

Synthesis, structural analysis, antimicrobial evaluation and synergistic studies of imidazo[1,2-a]pyrimidine chalcogenides

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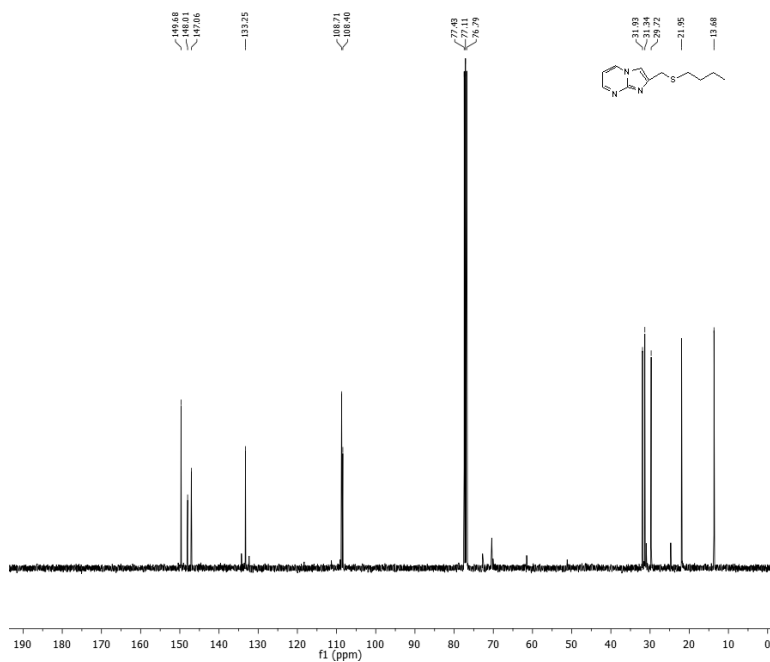
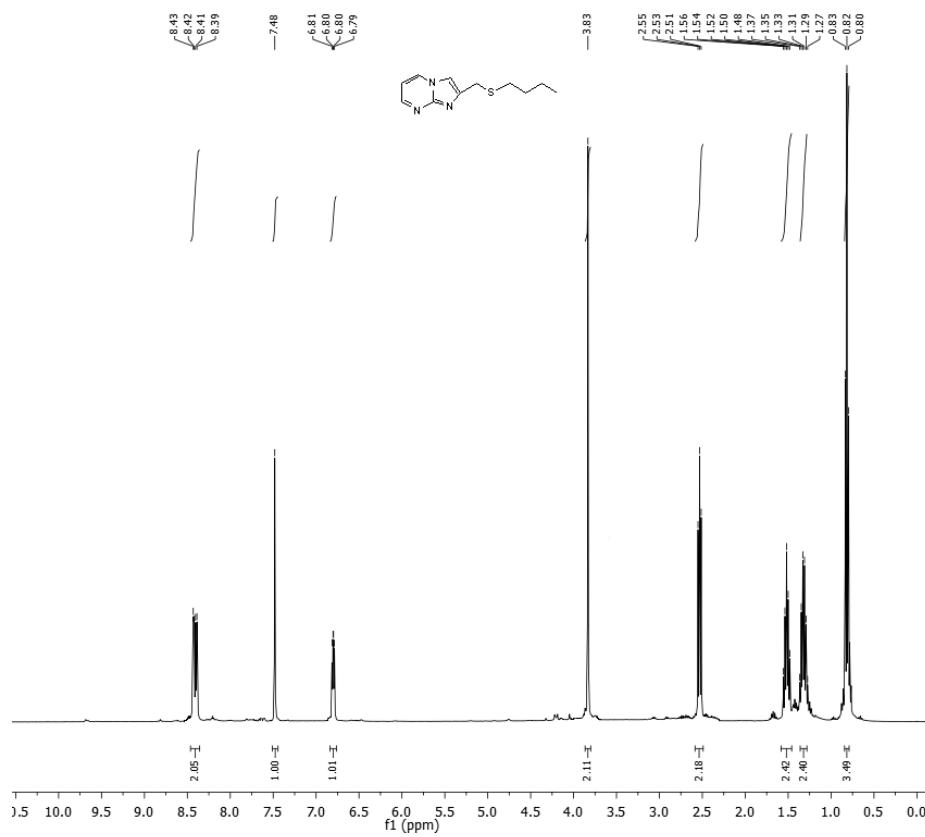
^cIndian Institute of Science Education and Research (IISER), Bhopal, India

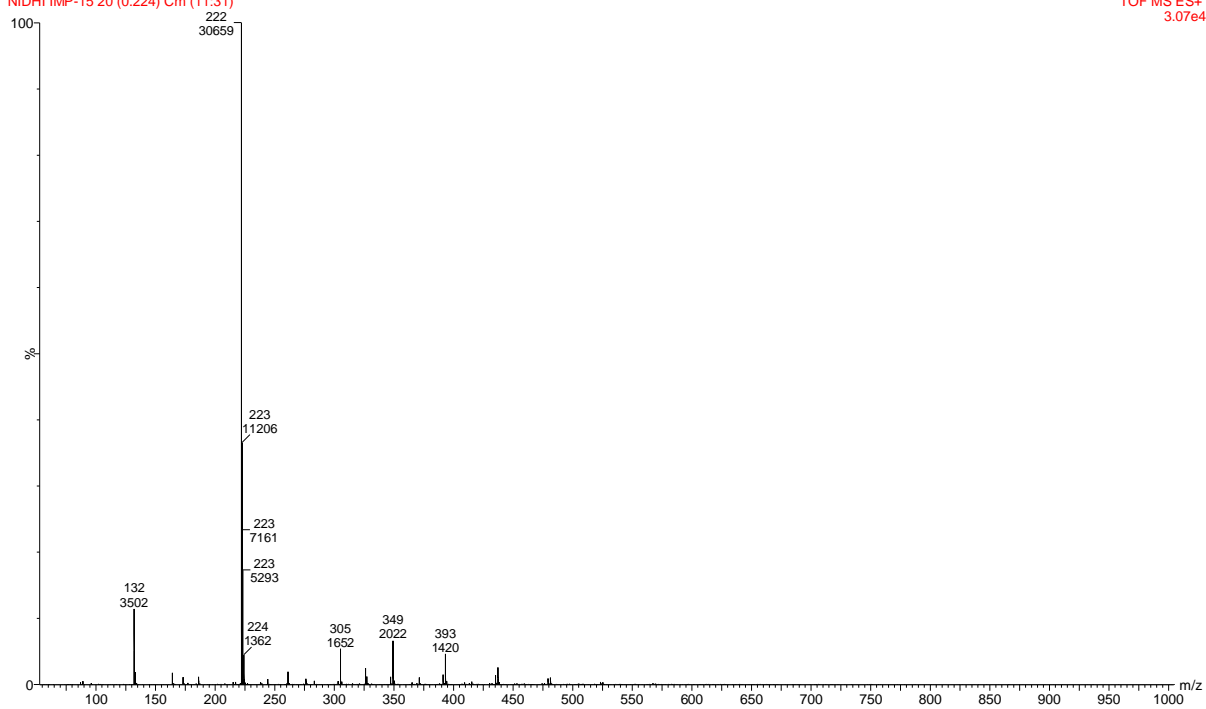
^dDepartment of Applied Sciences, University Institute of Engineering and Technology (UIET), Panjab University, Chandigarh, 160014, India

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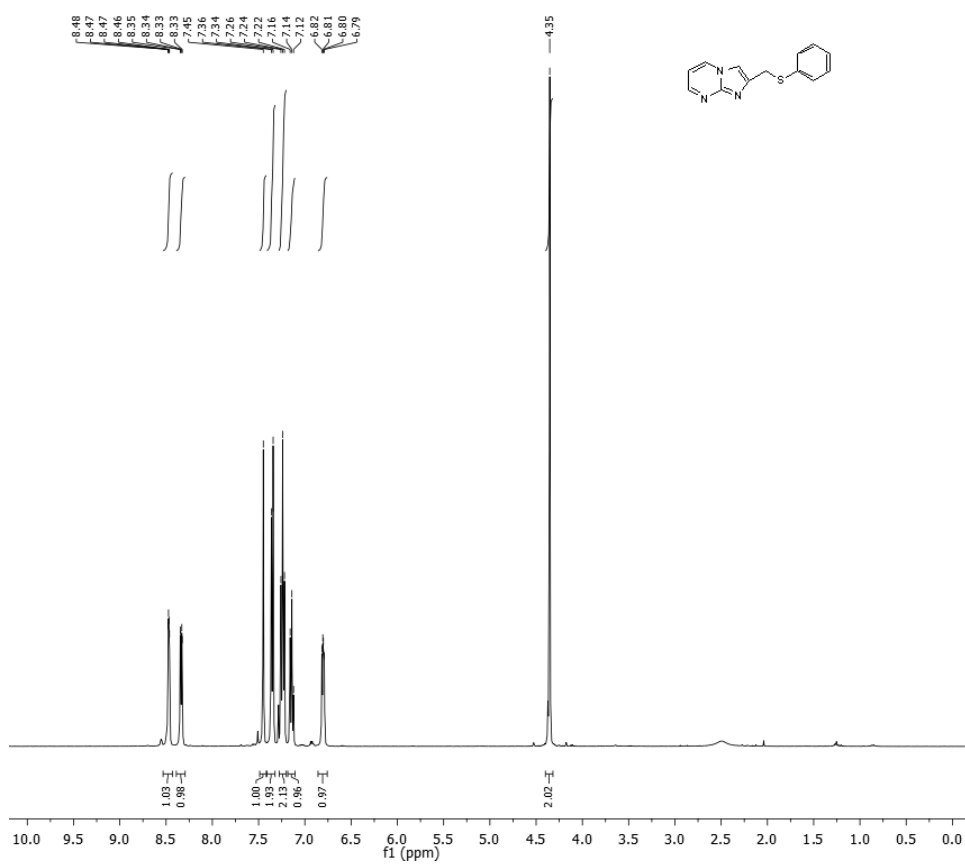
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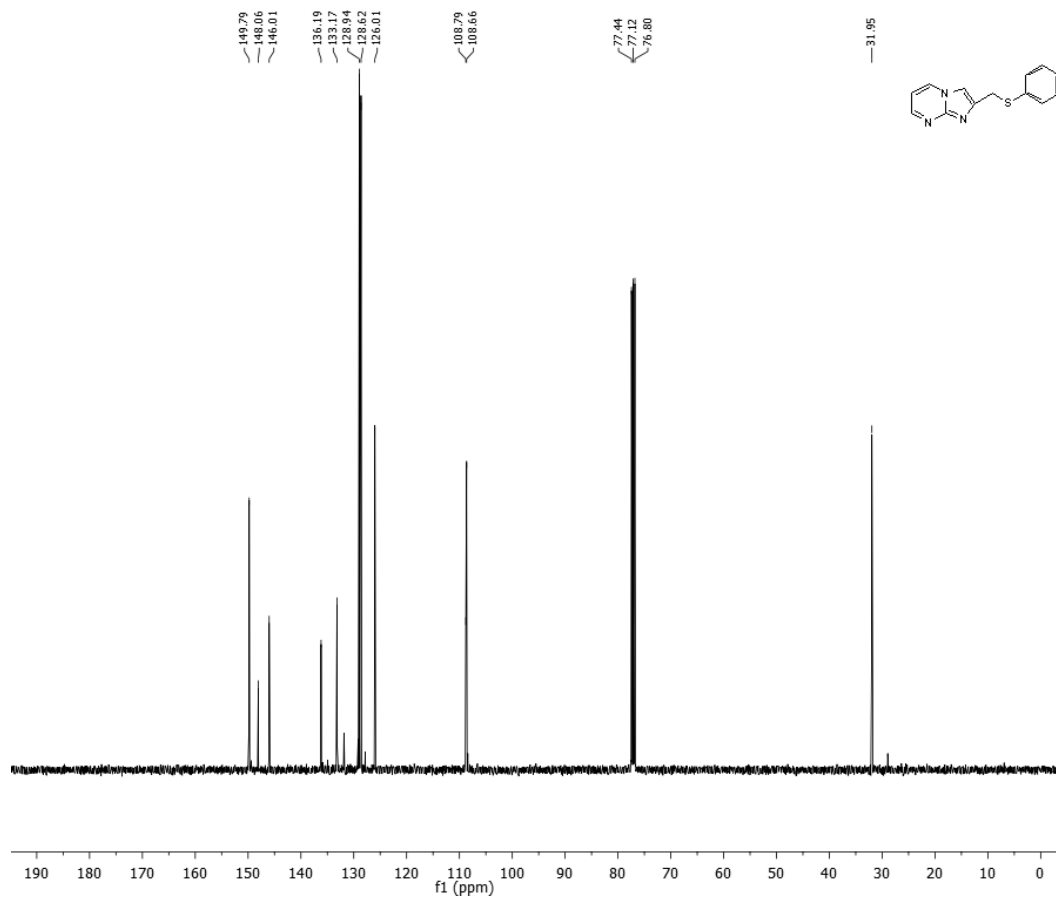
2-((Butylthio)methyl)imidazo[1,2-a]pyrimidine (4) ¹H, ¹³C, Mass Spectra





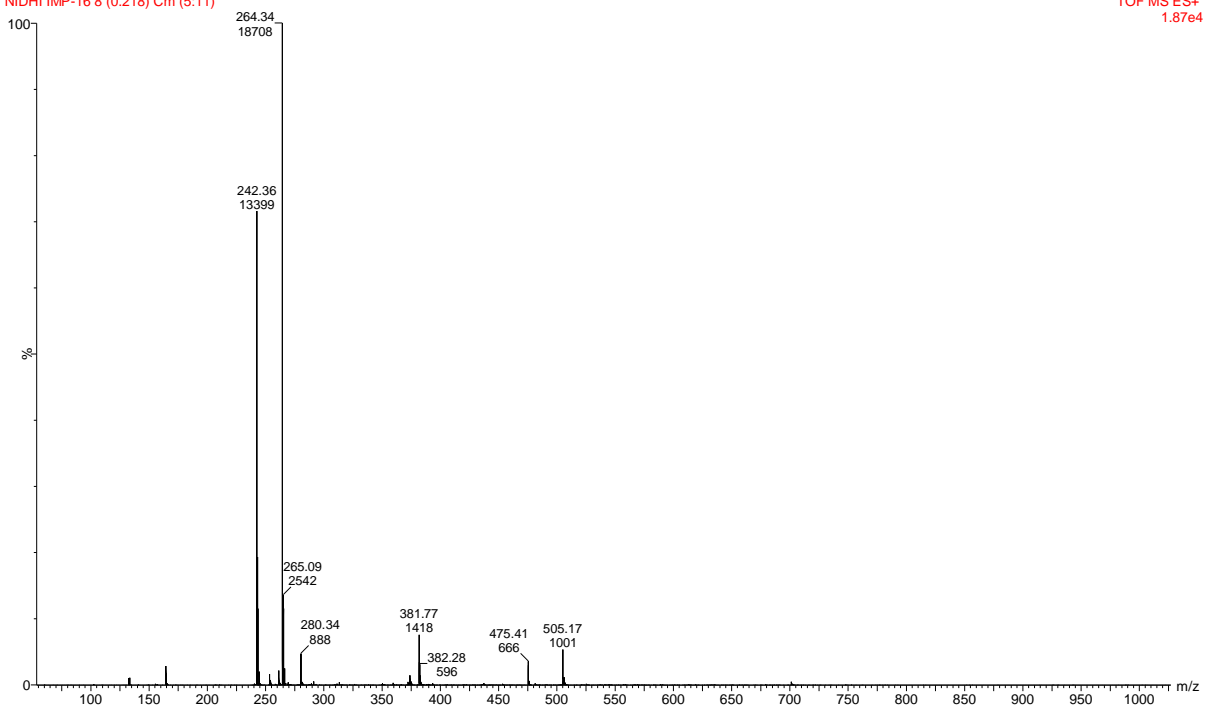
2-((Phenylthio)methyl)imidazo[1,2-a]pyrimidine (5) ¹H, ¹³C, Mass Spectra



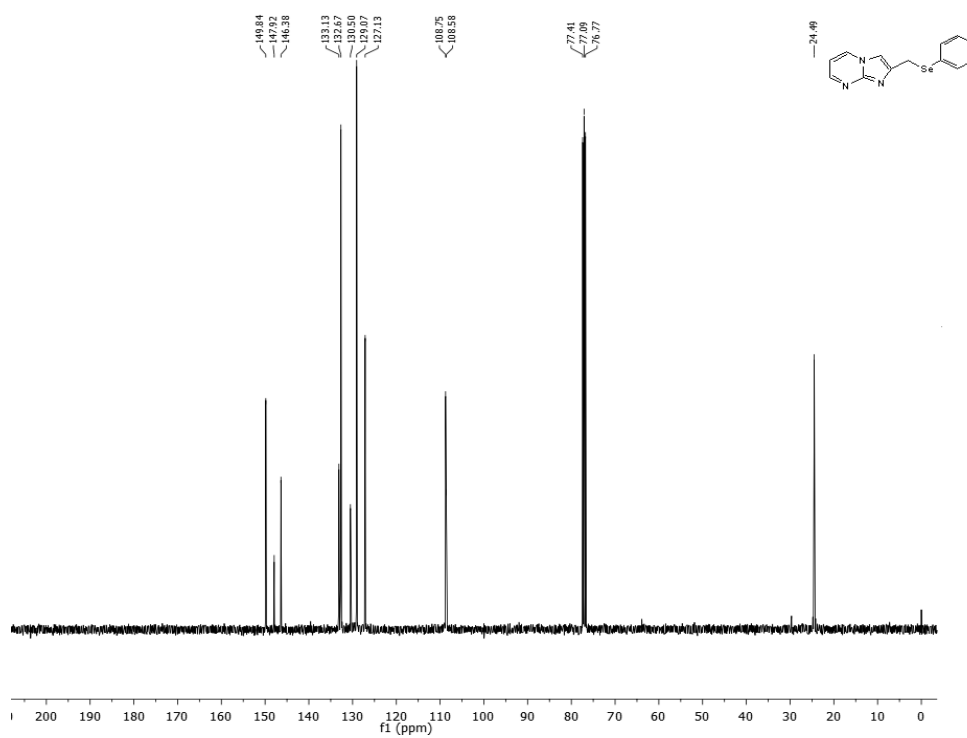
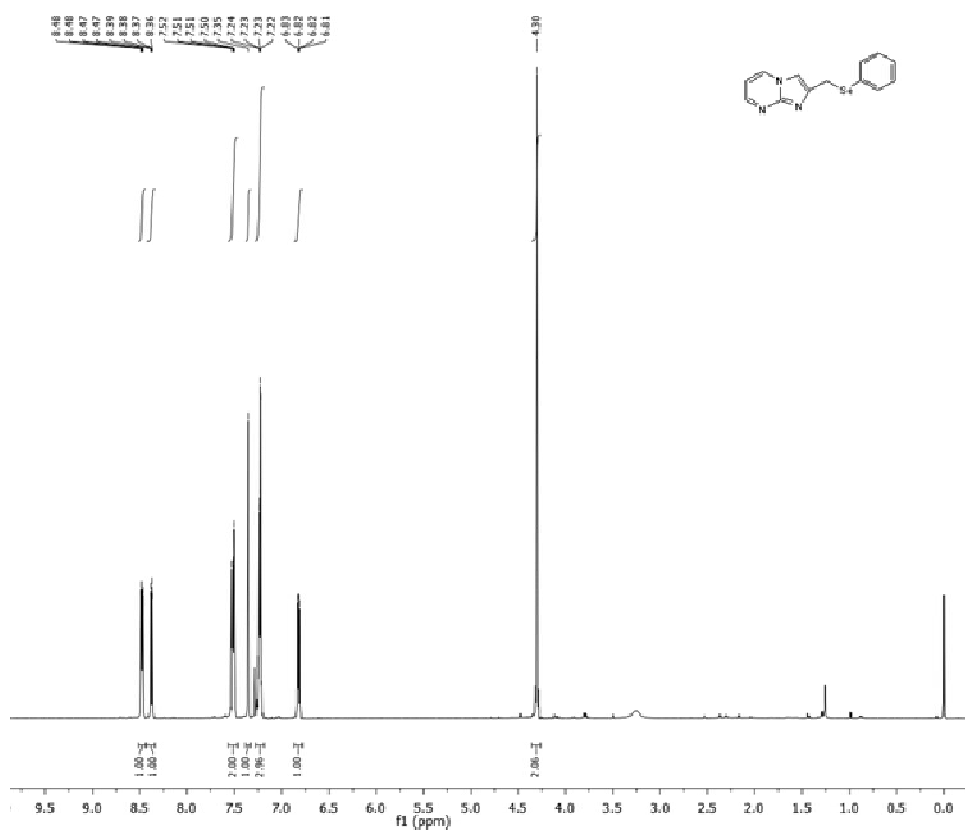


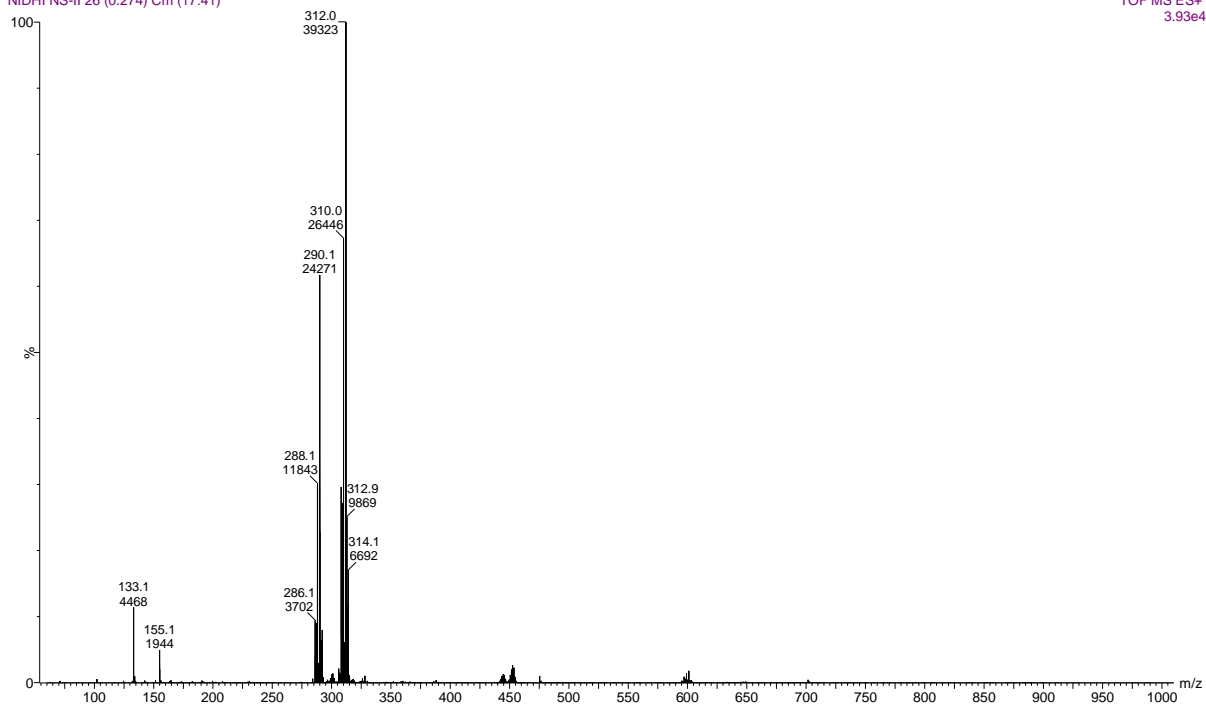
WATERS, Q-TOF MICROMASS (LC-MS)
 NIDHI IMP-16.8 (0.218) Cm (5:11)

SAIF/CIL, PANJAB UNIVERSITY, CHANDIGARH
 TOF MS ES+
 1.87e4

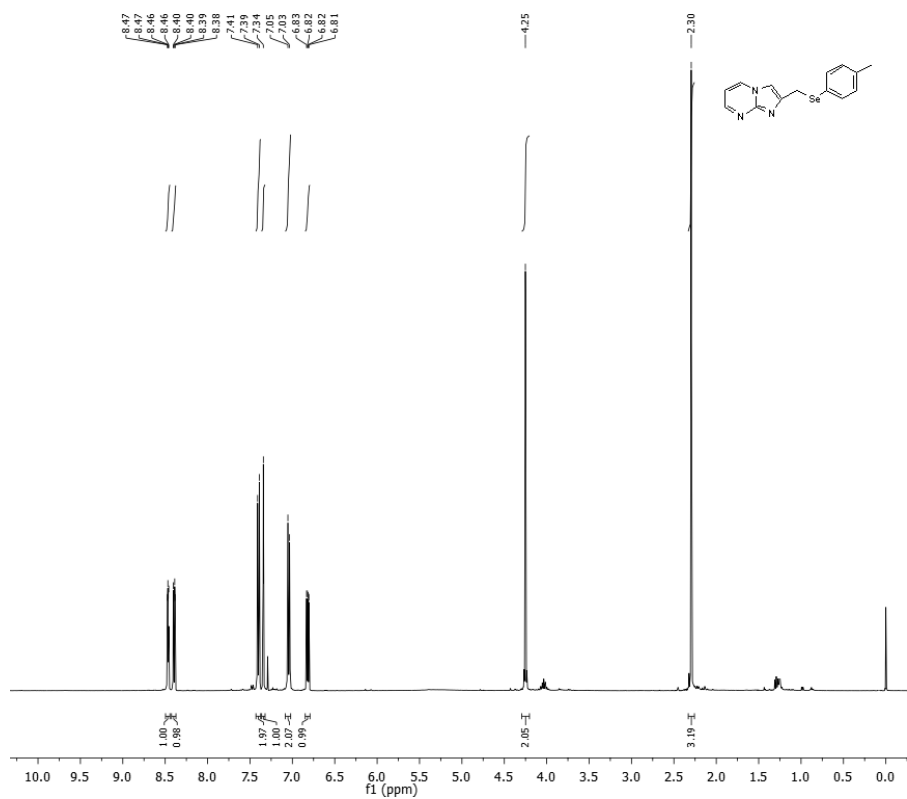


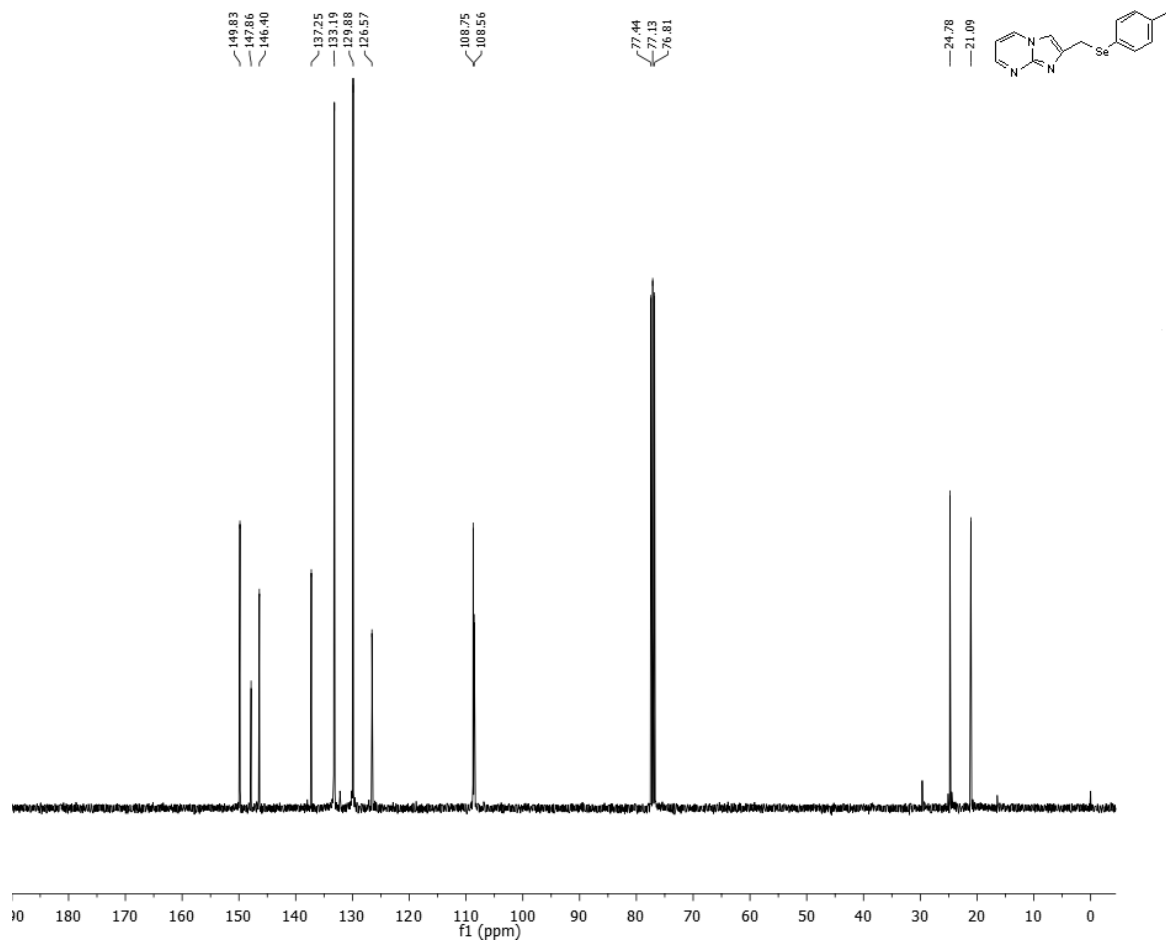
2-((Phenylselanyl)methyl)imidazo[1,2-a]pyrimidine (6) ¹H, ¹³C, Mass Spectra





2-((Tolylselanyl)methyl)imidazo[1,2-a]pyrimidine (7) ¹H, ¹³C, Mass Spectra



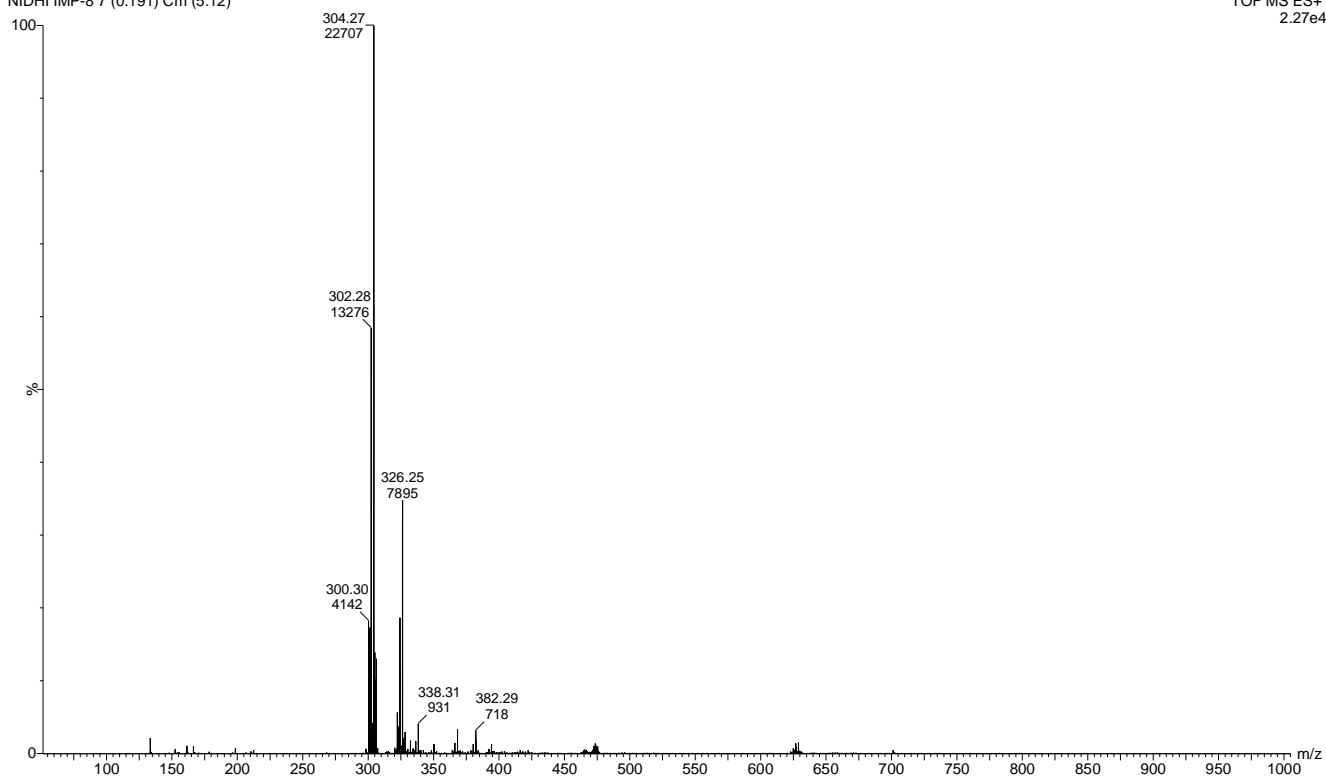


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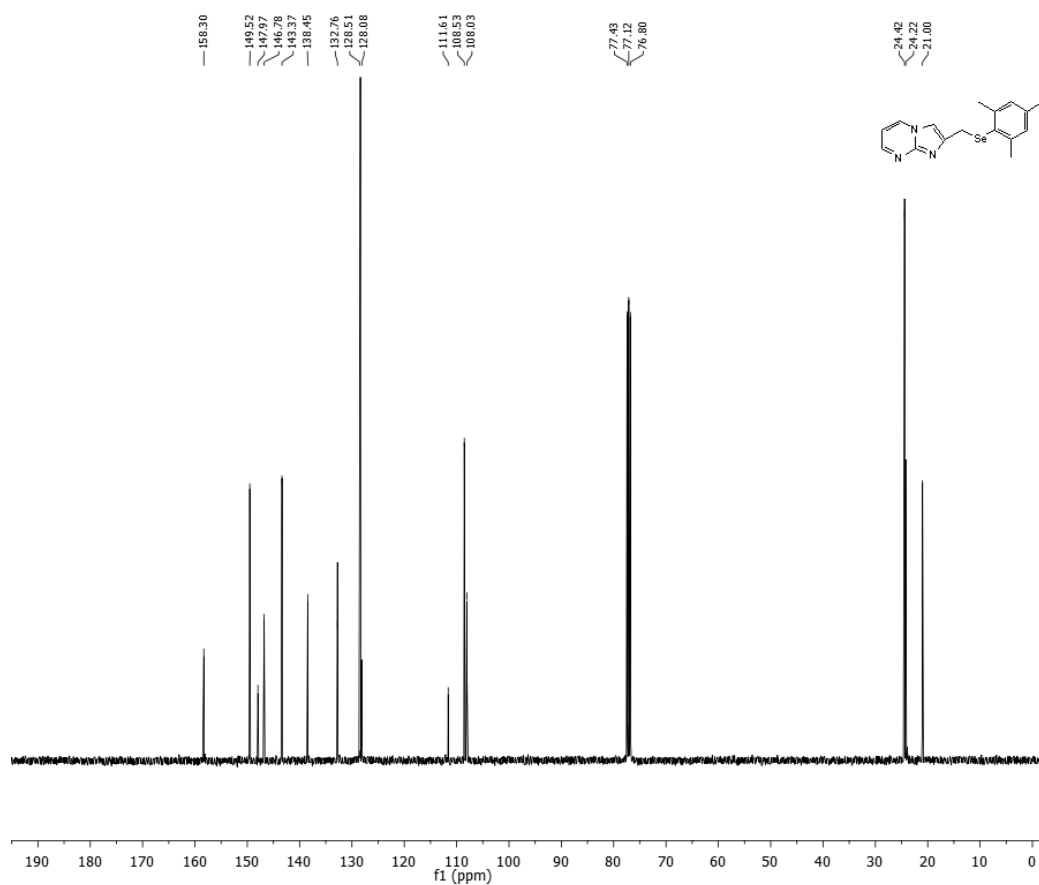
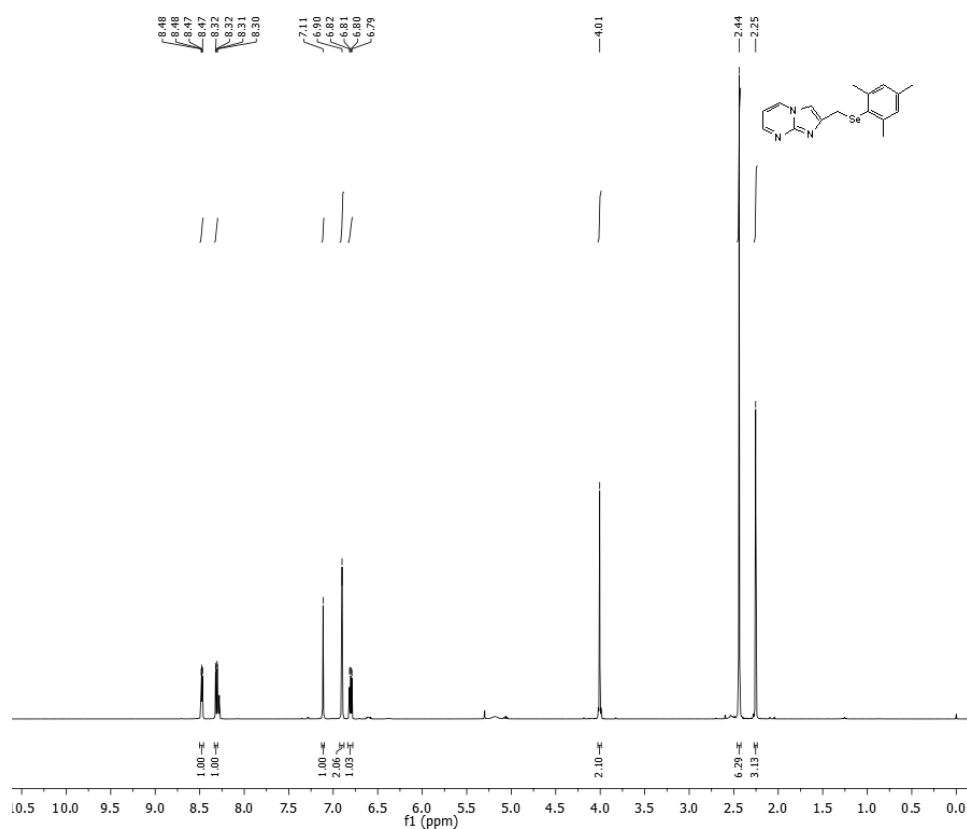
NIDHI IMP-8 7 (0.191) Cm (5:12)

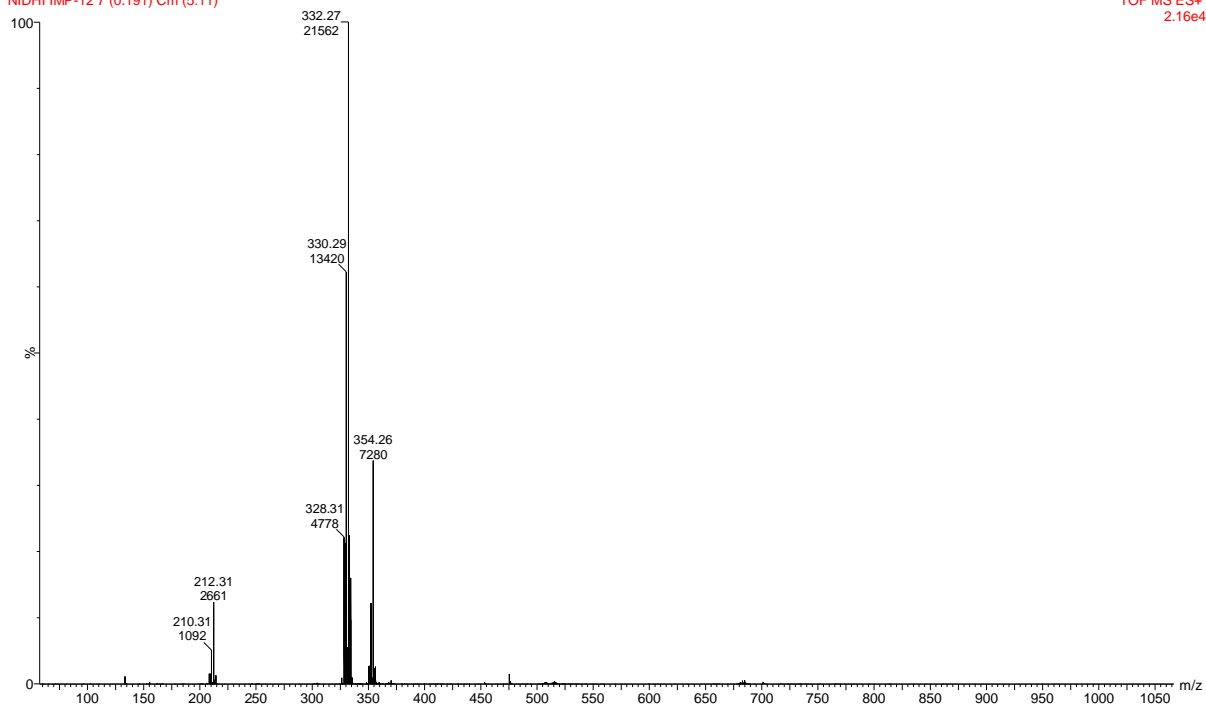
SAIF/CIL, PANJAB UNIVERSITY, CHANDIGARH

TOF MS ES+
2.27e4

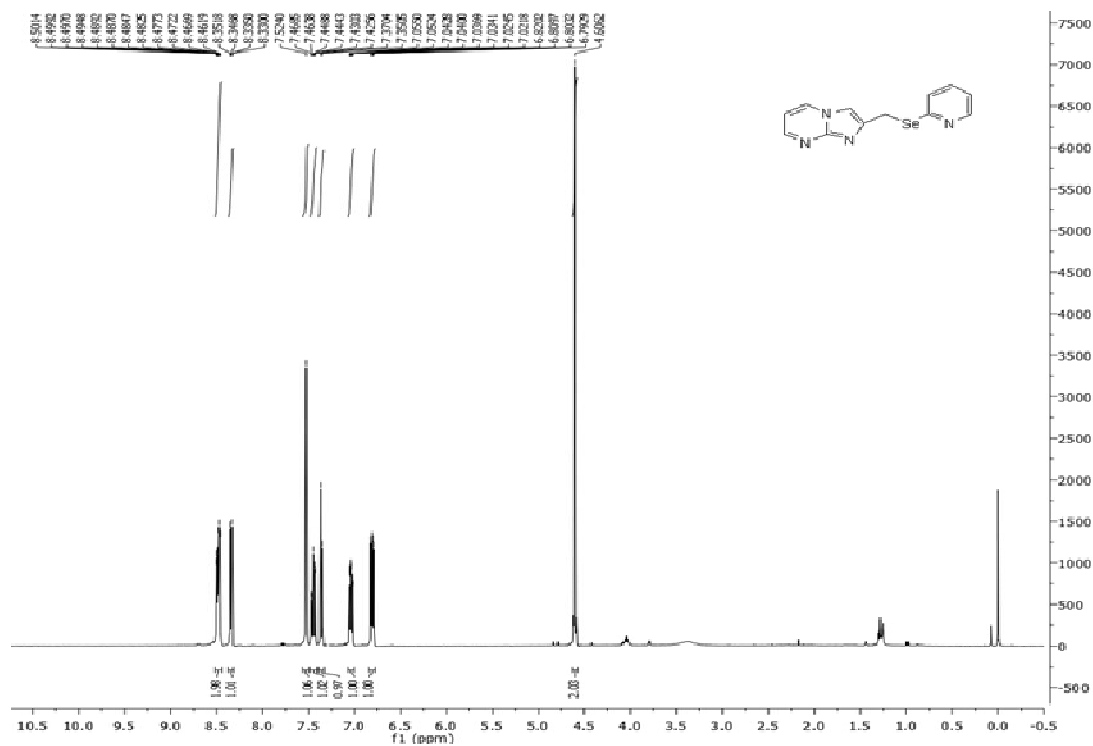


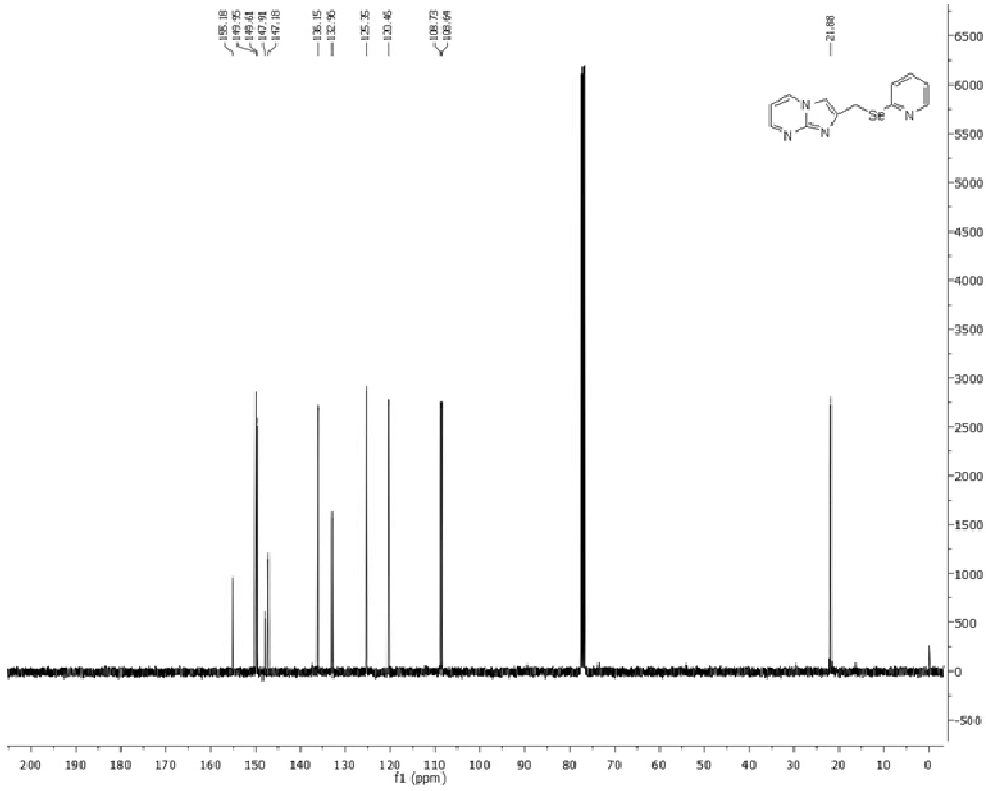
2-((Mesitylselenyl)methyl)imidazo[1,2-a]pyrimidine (8) ^1H , ^{13}C , Mass Spectra





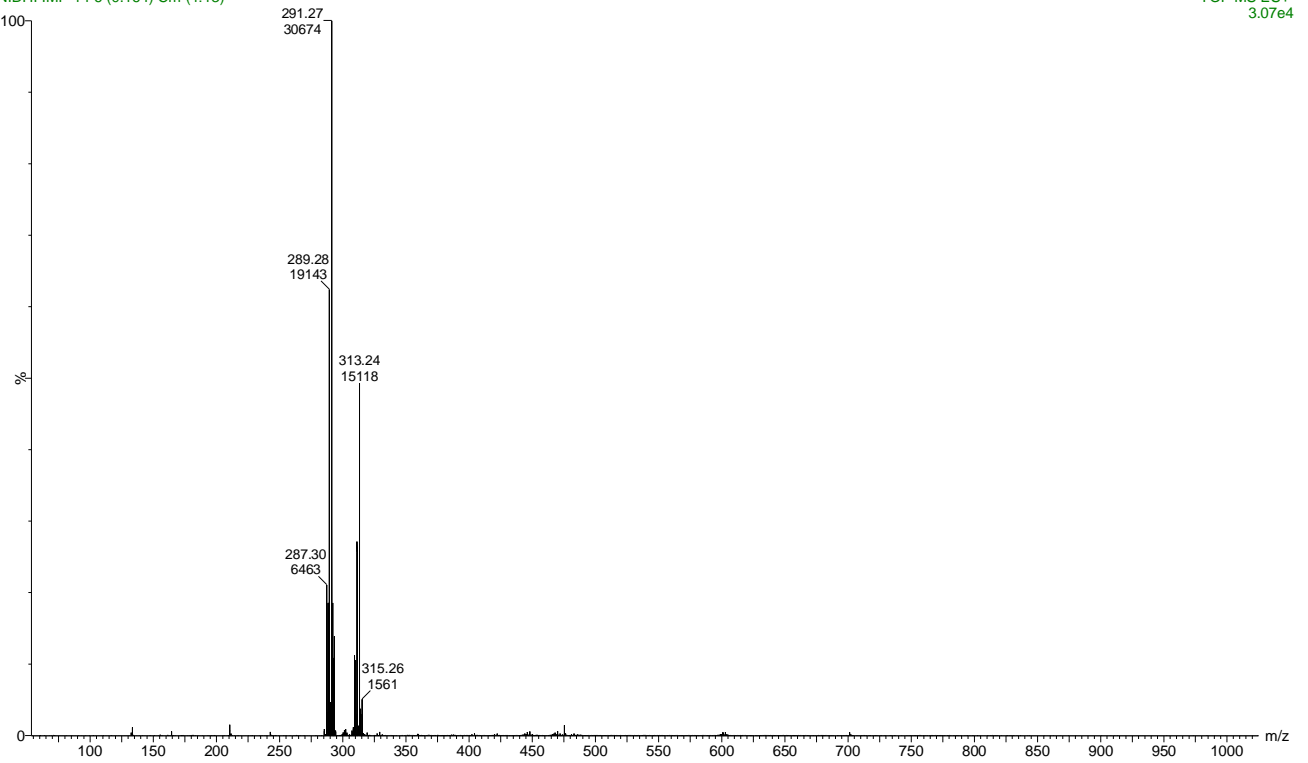
2-((Pyridylselanyl)methyl)imidazo[1,2-a]pyrimidine (9) ^1H , ^{13}C , Mass Spectra





WATERS, Q-TOF MICROMASS (LC-MS)

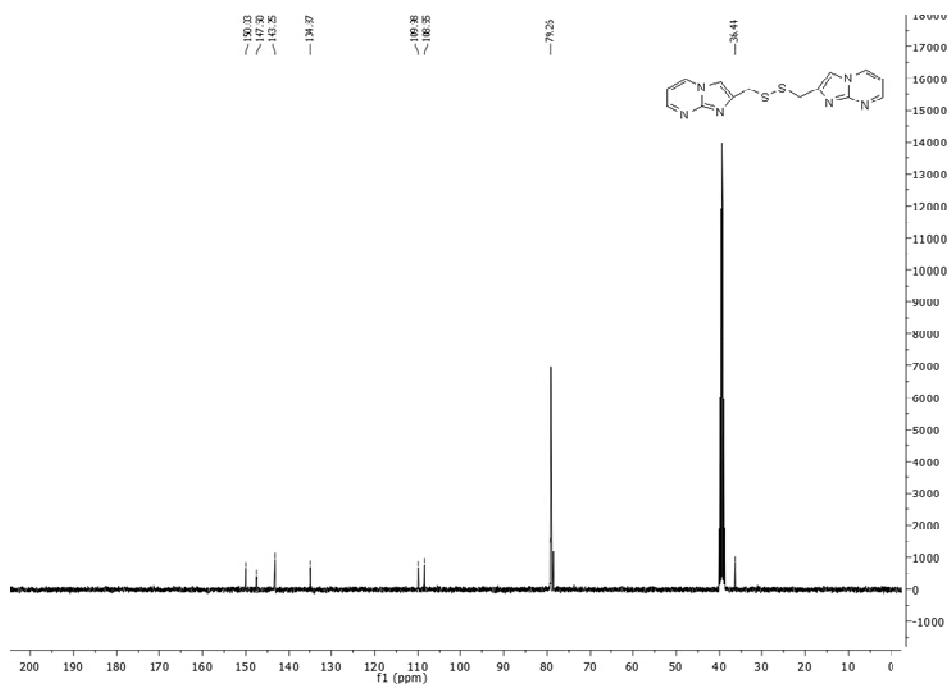
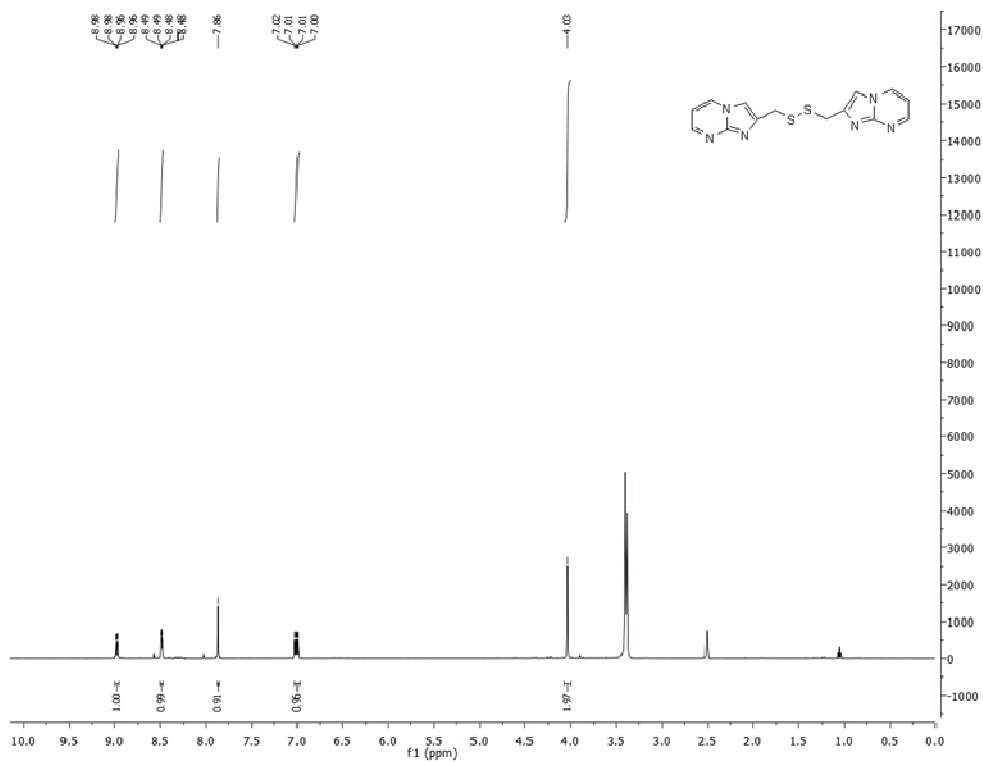
NIDHI IMP-14 6 (0.164) Cm (4:13)

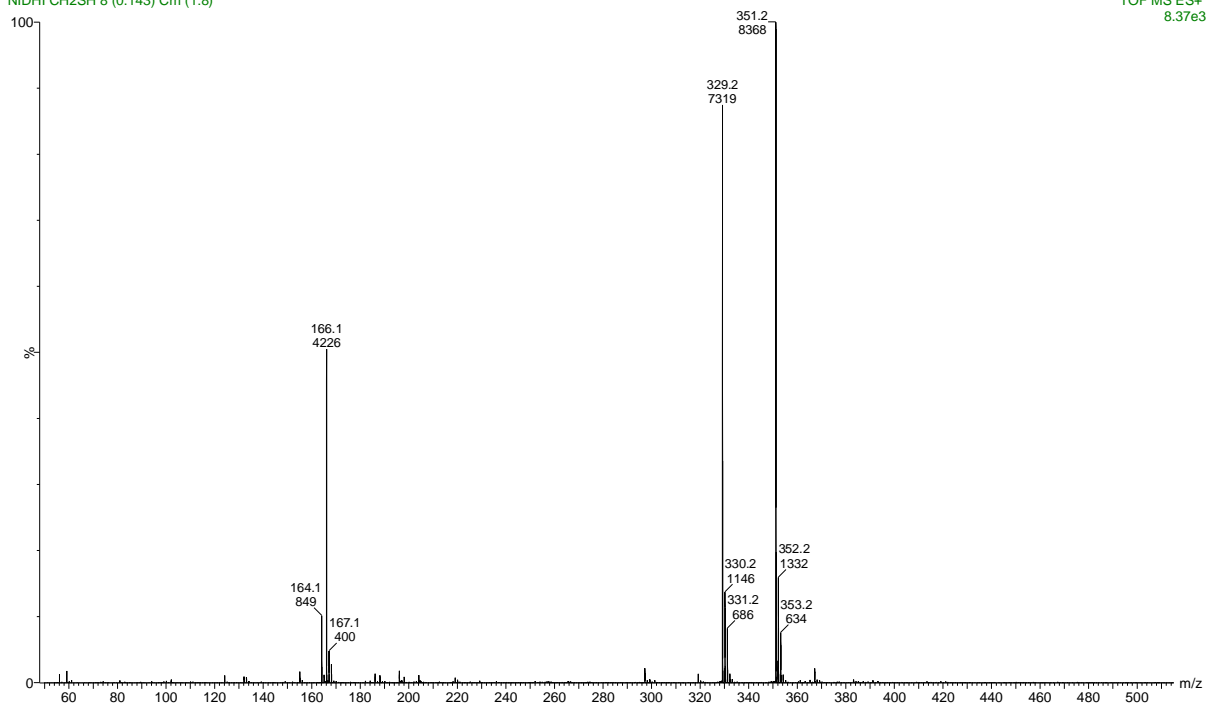


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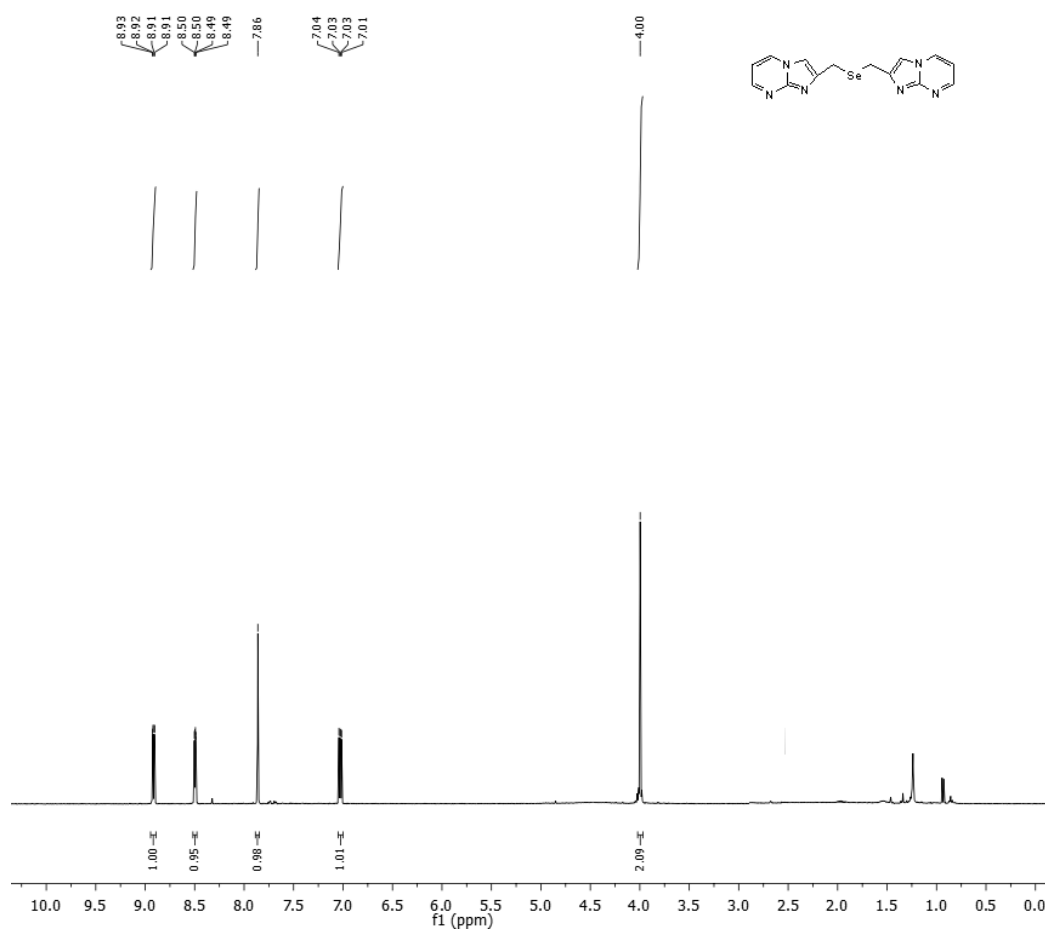
TOF MS ES+
3.07e4

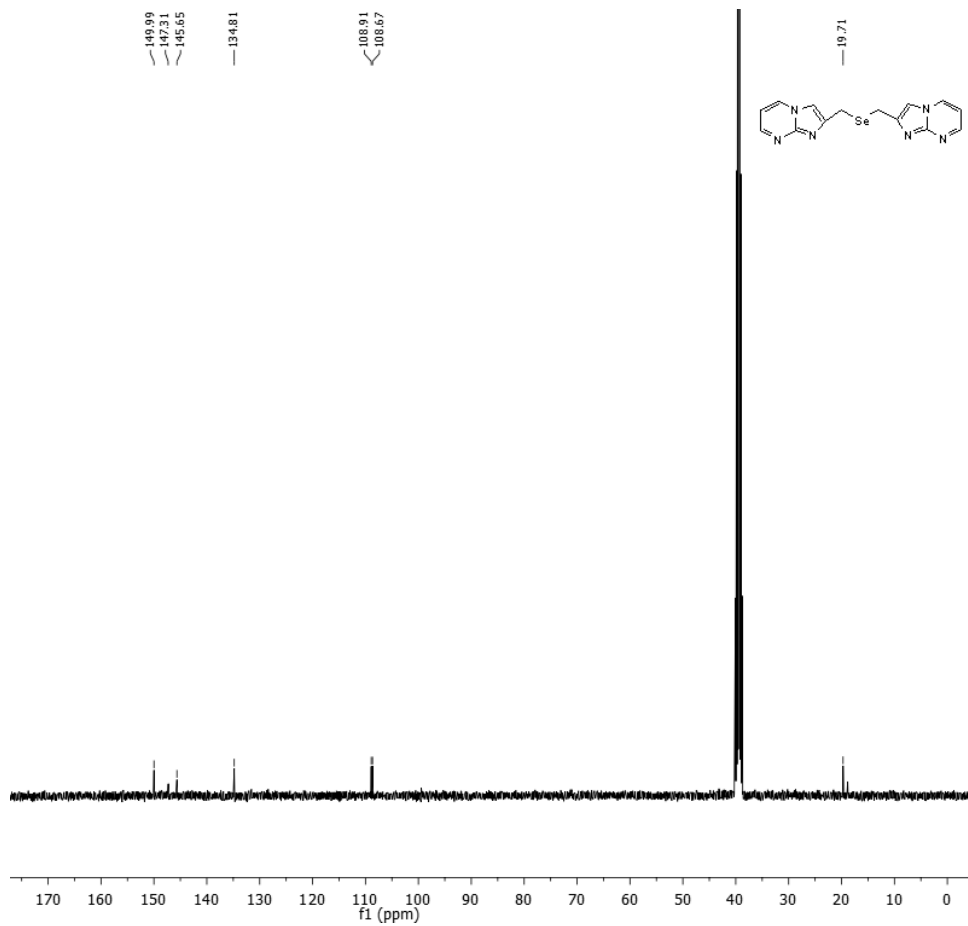
Bis(imidazo[1,2-a]pyrimidin-2-ylmethyl)disulfide (10) ¹H, ¹³C, Mass Spectra





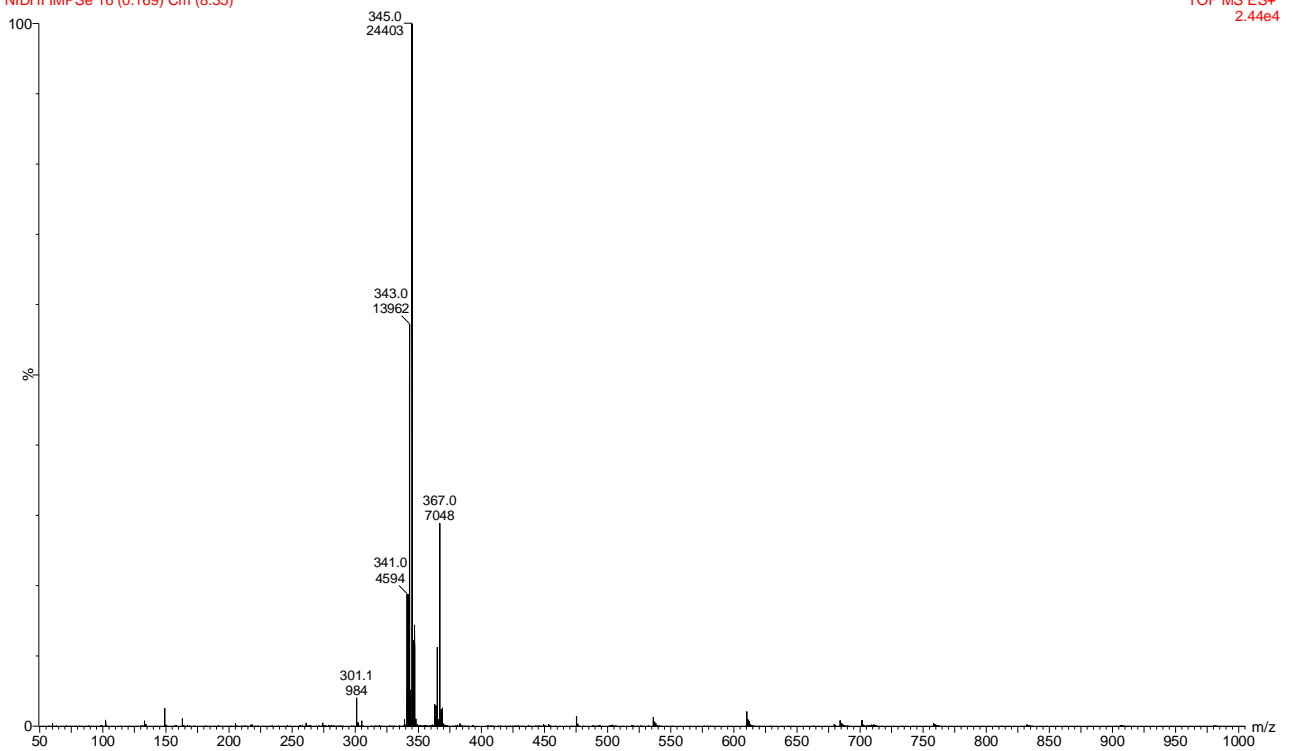
Bis(imidazo[1,2-a]pyrimidin-2-ylmethyl)selane (11) ¹H, ¹³C, Mass Spectra



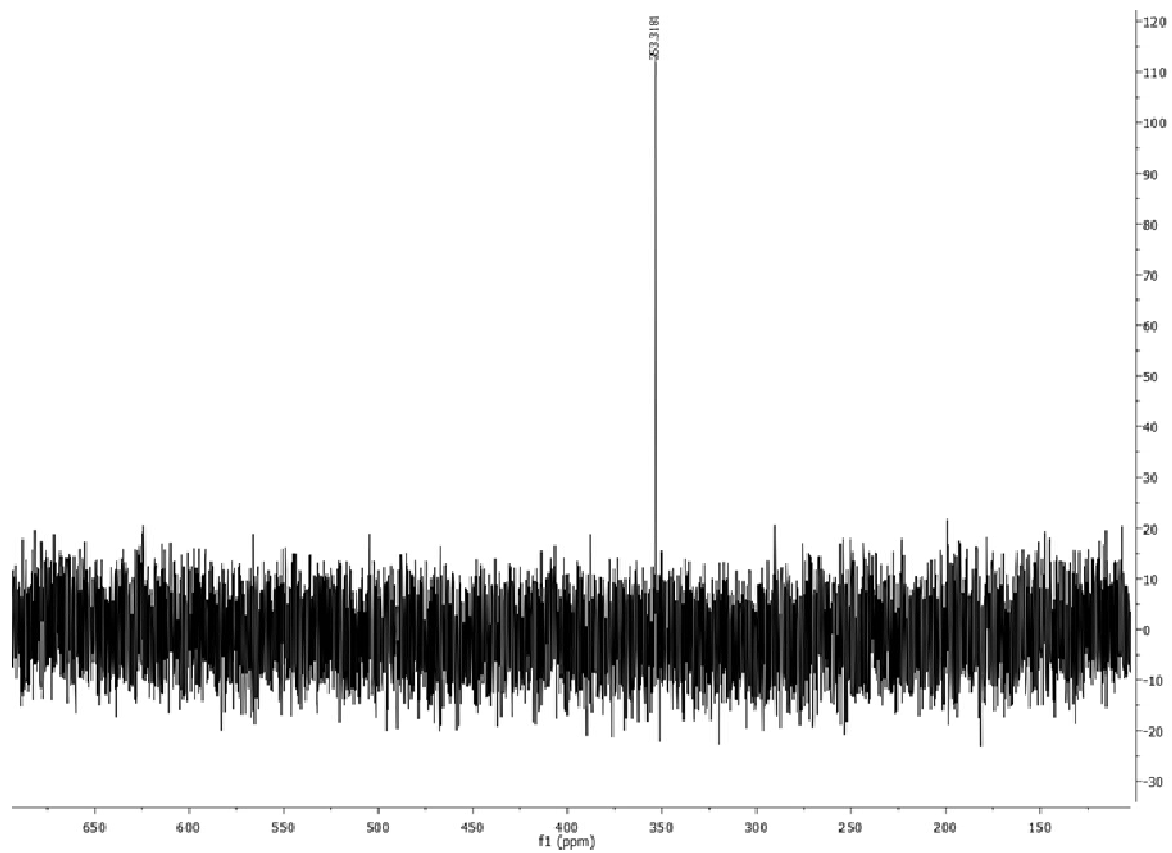


WATERS, Q-TOF MICROMASS (LC-MS)
 NIDHI IMPSe 16 (0.169) Cm (8:35)

SAIF/CIL,PANJAB UNIVERSITY,CHANDIGARH
 TOF MS ES+
 2.44e4



2-((Phenylselenanyl)methyl)imidazo[1,2-a]pyrimidine (6) ^{77}Se NMR



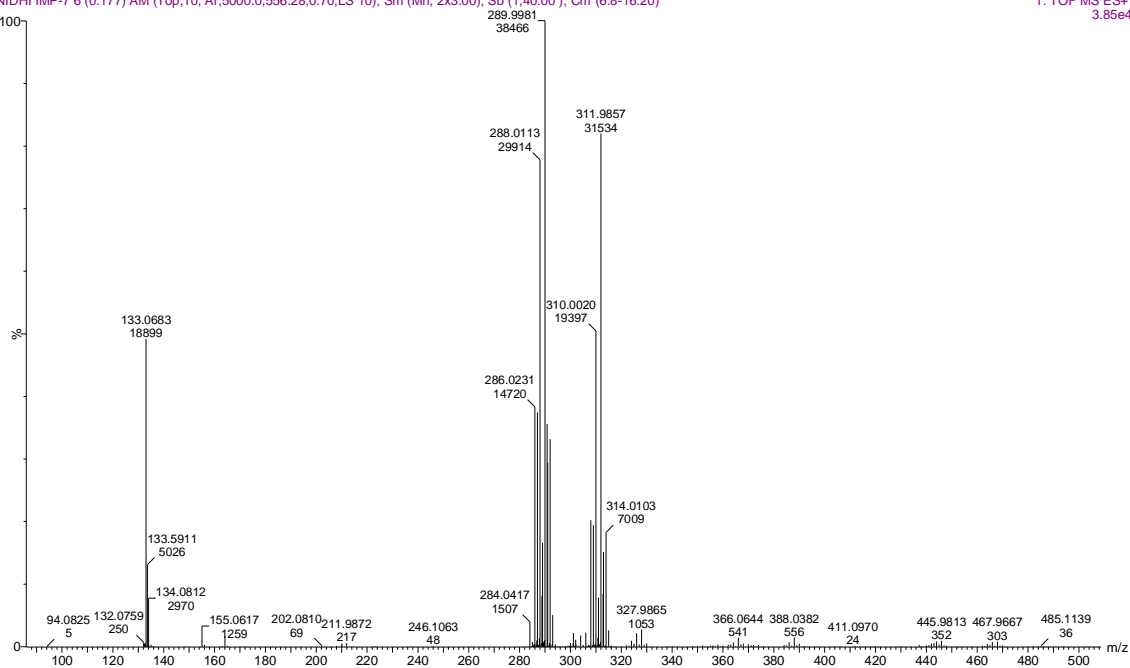
2-((Phenylselanyl)methyl)imidazo[1,2-a]pyrimidine (6) HRMS

WATERS, Q-TOF MICROMASS (ESI-MS)

NIDHI IMP-7 6 (0.177) AM (Top,10, Ar,5000.0,556.28,0.70,LS 10); Sm (Mn, 2x3.00); Sb (1,40.00); Cm (6:8-16:20)

SAIF/CIL,PANJAB UNIVERSITY,CHANDIGARH

1: TOF MS ES+
3.85e4



2-((Tolylselanyl)methyl)imidazo[1,2-a]pyrimidine (7)

WATERS, Q-TOF MICROMASS (ESI-MS)

NIDHI IMP-8 5 (0.153) AM (Top,10, Ar,5000.0,556.28,0.70,LS 10); Sm (Mn, 2x3.00); Sb (1,40.00); Cm (5:8-21:28)

SAIF/CIL,PANJAB UNIVERSITY,CHANDIGARH

1: TOF MS ES+
7.00e4

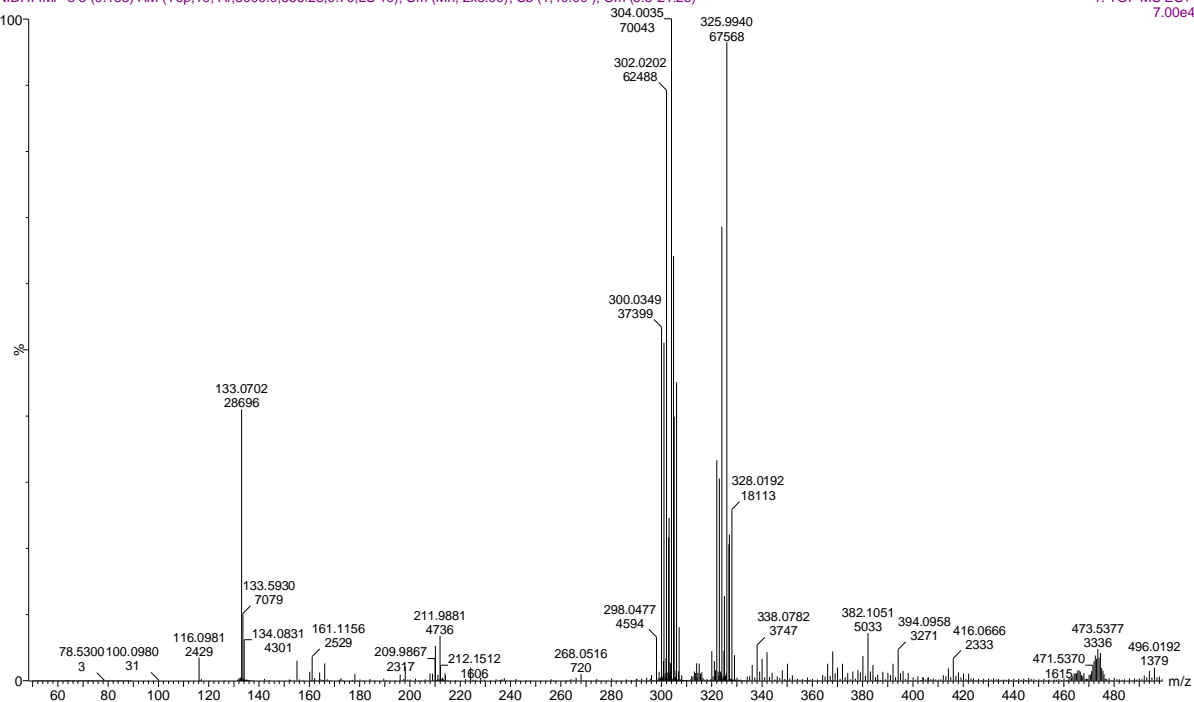


Table S1 Crystal data and structure refinement for **6**

Empirical formula	C13 H11 N3 Se
Formula weight	288.21
Temperature	296(2) K
Wavelength	0.71073 Å
Crystal system	Trigonal
Space group	R -3
Unit cell dimensions	a = 31.935(2) Å, α = 90° b = 31.935(2) Å, β = 90° c = 6.3894(5) Å, γ = 120°
Volume	5643.3(9) Å ³
Z	18
Density (calculated)	1.526 g/cm ³
Absorption coefficient	2.974 mm ⁻¹
F(000)	2592
Theta range for data collection	2.209 to 26.053°
Index ranges	-39<=h<=39, -39<=k<=39, -7<=l<=7
Reflections collected	27802
Independent reflections	2467 [R(int) = 0.0928]
Completeness to theta = 25.242°	99.5 %
Absorption correction	None
Refinement method	Full-matrix least-squares on F2
Data / restraints / parameters	2467 / 0 / 154
Goodness-of-fit on F2	1.102
Final R indices [I>2sigma(I)]	R1 = 0.0442, wR2 = 0.1088
R indices (all data)	R1 = 0.0760, wR2 = 0.1186
Extinction coefficient	n/a
Largest diff. peak and hole	0.592 and -0.281 e Å ⁻³

Table S2 Selected bond lengths (Å), bond angles (°) and torsional angles (°) for compound **6**

Se1-C8	1.917	C8-C9	1.378
Se1-C1	1.966	C8-C13	1.383
N3-C3	1.371	C6-C5	1.402
N3-C4	1.372	C2-C1-Se1	113.100
N3-C7	1.391	C8-Se1-C1	101.280
N1-C7	1.316	C2-C3-H3	127.100
N1-C2	1.376	N3-C3-H3	127.100
N2-C6	1.318	C1-Se1-C8-C13	37.700
N2-C7	1.346	C1-Se1-C8-C9	-144.600
C2-C3	1.356	Se1-C8-C9-C10	-178.500
C5-C4	1.338	C8-Se1-C1-C2	68.500
C13-C12	1.368	N1-C2-C1-Se1	74.300
C9-C10	1.363	C3-C2-C1-Se1	-104.600
C12-C11	1.376	Se1-C8-C13-C12	178.700
C11-C10	1.384	C3-N3-C7-N1	-0.200
C2-C1	1.477	C3-N3-C7-N2	-179.100

Compound	Energy (a.u.)	Band gap (eV)	Se1/S1-C8 Bond length (Å)	Se1-C1 Bond length (Å)	C8-Se1-C1 Bond angle (°)	C8-Se1-C1-C2 Dihedral angle (°)
4	-990.6457	4.1676	1.840	1.850	100.53	-73.31
5	-1064.4399	3.9608	1.793	1.858	103.76	-99.03
6	-3065.6345	3.9197	1.920	2.000	99.70	-101.40
7	-3104.9527	3.8906	1.925	2.000	99.74	-101.13
8	-3183.5905	3.9017	1.930	2.000	99.63	-109.90
9	-3081.6726	4.2019	1.920	1.972	101.30	-70.19
10	-3268.5440	3.5740	1.940	1.940	109.40	-60.00
11	-1665.5157	3.5541	1.780	1.780	-	-

Table S3 Calculations of theoretical parameters using density functional theory (DFT)

Table S4 Optimized structures of compounds 4-11 with DFT

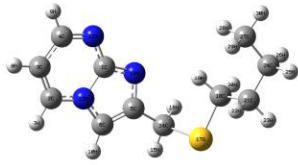

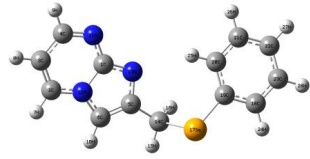
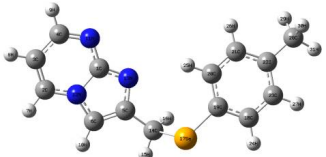
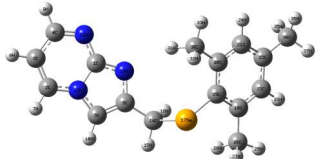
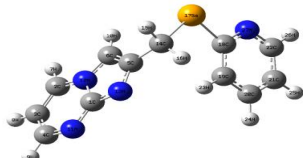
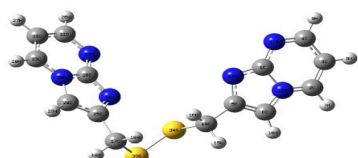
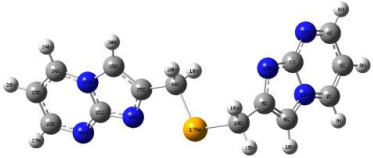
Compound	Optimized structure
4	
5	
6	
7	
8	
9	
10	
11	

Fig. S1 HOMO-LUMO diagrams of compounds 4-11

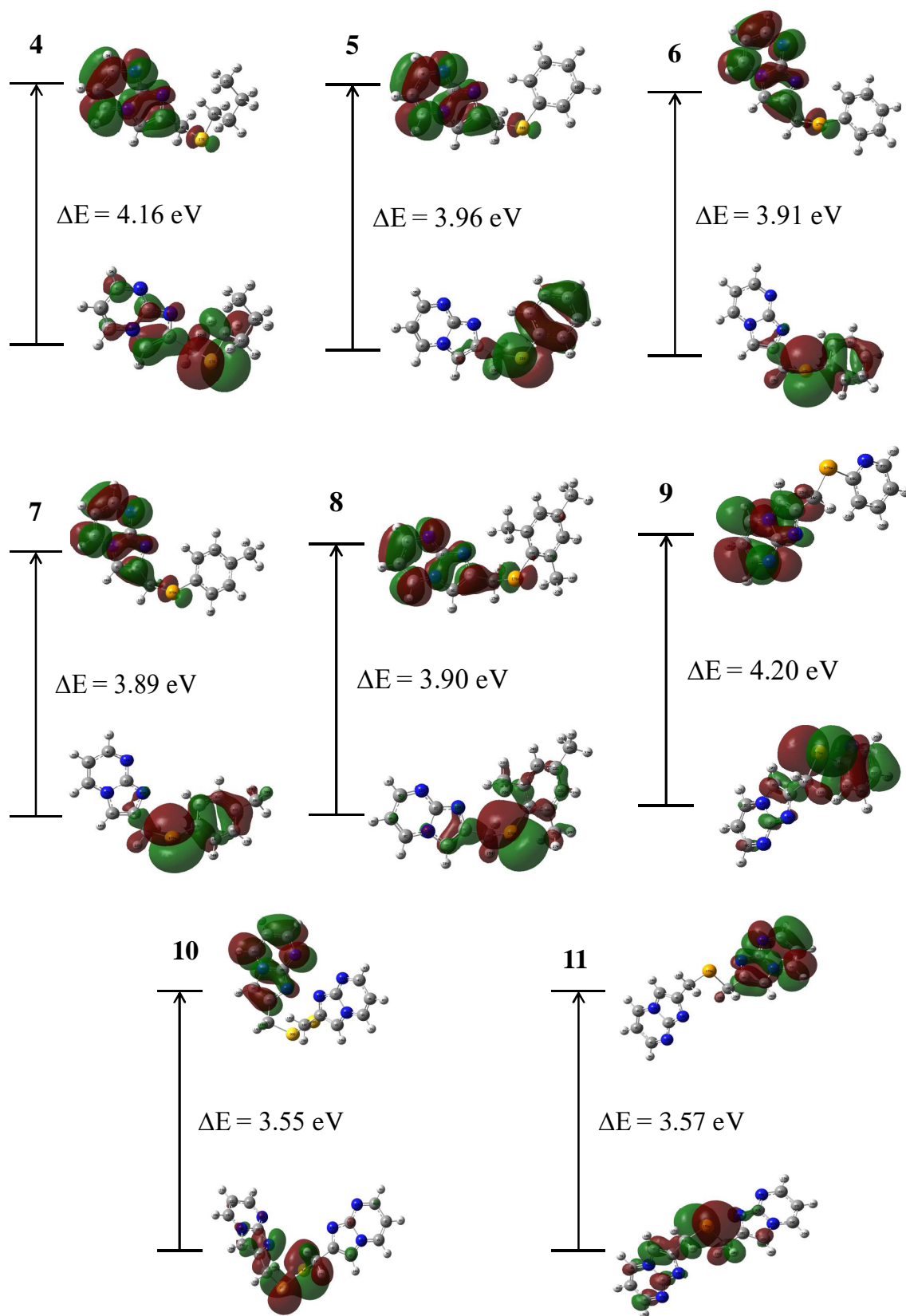


Table S5 MIC values of compounds **4-11** and standard antifungal agent (Amp B) against various fungal stains (in ug/ml)^a

Compound	<i>Candida albicans</i>	<i>Candida glabrata</i>	<i>Candida tropicalis</i>	<i>Candida krusei</i>	<i>Candida parapsolisis</i>	<i>Candida kyfer</i>	<i>Cryptococcus neoformans</i>	<i>Aspergillus niger</i>	<i>Neurospora crassa</i>
4	>27.62	>27.62	>27.62	>27.62	>27.62	>27.62	>27.62	NT	NT
5	>30.12	>30.12	>30.12	>30.12	>30.12	>30.12	>30.12	NT	NT
6	36.00	>36.00	>36.00	>36.00	>36.00	>36.00	>36.00	>36.00	18
7	>37.75	>37.75	>37.75	>37.75	>37.75	18.87	>37.75	37.75	18.87
8	>41.25	NT	NT	NT	NT	>41.25	>41.25	>41.25	20.62
9	>36.12	NT	NT	NT	NT	>36.12	>36.12	>36.12	18.06
10	>42.87	NT	NT	NT	NT	>42.87	>42.87	>42.87	>42.87
11	NT	NT	NT	NT	NT	NT	NT	NT	NT
Amp B	0.78	-	-	-	-	12.50	0.20	0.39	0.39

^aNT = Not Tested

Table S6 MIC values of compounds **4-11** and standard antibacterial agent (RIF) against various bacterial strains (in µg/ml)

Compounds	<i>Escherichia coli</i>	<i>Staphylococcus aureus</i>	<i>Vibrio cholera</i>	<i>Enterococcus faecalis</i>	<i>Listeria monocytogens</i>
4	>27.62	>27.62	>27.62	>27.62	>27.62
5	>30.12	>30.12	>30.12	>30.12	>30.12
6	>36.00	>36.00	>36.00	>36.00	>36.00
7	18.87	>37.75	>37.75	37.75	>37.75
8	>41.25	>41.25	>41.25	>41.25	>41.25
9	>36.12	>36.12	>36.12	>36.12	>36.12
10	>42.87	>42.87	>42.87	>42.87	42.87
11	NT	NT	NT	NT	NT
RIF	4.00	-	-	12.50	78.80