

Supporting Information

Design and Synthesis of Non-Covalent Imidazo[1,2-*a*]quinoxaline-Based Inhibitors of EGFR and Their Anti-Cancer Assessment

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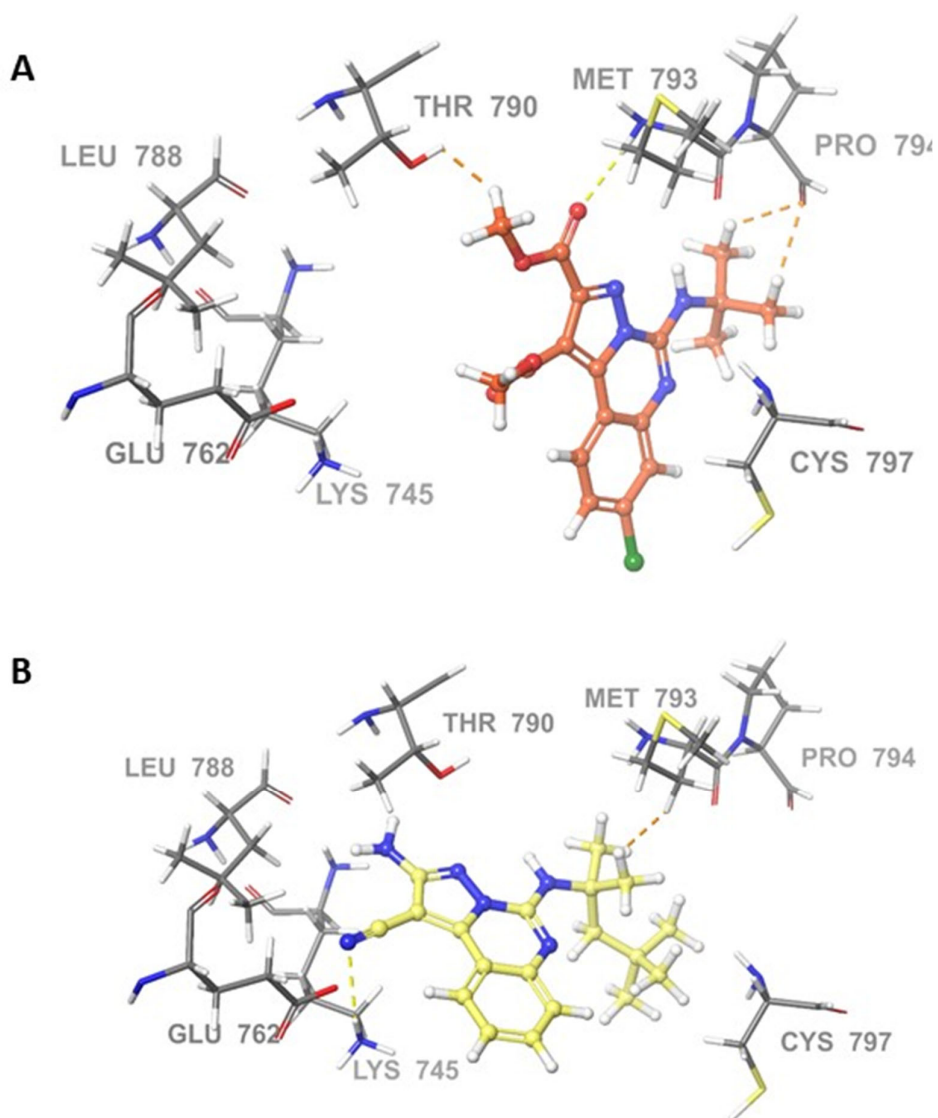


Figure S1. 3D docking pose of **A. I** and **B. II**, at EGFR ATP kinase domain.

Table S1. Docking scores of **I**, **II**, erlotinib and designed compound **6b**.

Entry	Compound	Dock score
1	6b	-5.463
2	I	-4.950
3	II	-4.067
4	Erlotinib	-4.985

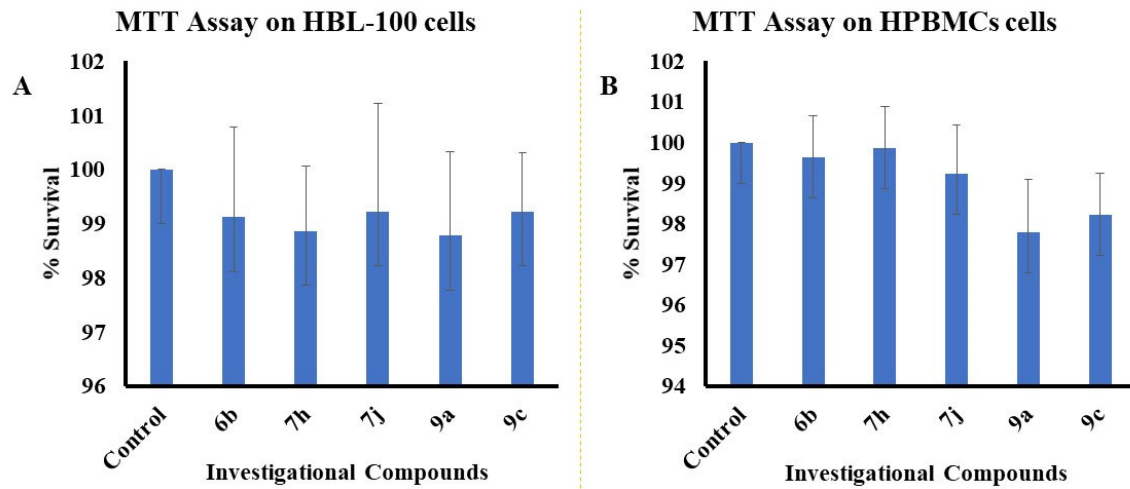


Figure S2. Bar graphs representing percentage survival of normal cells, **A.** HBL-100 (breast); **B.** HPBMCs (Human Peripheral Blood Mononuclear Cells), upon treatment with investigational compounds **6b**, **7h**, **7j**, **9a** and **9c** at 10 μ M concentration previously incubated for 24 h.

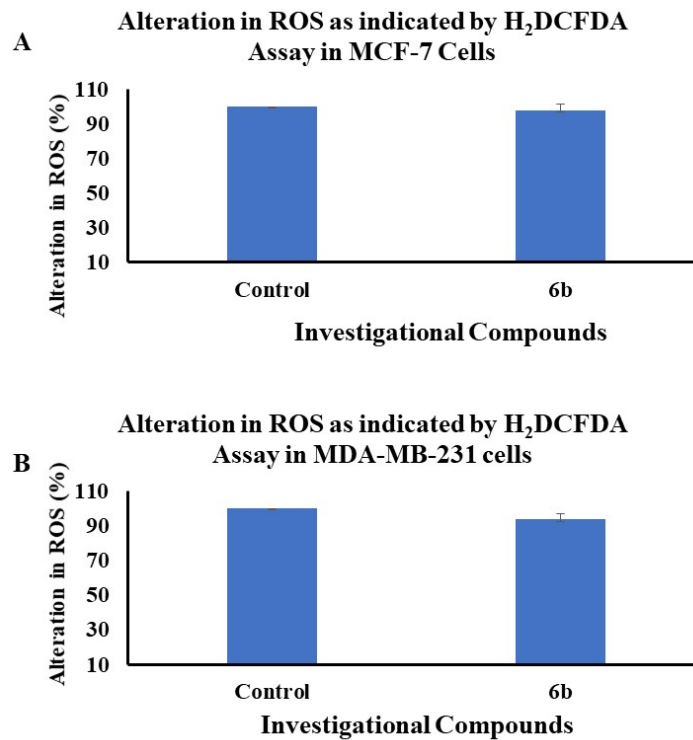
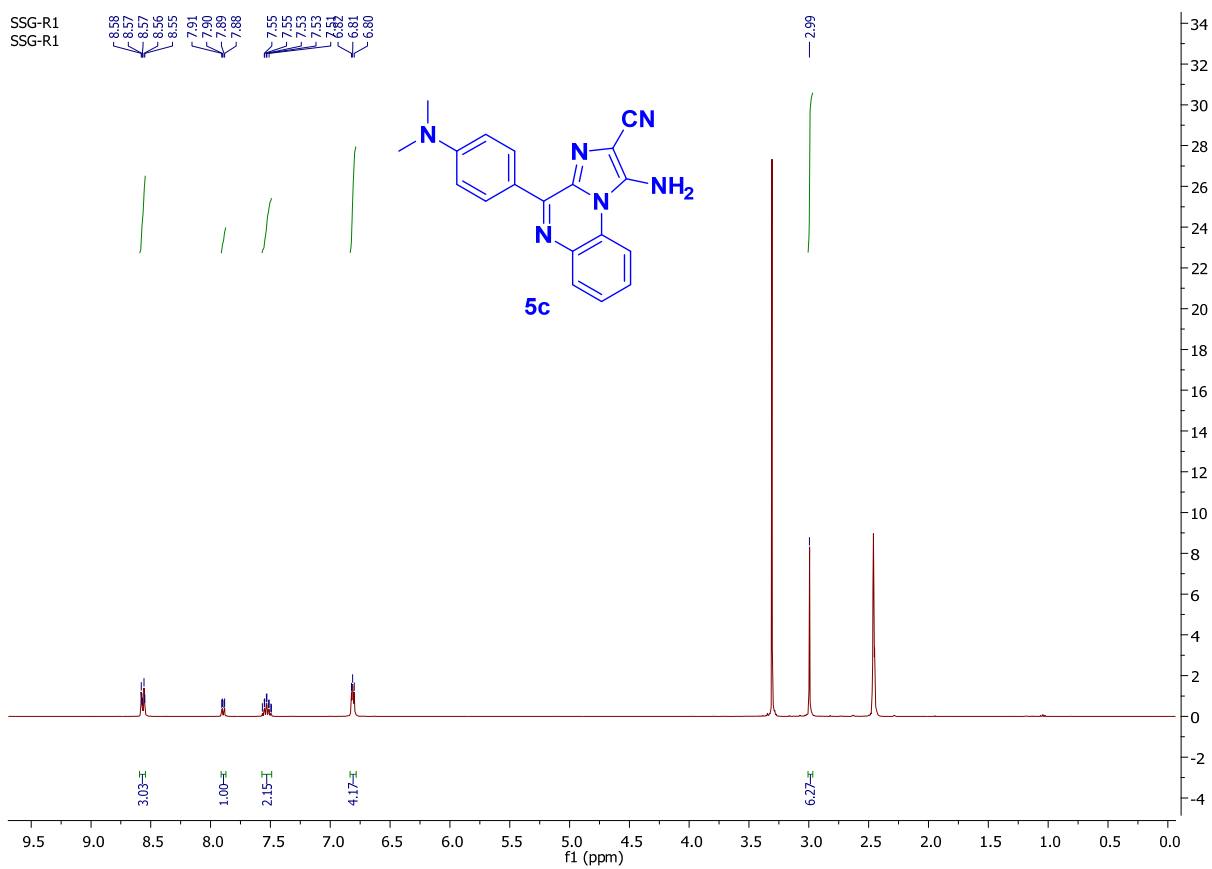


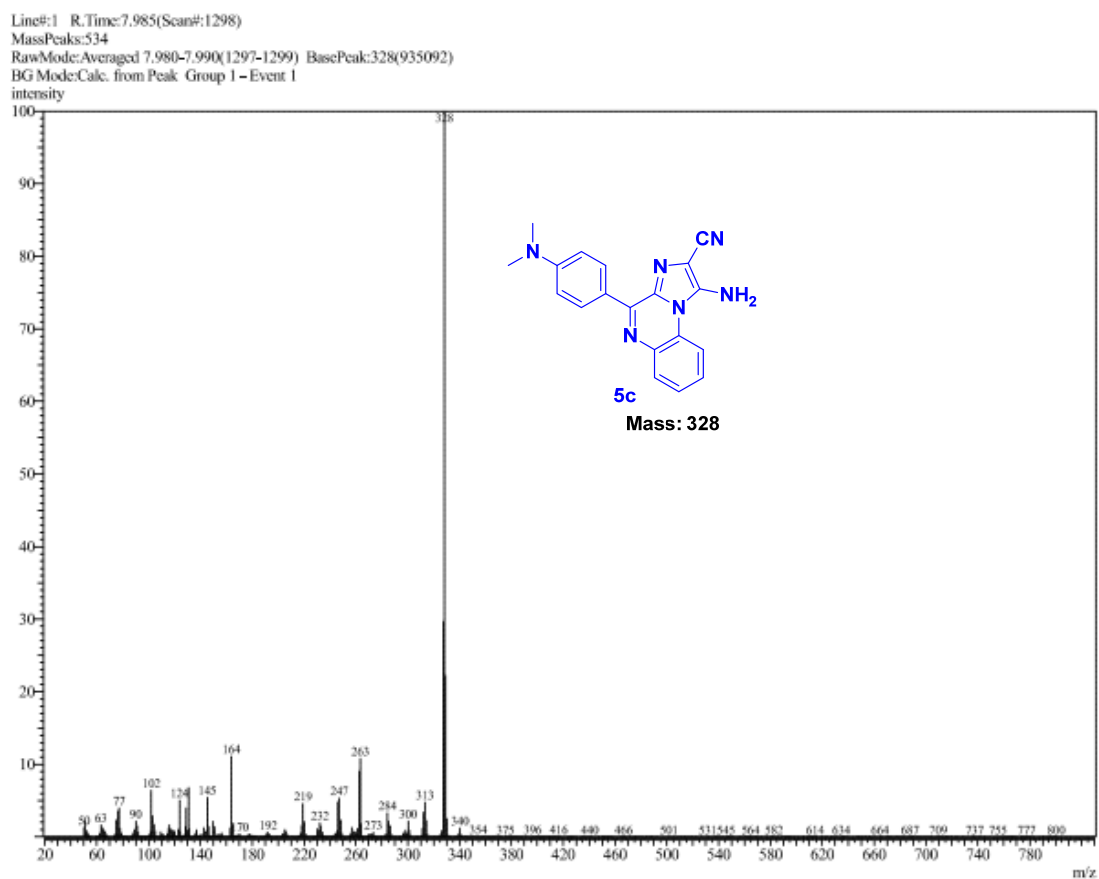
Figure S3. Bar graphs representing alteration in ROS as indicated by H₂DCFDA assay in **A.** MCF-7; **B.** MDA-MB-231 cells. The assays were performed in cancer cells previously treated with investigational compound **6b** at sub-IC₅₀ concentration for 48 h before analysis.

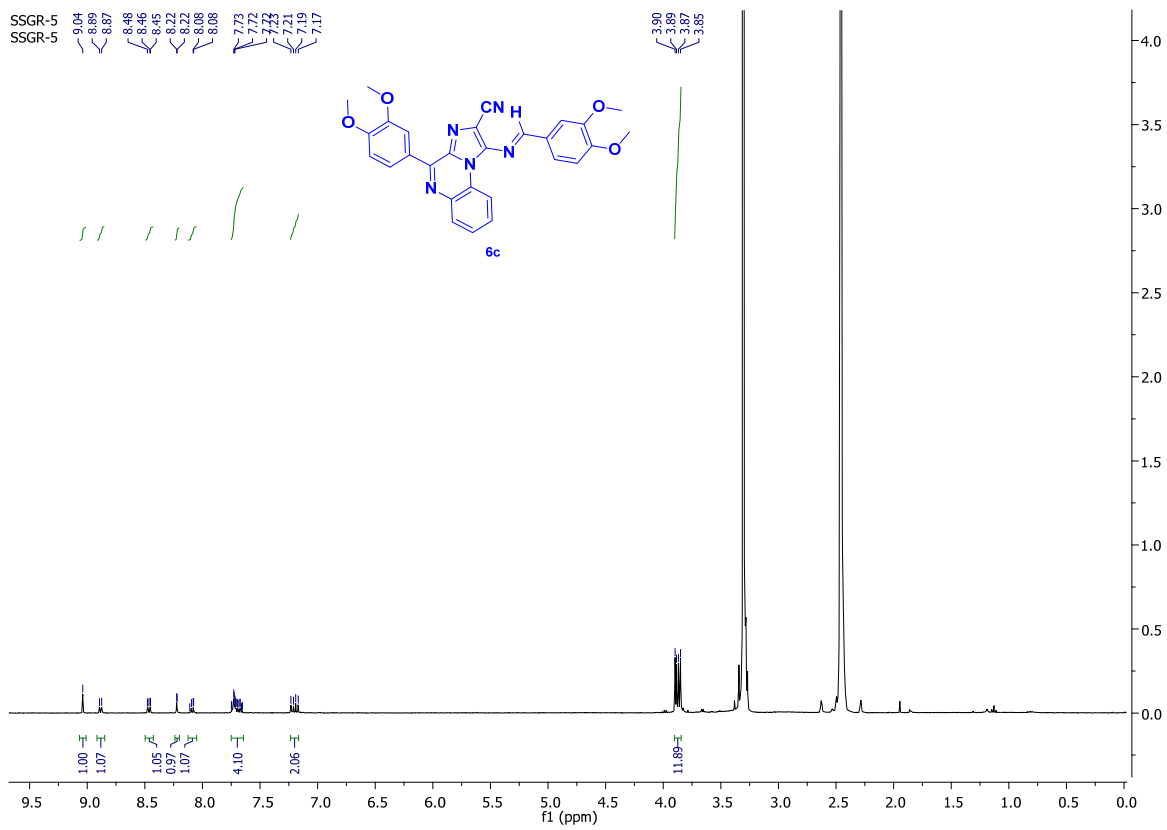
Spectra for representative compounds (Figures S4–S11)

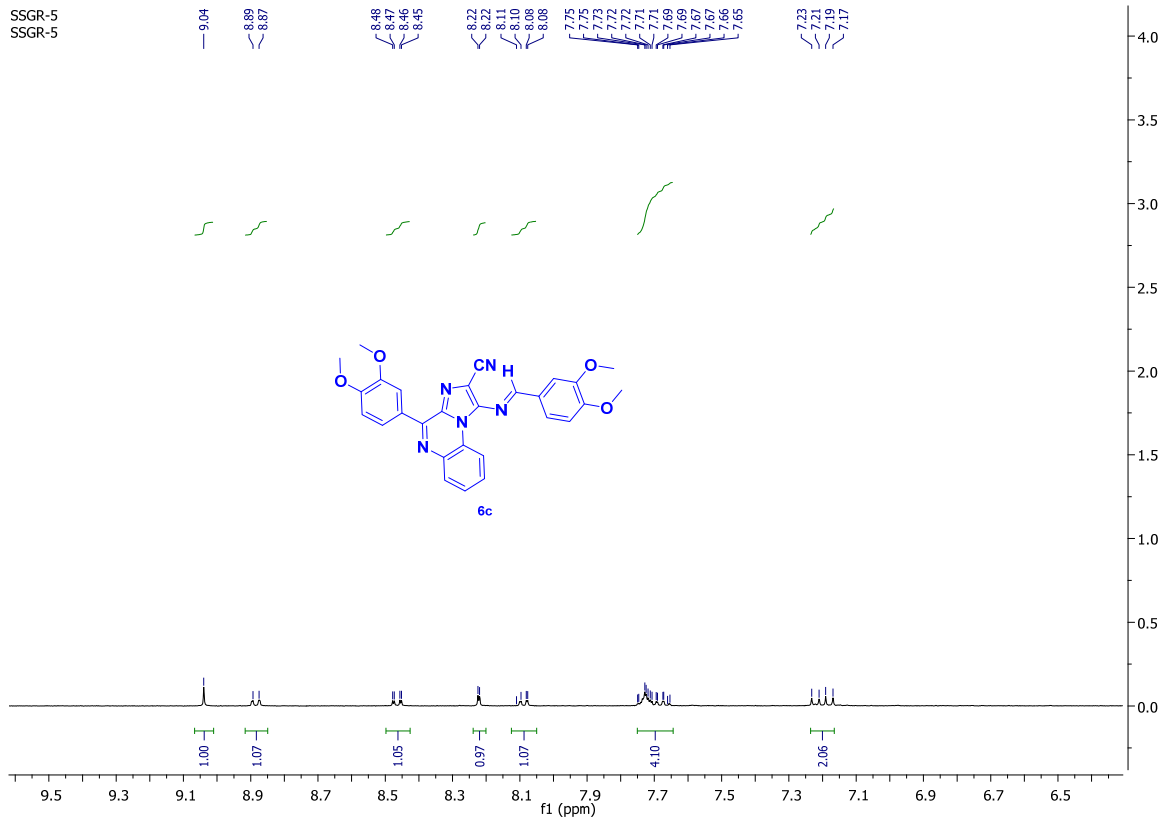
 $^1\text{H-NMR}$ 

Mass Spectrometry

Spectrum



¹H-NMR



Mass Spectrometry

Elemental Composition Report

Single Mass Analysis

Tolerance = 100.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5

Monoisotopic Mass, Even Electron Ions

16 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 9-30 H: 6-25 N: 0-5 O: 0-4

Sample Name : SBD-3

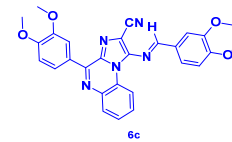
Test Name : HRMS-1

130120-SBD-3- 18 (0.183)

IITRPR

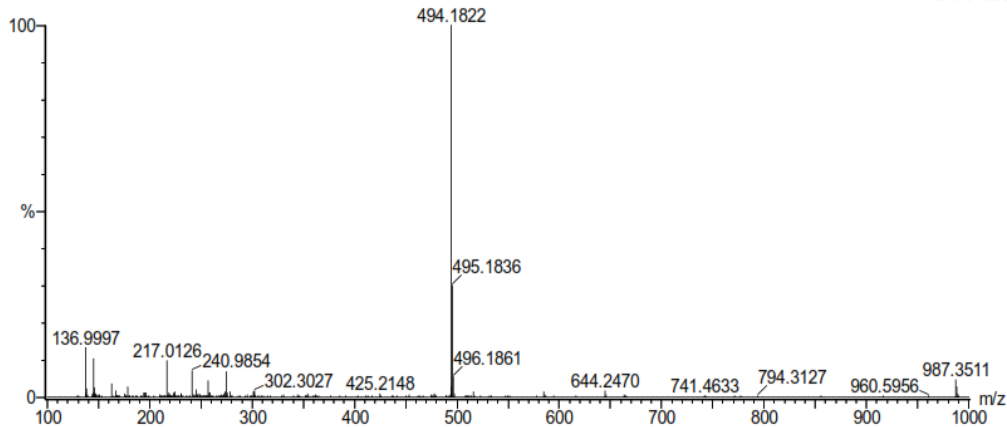
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Exact Mass: 493.1750

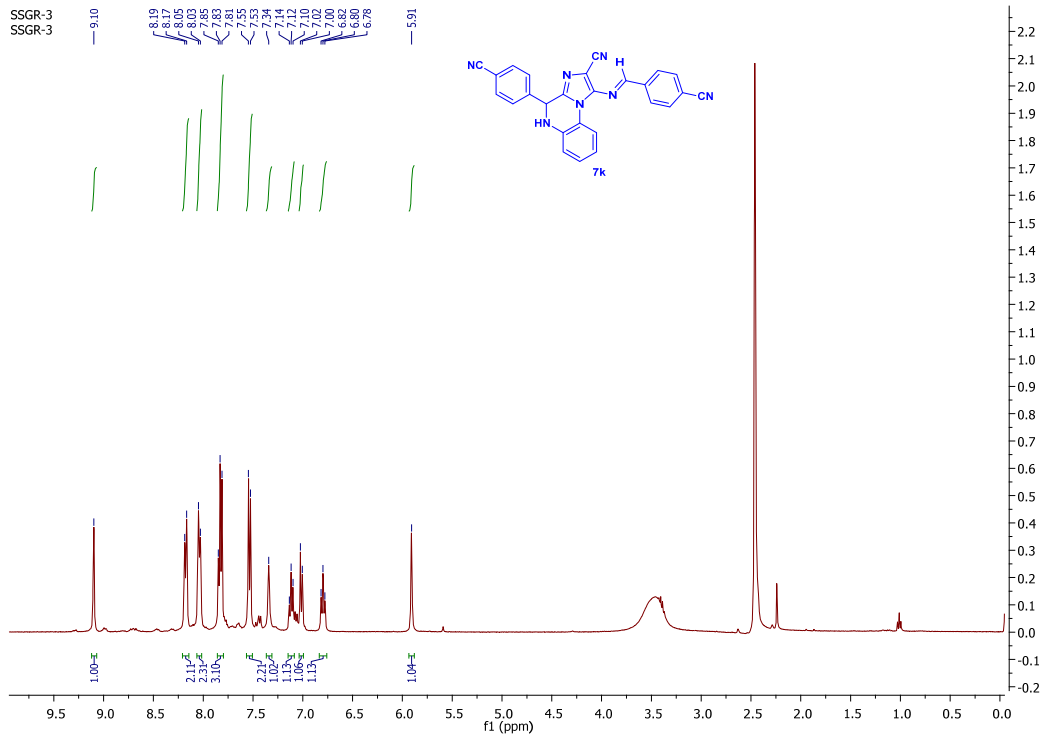


XEVO G2-XS QTOF

1: TOF MS ES+
6.47e+006



¹H-NMR



Mass Spectrometry

Elemental Composition Report

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5

Monoisotopic Mass, Even Electron Ions

8 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 9-30 H: 6-25 N: 0-7

Sample Name : SBD-2

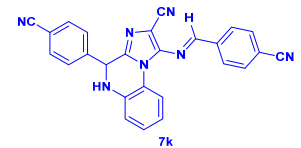
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130120-SBD-2 16 (0.165)

IITRPR

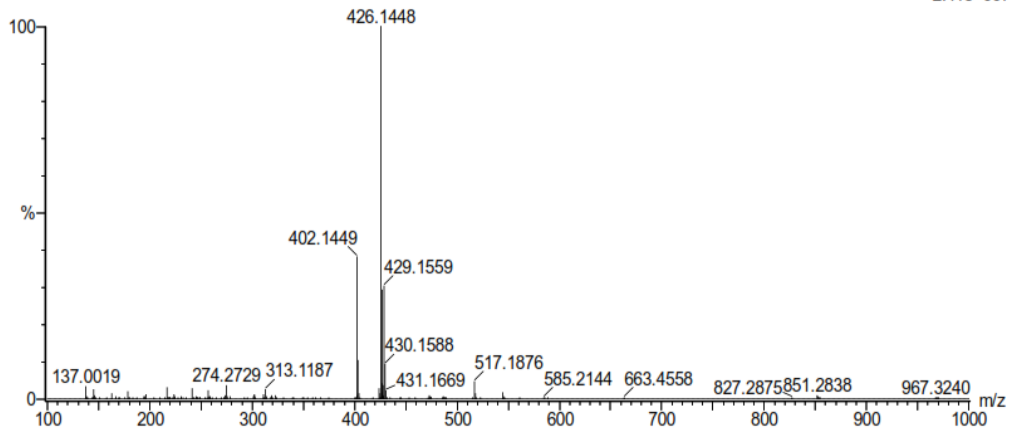
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Exact Mass: 425.1389



XEVO G2-XS QTOF

1: TOF MS ES+
2.11e+007



¹H-NMR