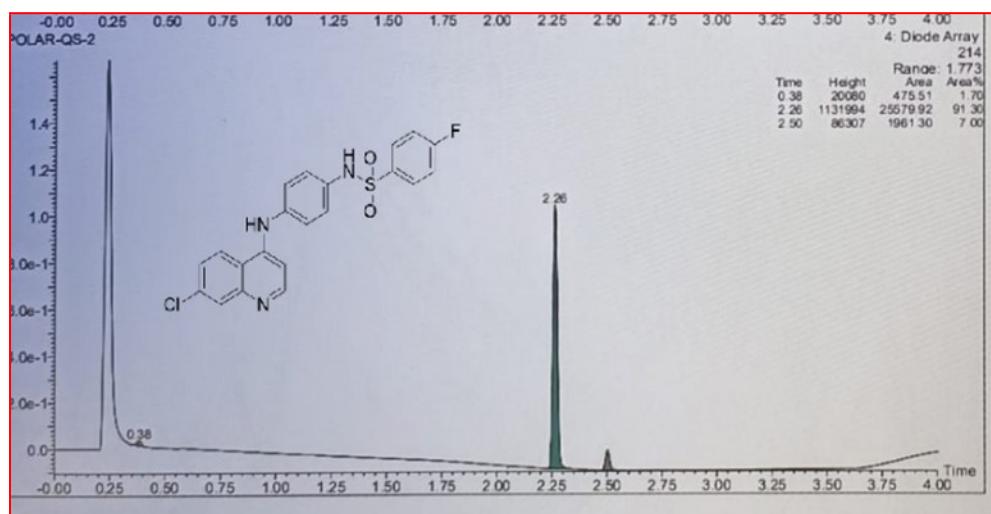


Figure - S40. Purity of compound (QS-2)

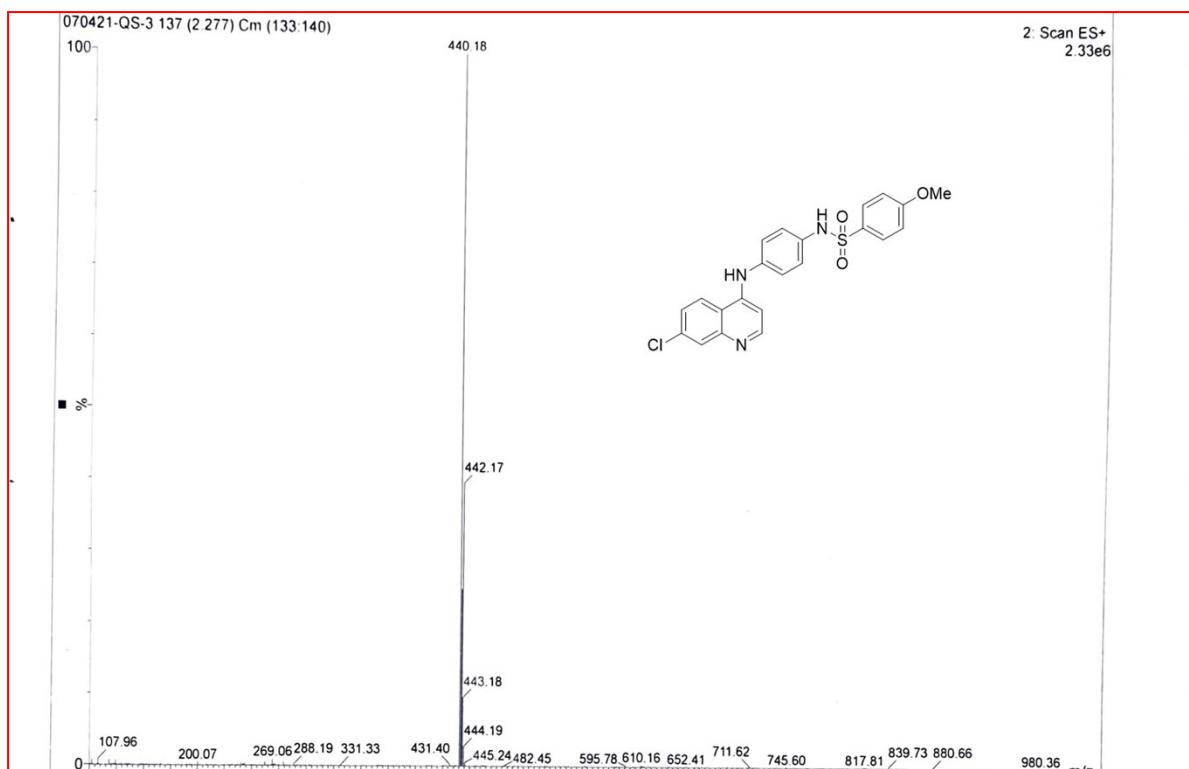


Figure - S41. Mass spectrum of compound (QS-3)

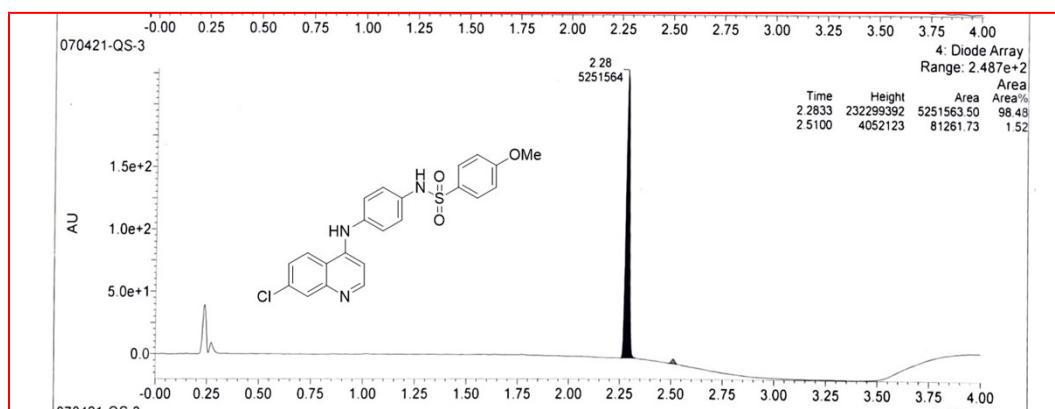


Figure - S42. Purity of compound (QS-3)

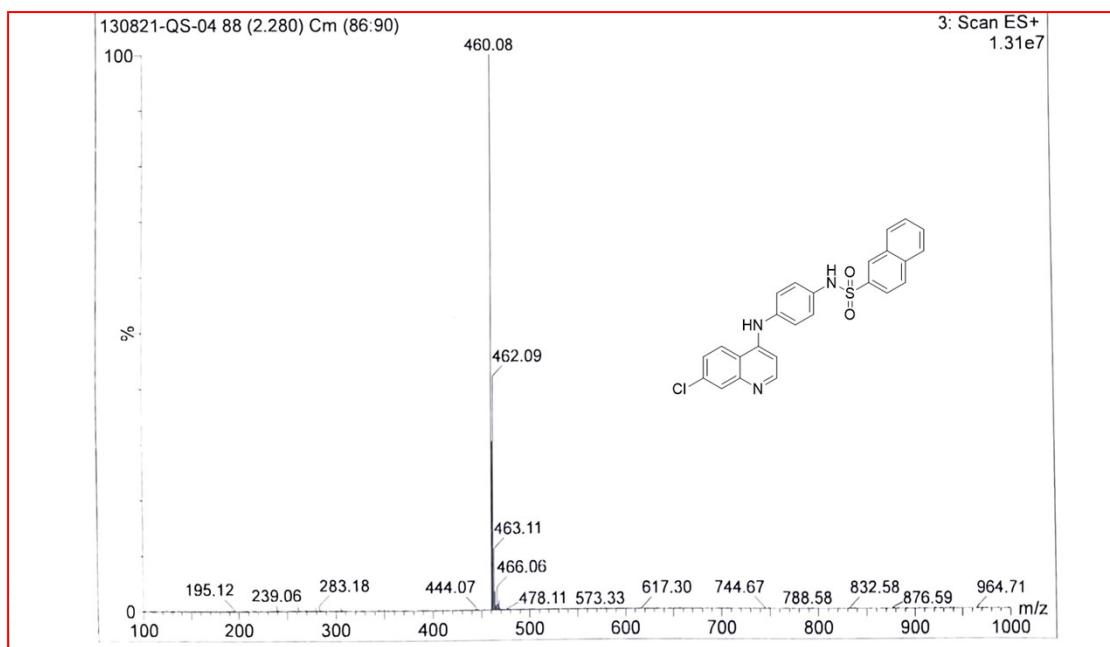


Figure - S43. Mass spectrum of compound (QS-4)

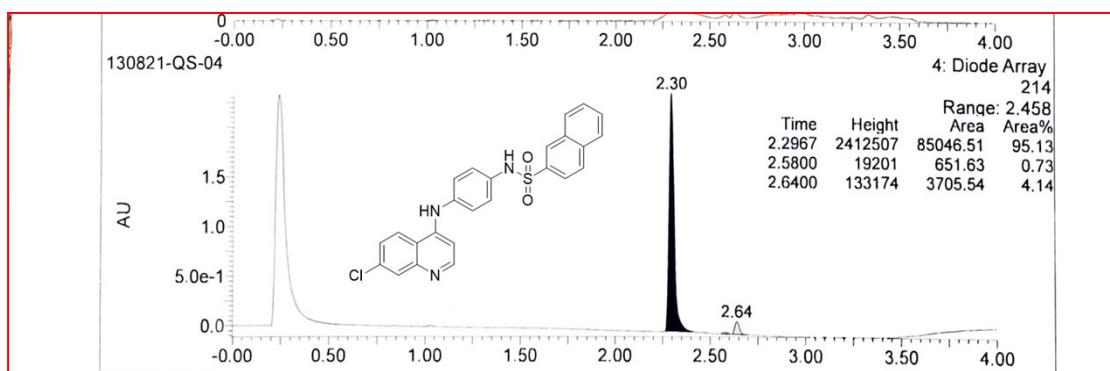


Figure - S44. Purity of compound (QS-4)

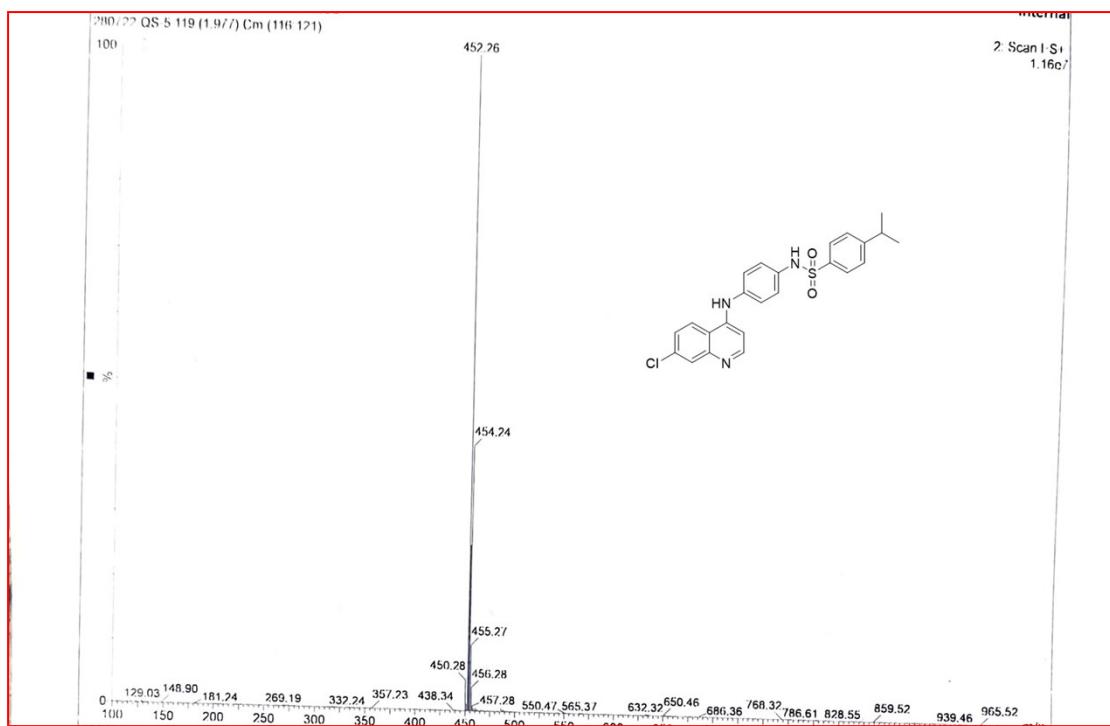


Figure - S45. Mass spectrum of compound (QS-5)

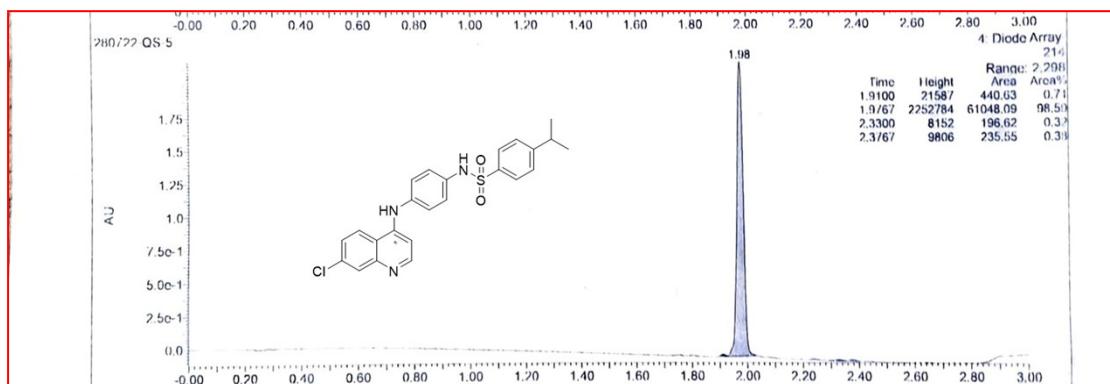


Figure - S46. Purity of compound (QS-5)

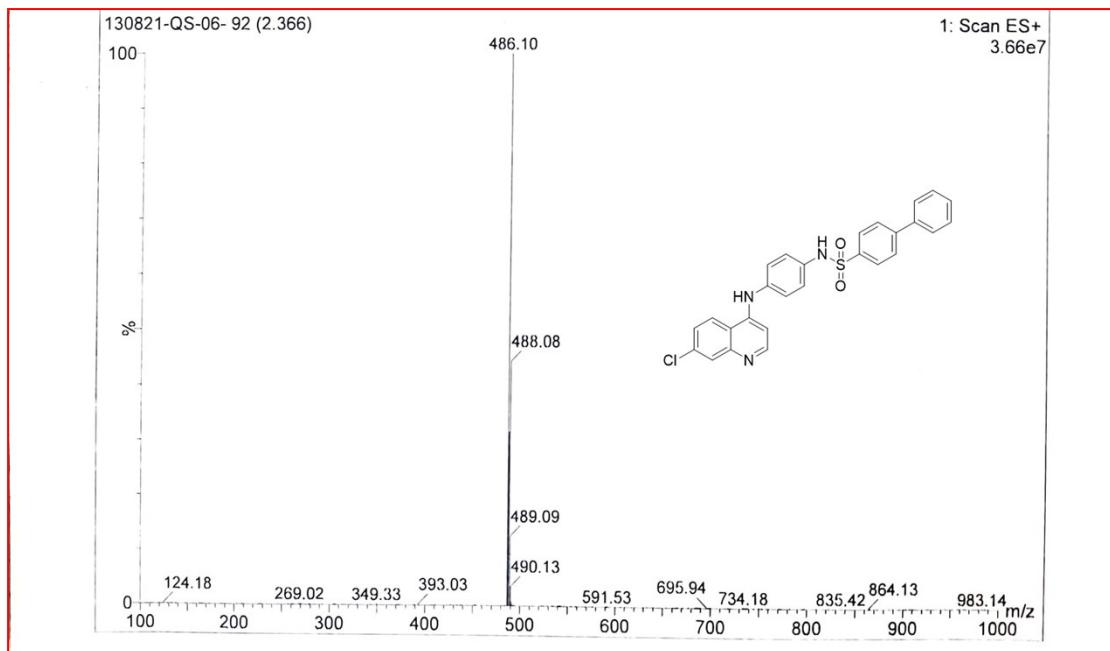


Figure - S47. Mass spectrum of compound (QS-6)

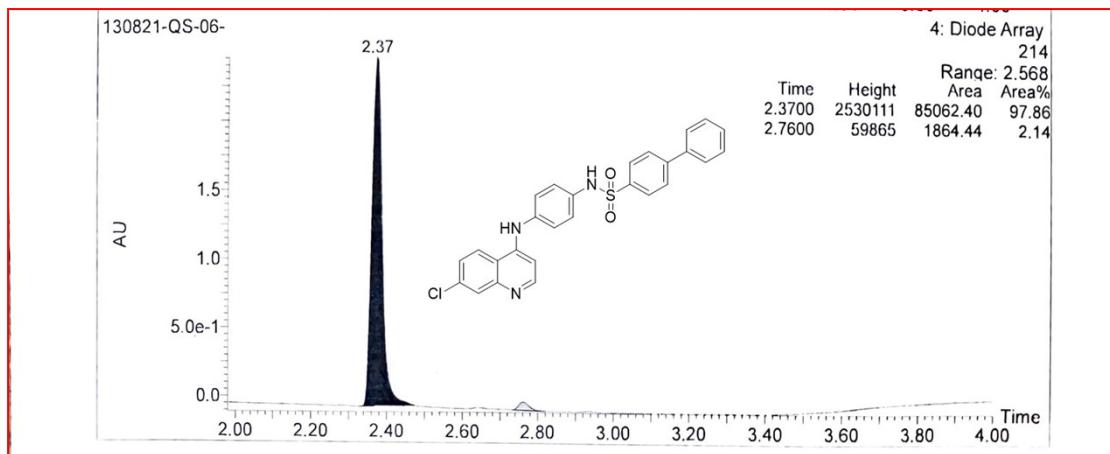


Figure - S48. Purity of compound (QS-6)

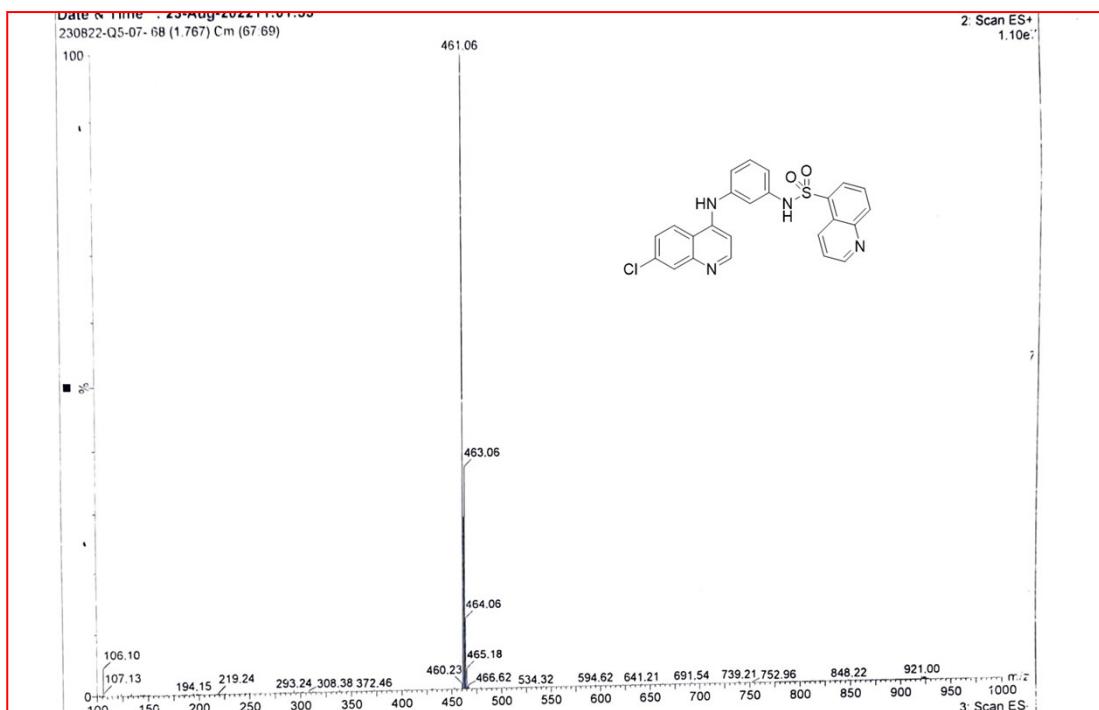


Figure - S49. Mass spectrum of compound (QS-7)

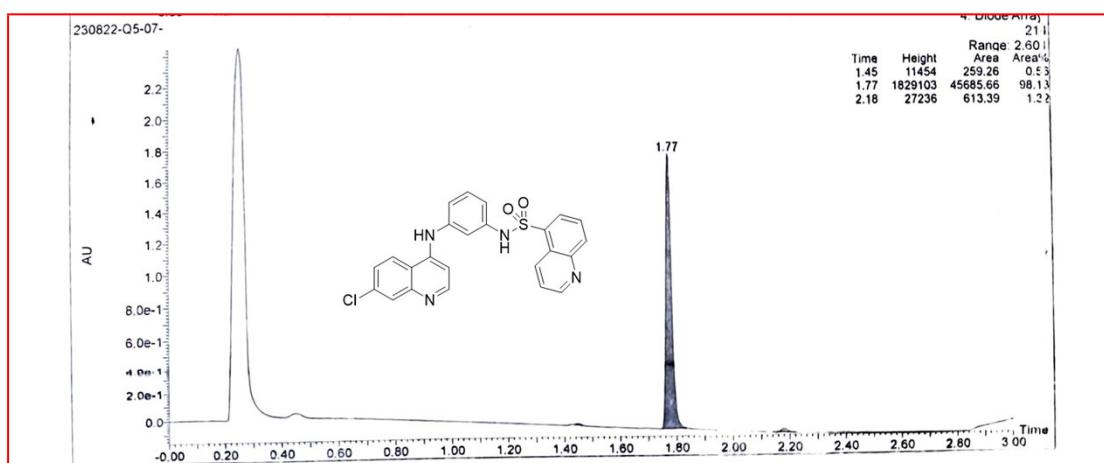


Figure - S50. Purity of compound (QS-7)

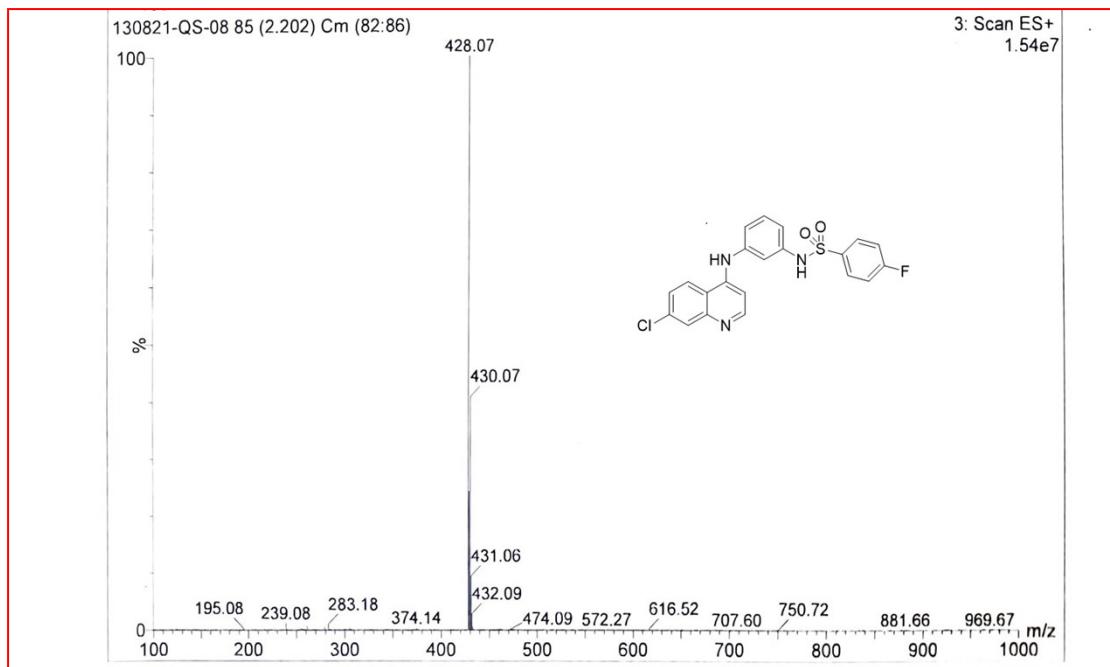


Figure - S51. Mass spectrum of compound (QS-8)

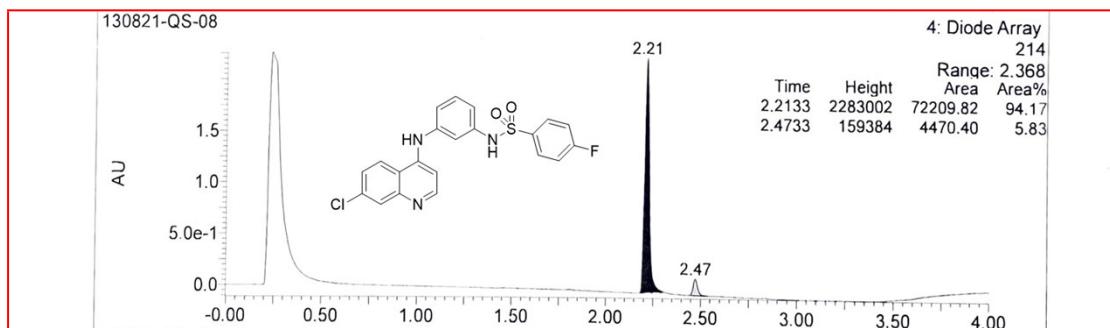


Figure - S52. Purity of compound (QS-8)

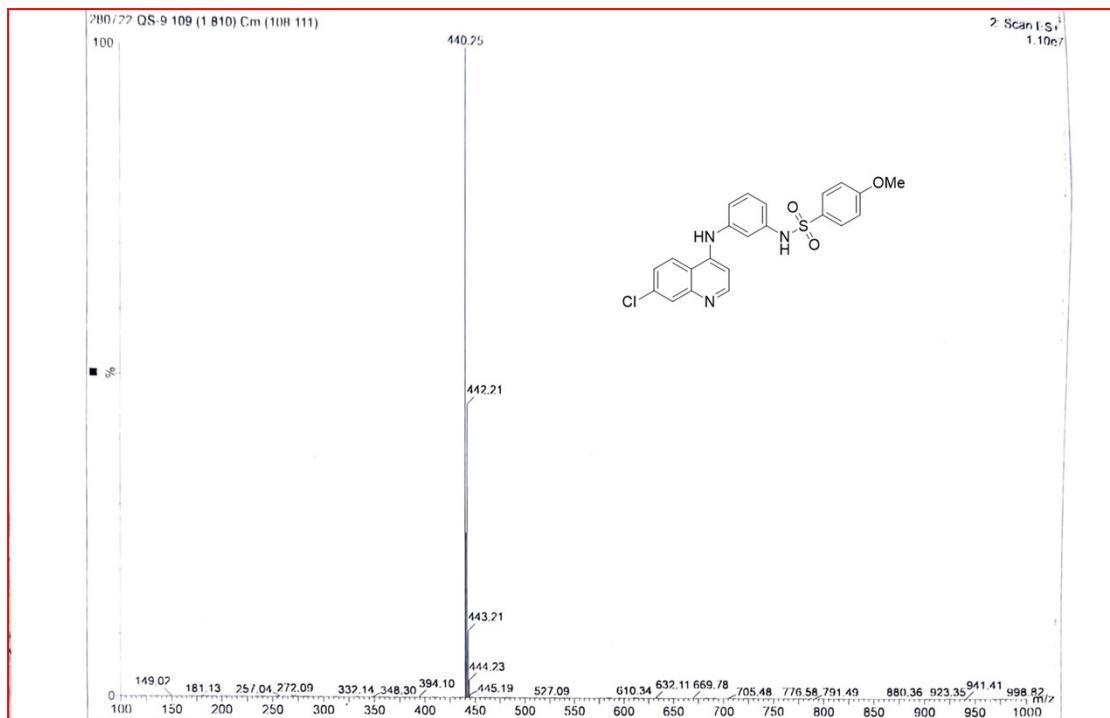


Figure - S53. Mass spectrum of compound (QS-9)

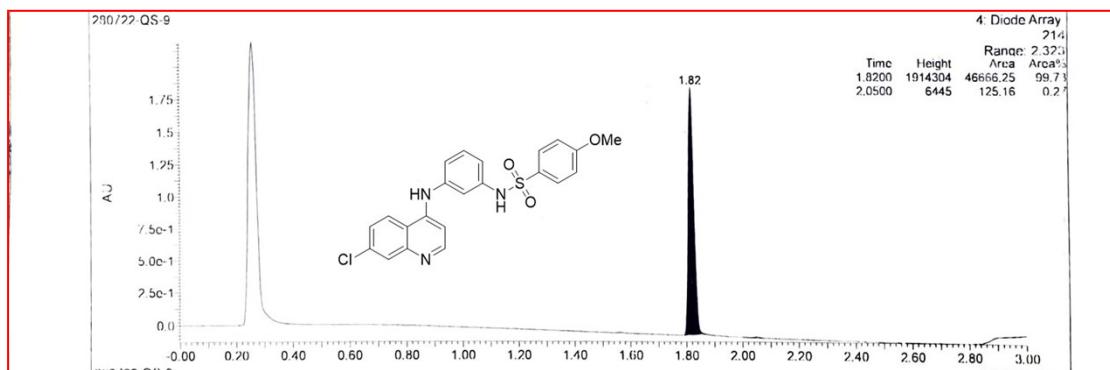


Figure - S54. Purity of compound (QS-9)

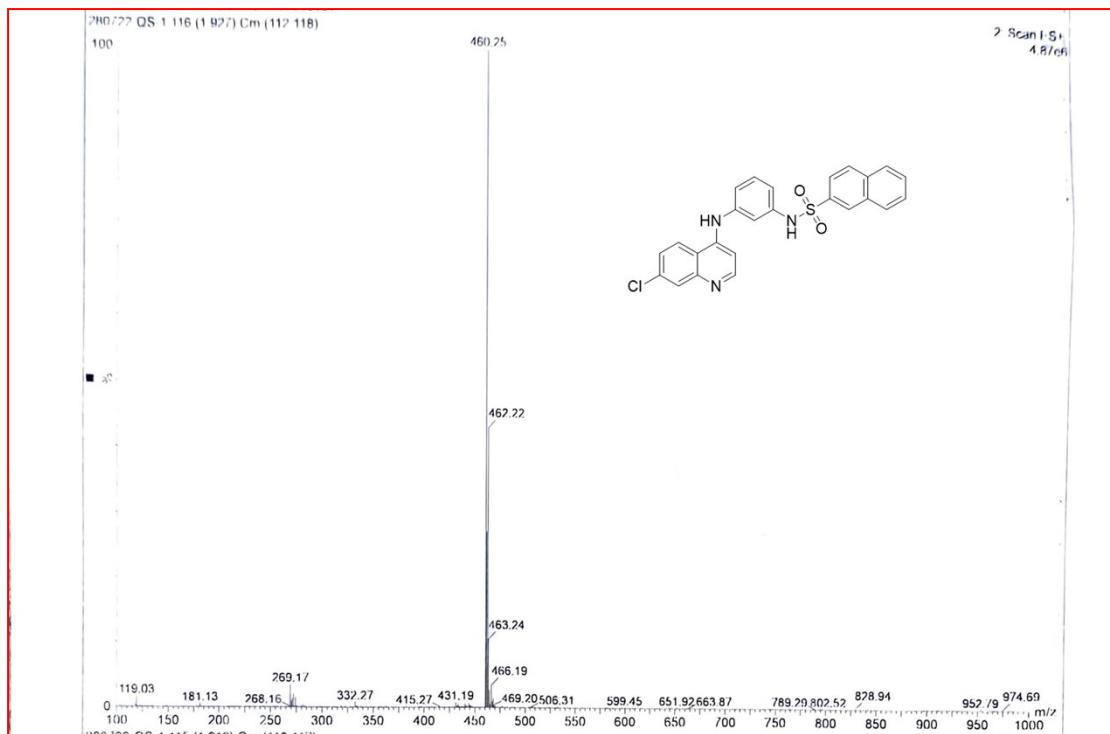


Figure - S55. Mass spectrum of compound (QS-10)

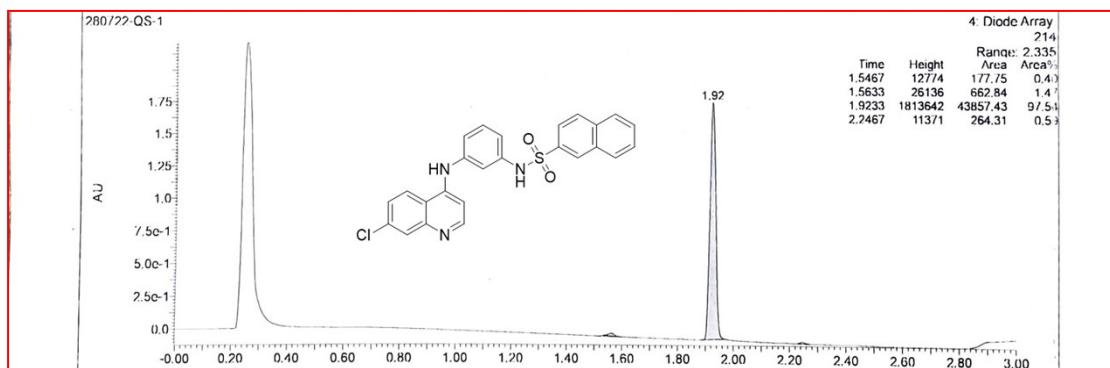


Figure - S56.Purity of compound (QS-10)

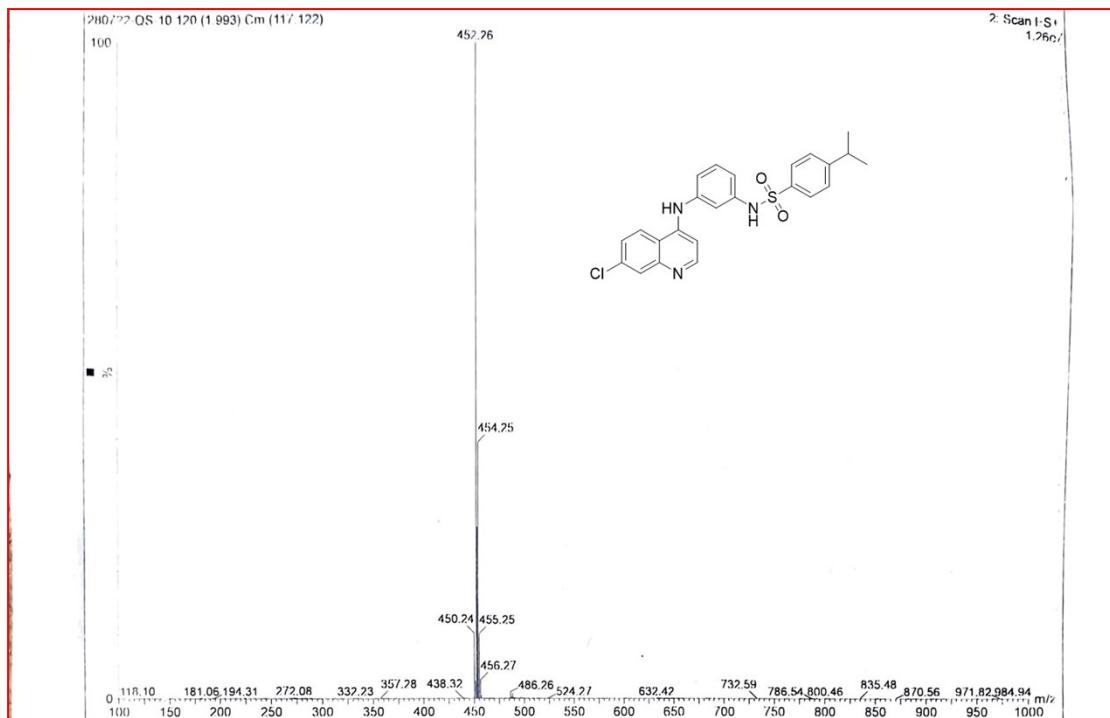


Figure - S57. Mass spectrum of compound (QS-11)

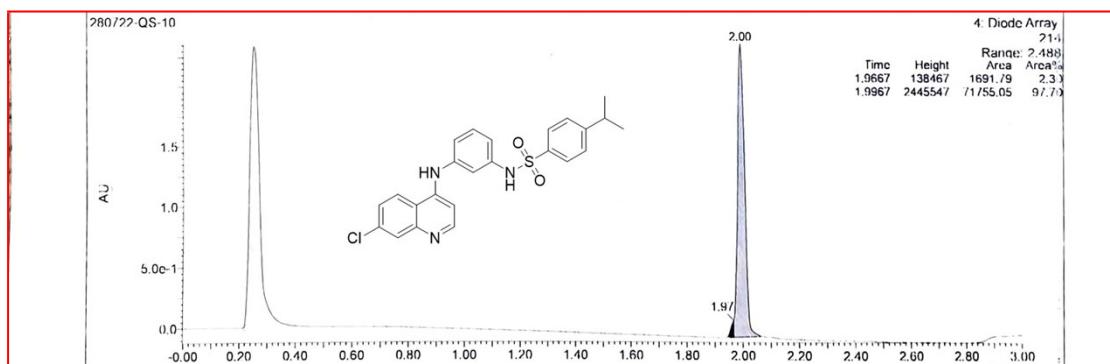


Figure - S58. Purity of compound (QS-11)

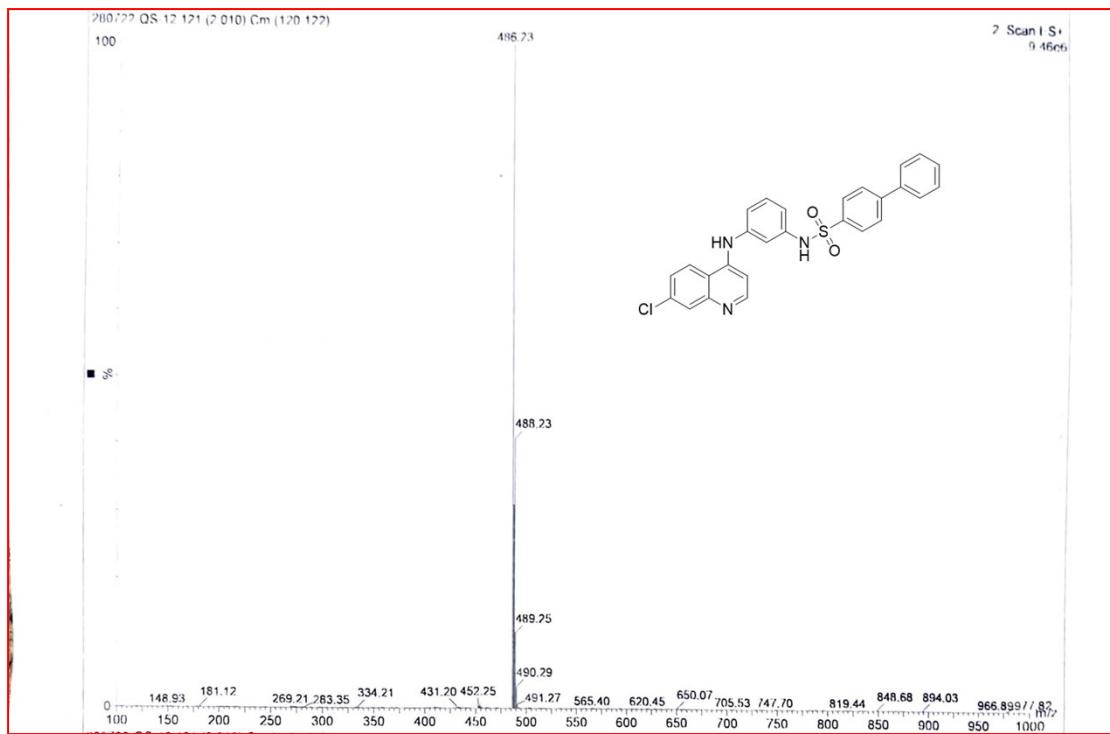


Figure - S59. Mass spectrum of compound (QS-12)

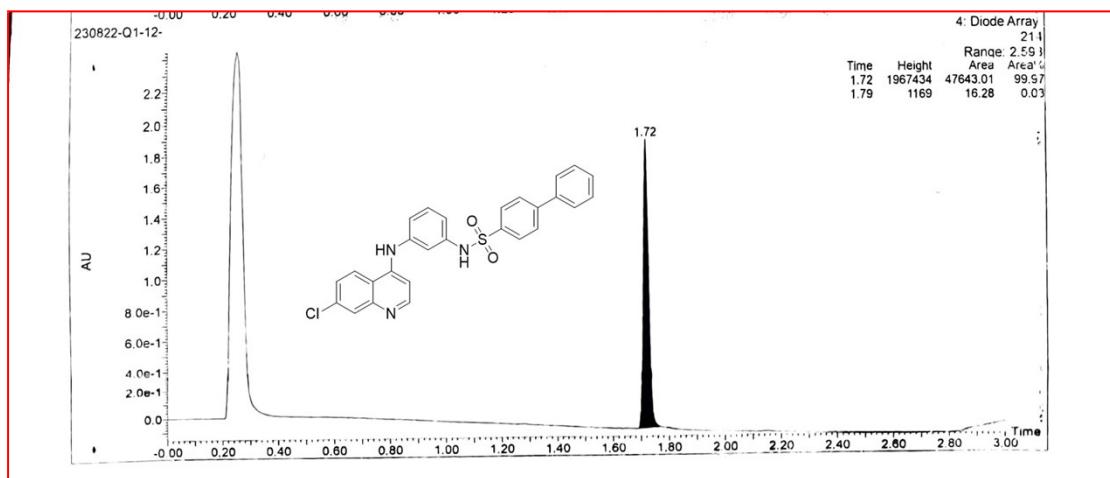


Figure - S60. Purity of compound (QS-12)

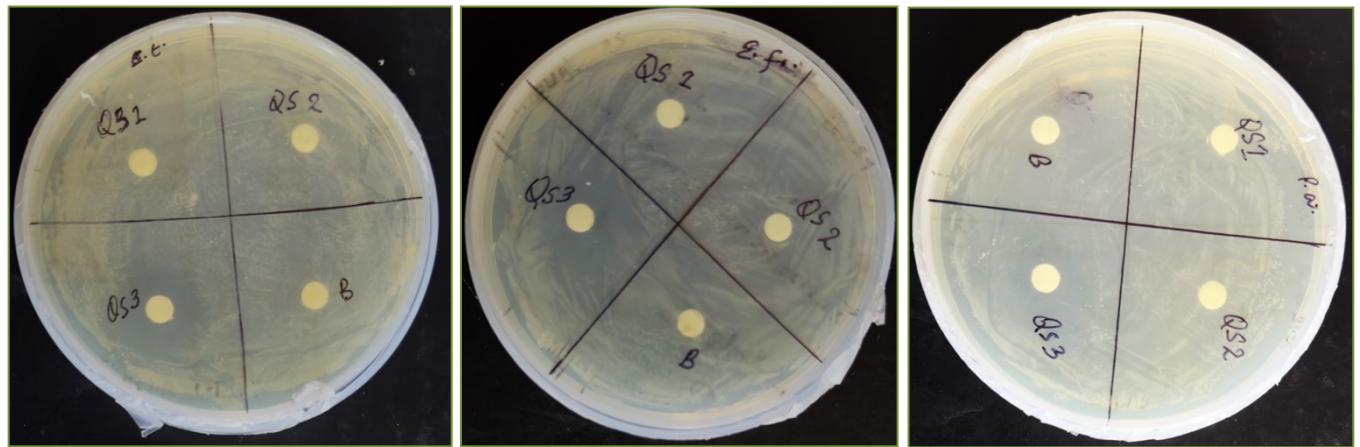


Figure – S61. Clearance zone in the bacterial lawn on treating with compounds

Combination Assay raw data:

Reading 1:

Bacterial Strain	MIC in combination ($\mu\text{g/mL}$)	
	Comp	CIP
<i>E. coli</i>	4	0.125
<i>E. faecalis</i>	2	0.125
<i>P. aeruginosa</i>	4	0.5
<i>S. typhi</i>	2	0.25

Reading 2:

Bacterial Strain	MIC in combination ($\mu\text{g/mL}$)	
	Comp	CIP
<i>E. coli</i>	2	0.25
<i>E. faecalis</i>	2	0.125
<i>P. aeruginosa</i>	8	0.25
<i>S. typhi</i>	2	0.25

Reading 3:

Bacterial Strain	MIC in combination ($\mu\text{g/mL}$)	
	Comp	CIP

<i>E. coli</i>	2	0.125
<i>E. faecalis</i>	8	0.5
<i>P. aeruginosa</i>	4	0.5
<i>S. typhi</i>	2	0.25

Disk diffusion Assay raw data:

Reading 1:

Isolates	Zone of Inhibition (in mm) at different concentrations of Test Compound.		
	½ MIC	MIC	2MIC
<i>E. coli</i>	6	7	7
<i>E. faecalis</i>	7	6	7
<i>P. aeruginosa</i>	6	8	12
<i>S. typhi</i>	7	5	8

Reading 2:

Isolates	Zone of Inhibition (in mm) at different concentrations of Test Compound.		
	½ MIC	MIC	2MIC
<i>E. coli</i>	6	9	8
<i>E. faecalis</i>	8	10	8
<i>P. aeruginosa</i>	5	6	8
<i>S. typhi</i>	7	9	5

Reading 3:

Isolates	Zone of Inhibition (in mm) at different concentrations of Test Compound.		
	½ MIC	MIC	2MIC
<i>E. coli</i>	6	6	7
<i>E. faecalis</i>	6	8	9
<i>P. aeruginosa</i>	7	10	10
<i>S. typhi</i>	7	7	11

Environmental strains raw data:

Isolates Name	Reading 1	Reading 2	Reading 3	Reading 4
EJH01	0	0	0	0
EJH02	0	0	0	0
EJH04	0	0	0	0
EJH05	0	0	0	0

EJH07	0	0	0	0
EJH11	0	0	0	0
EJH12	0	0	0	0
EJH13	0	0	0	0
EJH14	0	0	0	0
EJH15	0	0	0	0
EJH18	0	0	0	0
EJH19	0	0	0	0
EJH20	0	0	0	0
EJH21	0	0	0	0
EJH22	0	0	0	0
EJH23	0	0	0	0
EJH24	0	0	0	0
EJH25	0	0	0	0
A200	0	0	0	0
AA 201	0	0	0	0
AA 202	8	9	6	9
AA 209	0	0	0	0
AA 216	0	0	0	0
AA 221	0	0	0	0
AA 224	0	0	0	0
AA 237	0	0	0	0
AA 240	0	0	0	0
AA 243	0	0	0	0
AA 245	0	0	0	0
AA 248	0	0	0	0
AA 261	0	0	0	0
AA 269	0	0	0	0
AA 273	0	0	0	0
AA 276	0	0	0	0
AA 290	8	10	7	7

Hemolysis raw data:

Control	0.05	0.1	0.25	0.5	1
0.6	1.23	2.62	2.8	2.9	3.24
0.88	1.43	1.9	2.5	3.1	3.9
0.9	1.33	2.26	2.6	3.2	4.2

MIC raw data:

QS-3	<i>E. coli</i>	<i>E. faecalis</i>	<i>P. aeruginosa</i>	<i>K. pneumoniae</i>	<i>S. typhi</i>
Reading 1	128	128	64	256	512
Reading 2	128	64	64	128	512
Reading 3	128	128	32	256	512