

Electronic Supplementary Information

A new three-component reaction: Green synthesis of novel isoindolo[2,1-*a*]quinazoline derivatives as potent inhibitors of TNF- α

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Experimental

Chemistry

General methods: Unless stated otherwise, reactions were performed under nitrogen atmosphere using oven dried glassware. Reactions were monitored by thin layer chromatography (TLC) on silica gel plates (60 F254), visualizing with ultraviolet light or iodine spray. Flash chromatography was performed on silica gel (230-400 mesh) using distilled hexane, ethyl acetate, dichloromethane. ^1H NMR and ^{13}C NMR spectra were determined in CDCl_3 or $\text{DMSO}-d_6$ solution by using 400 or 500 and 50 or 100 MHz spectrometers, respectively. Proton chemical shifts (δ) are relative to tetramethylsilane (TMS, $\delta = 0.00$) as internal standard and expressed in ppm. Spin multiplicities are given as s (singlet), d (doublet), t (triplet) and m (multiplet) as well as b (broad). Coupling constants (J) are given in hertz. Infrared spectra were recorded on a FT-IR spectrometer. Melting points were determined using melting point apparatus and are uncorrected. MS spectra were obtained on a mass spectrometer. High-resolution mass spectra (HRMS) were recorded using electron ionization (EI) mass spectrometry.

Preparation of **4** via camphor sulfonic acid catalyzed MCR:

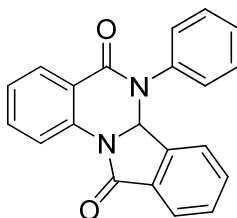
A mixture of isatoic anhydride **1** (1.0 g, 6.13 mmol), amine **2** (6.7mmol), 2-carboxy benzaldehyde **3** (1.01 g, 6.74 mmol) and catalytic amount of anhydrous camphor sulfonic acid (0.07 g, 0.31mmol) in ethanol (10 mL) was heated to 80-85 °C with stirring for 7-16 h (initially effervescence was observed due to the generation of CO_2 gas). After completion of the reaction the mixture was cooled and filtered to give the desired product. The product isolated was found to be pure.

Preparation of **4** via Montmorillonite K10 catalyzed MCR:

A mixture of isatoic anhydride **1** (1.0 g, 6.13 mmol), amine **2** (6.7 mmol), 2-formylbenzoic acid **3** (1.01 g, 6.74 mmol) and catalytic amount of montmorillonite K10 (0.05 g, 5%) in ethanol (10 mL) was stirred at 80-85 °C according to the time mentioned in Table 2 (initially effervescence was observed due to the generation of CO_2 gas). After completion of the reaction the mixture was cooled and diluted with ethanol (20 mL). The catalyst was filtered off. The filtrate was

collected and concentrated. The solid separated was filtered and dried to give the desired product.

6-phenyl-6,6a-dihydroisindolo[2,1-a]quinazoline-5,11-dione (4a)



White color solid, Mp = 183-185 °C

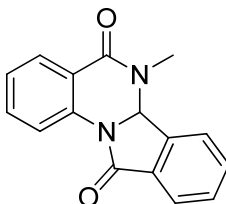
^1H NMR (DMSO- d_6 , 400 MHz): 8.13 (d, $J=8.3$ Hz, 1H), 8.04 (dd, $J=7.8$, 1.6 Hz, 1H), 7.89 (d, $J=8.3$ Hz, 1H), 7.80-7.76 (m, 2H), 7.60-7.36 (m, 6H), 7.02 (s, 1H), 6.8 (br s, 1H), 6.10 (d, $J=7.8$ Hz, 1H)

^{13}C NMR (DMSO- d_6 , 50 MHz): 164.6, 163.2, 139.0, 138.5, 136.8, 133.6, 132.3, 131.4, 130.2, 129.3 (2C), 128.7 (2C), 128.6 (2C), 125.0 (2C), 123.9, 120.2, 120.0, 71.5

IR (KBr): 3421, 3054, 1723, 1655, 1487, 1464 cm^{-1}

HRMS (ESI): calcd for $\text{C}_{21}\text{H}_{15}\text{N}_2\text{O}_2$ ($\text{M}+\text{H}$) $^+$ 327.1134, found 327.1130

6-methyl-6,6a-dihydroisindolo[2,1-a]quinazoline-5,11-dione (4b):



White color solid, Mp = 187-189 °C

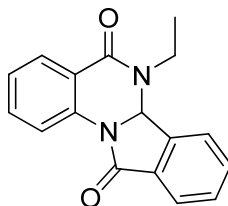
^1H NMR (CDCl_3 , 400 MHz): 8.15 (dd, $J=7.8$, 1.0 Hz, 1H), 8.11 (dd, $J=7.8$, 1.0 Hz, 1H), 8.03 (dd, $J=7.8$, 1.0 Hz, 1H), 7.78-7.61 (m, 4H), 7.35-7.31 (m, 1H), 6.12 (s, 1H), 3.32 (s, 3H, CH_3)

^{13}C NMR (CDCl_3 , 50 MHz): 164.8, 163.8, 137.9, 136.5, 133.3, 132.6, 132.4, 130.4, 128.9, 125.5, 125.0, 124.9, 120.0 (2C), 71.1, 29.9

IR (KBr): 3417, 1720, 1652, 1605, 1474, 1466 cm^{-1}

HRMS (ESI): calcd for $\text{C}_{16}\text{H}_{13}\text{N}_2\text{O}_2$ ($\text{M}+\text{H}$) $^+$ 265.0977, found 265.0969

6-ethyl-6,6a-dihydroisoindolo[2,1-a]quinazoline-5,11-dione (4c):



White color solid, Mp = 155-157 °C

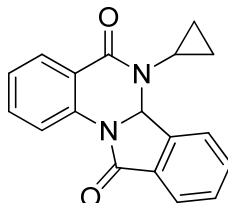
^1H NMR (DMSO- d_6 , 400 MHz): 8.01 (d, $J=7.4$ Hz, 1H), 7.99-7.95 (m, 3H), 7.87-7.84 (m, 1H), 7.77 (d, $J=7.4$ Hz, 1H), 7.74-7.69 (m, 1H), 7.40-7.36 (m, 1H), 6.60 (s, 1H), 3.90-3.83 (m, 1H), 3.72-3.67 (m, 1H), 1.03 (t, $J=7.4$ Hz, 3H, CH_3)

^{13}C NMR (DMSO- d_6 , 100 MHz): 164.3, 162.6, 138.4, 136.4, 133.2, 133.1, 131.8, 130.6, 128.3, 126.0, 125.0, 124.3, 120.3, 119.9, 69.7, 37.2, 13.4

IR (KBr): 3403, 1711, 1668, 1603, 1490, 1469 cm^{-1}

HRMS (ESI): calcd for $\text{C}_{17}\text{H}_{15}\text{N}_2\text{O}_2$ ($\text{M}+\text{H}$) $^+$ 279.1134, found 279.1135

6-cyclopropyl-6,6a-dihydroisoindolo[2,1-a]quinazoline-5,11-dione (4d):



White color solid, Mp = 155-158 °C

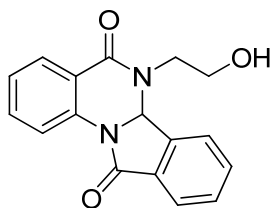
^1H NMR (CDCl_3 , 400 MHz): 8.19-8.14 (m, 2H), 8.05 (d, $J=6.9$ Hz, 1H), 7.92 (d, $J=6.9$ Hz, 1H), 7.70-7.60 (m, 3H), 7.32-7.28 (m, 1H), 6.18 (s, 1H), 2.73-2.68 (m, 1H), 1.19-1.11 (m, 1H), 0.90-0.84 (m, 1H), 0.71-0.64 (m, 1H), 0.08-0.02 (m, 1H)

^{13}C NMR (DMSO- d_6 , 100 MHz): 164.4, 164.3, 138.8, 136.7, 133.3, 132.3, 131.6, 130.2, 128.4, 127.3, 124.6, 123.7, 120.4, 119.2, 71.2, 25.9, 11.1, 9.1

IR (KBr): 3427, 1726, 1663, 1601, 1487, 1466, cm^{-1}

HRMS (ESI): calcd for $\text{C}_{18}\text{H}_{15}\text{N}_2\text{O}_2$ ($\text{M}+\text{H}$) $^+$ 291.1134, found 291.1132

6-(2-hydroxyethyl)-6,6a-dihydroisoindolo[2,1-a]quinazoline-5,11-dione (4e):



White color solid, Mp = 215-217 °C

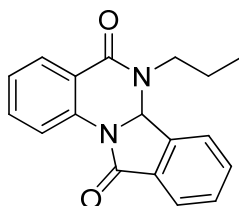
^1H NMR (CDCl_3 , 400 MHz): 8.12-8.08 (m, 2H), 8.05-8.01 (m, 2H), 7.73-7.61 (m, 3H), 7.34-7.30 (m, 1H), 6.42 (s, 1H), 4.18-4.03 (m, 2H), 3.94-3.89 (m, 1H), 3.81-3.74 (m, 1H), 2.48 (br s, 1H, OH)

^{13}C NMR ($\text{DMSO}-d_6$, 100 MHz): 164.3, 163.0, 138.5, 136.5, 133.2, 132.9, 132.0, 130.5, 128.3, 127.3, 125.0, 124.1, 120.3, 119.8, 71.0, 58.9, 44.9

IR (KBr): 3423, 3395, 1723, 1660, 1601, 1486, 1470, 1108 cm^{-1}

HRMS (ESI): calcd for $\text{C}_{17}\text{H}_{15}\text{N}_2\text{O}_3$ ($\text{M}+\text{H}$) $^+$ 295.1083, found 295.1092

6-propyl-6,6a-dihydroisoindolo[2,1-a]quinazoline-5,11-dione (4f):



White color solid, Mp = 127-130 °C

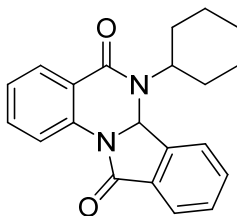
^1H NMR ($\text{DMSO}-d_6$, 500 MHz): 8.01-7.95 (m, 4H), 7.87-7.84 (m, 1H), 7.77-7.69 (m, 2H), 7.39-7.36 (m, 1H), 6.57 (s, 1H), 3.73-3.67 (m, 2H), 1.46-1.29 (m, 1H), 1.27-1.22 (m, 1H), 0.76 (t, J=7.3Hz, 3H, CH_3)

^{13}C NMR ($\text{DMSO}-d_6$, 100 MHz): 164.3, 162.7, 138.5, 136.4, 133.1, 133.2, 131.8, 130.7, 128.4, 126.0, 125.0, 124.3, 120.3, 119.9, 70.0, 43.7, 21.3, 10.8

IR (KBr): 3423, 2960, 1728, 1653, 1489, 1470 cm^{-1}

HRMS (ESI): calcd for $\text{C}_{18}\text{H}_{17}\text{N}_2\text{O}_2$ ($\text{M}+\text{H}$) $^+$ 293.1290, found 293.1286

6-cyclohexyl-6,6a-dihydroisoindolo[2,1-a]quinazoline-5,11-dione (4g):



White color solid, Mp = 148-150 °C

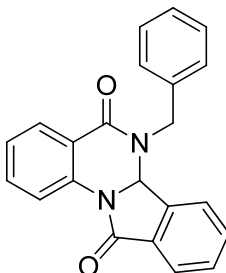
^1H NMR (DMSO- d_6 , 500 MHz): 8.01 (d, $J=7.9$ Hz, 1H), 8.00-7.95 (m, 2H), 7.87-7.86 (m, 2H), 7.80-7.68 (m, 2H), 7.38-7.34 (m, 1H), 6.58 (s, 1H), 3.58-3.53 (m, 1H), 2.51-2.39 (m, 2H), 2.15-2.05 (m, 1H), 1.91-1.01 (m, 7H)

^{13}C NMR (DMSO- d_6 , 100 MHz): 164.0, 163.4, 138.5, 136.3, 133.2, 133.0, 132.2, 130.7, 128.2, 126.9, 124.8, 124.4, 121.4, 119.3, 71.1, 57.6, 29.9, 28.1, 26.0, 25.6, 25.0

IR (KBr): 3428, 2965, 1735, 1659, 1495, 1478 cm^{-1}

HRMS (ESI): calcd for $\text{C}_{21}\text{H}_{21}\text{N}_2\text{O}_2$ ($\text{M}+\text{H}$) $^+$ 333.1603, found 333.1602

6-benzyl-6,6a-dihydroisoindolo[2,1-a]quinazoline-5,11-dione (4h):



White color solid, Mp = 148-150 °C

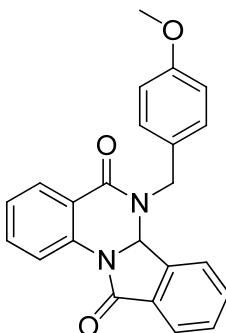
^1H NMR (DMSO- d_6 , 400 MHz): 8.06 (d, $J=7.9$ Hz, 2H), 7.87-7.84 (m, 1H), 7.77-7.72 (m, 2H), 7.65-7.59 (m, 2H), 7.44-7.40 (m, 1H), 7.22-7.05 (m, 5H), 6.72 (s, 1H), 5.11-5.00 (m, 2H)

^{13}C NMR (DMSO- d_6 , 100 MHz): 164.3, 163.3, 138.0, 137.3, 136.7, 133.5, 132.9, 131.7, 130.5, 128.6, 128.4 (2C), 126.6, 126.0, 125.9 (2C), 125.1, 124.1, 120.0, 119.9, 70.2, 45.7

IR (KBr): 3422, 3055, 1723, 1660, 1486, 1470 cm^{-1}

HRMS (ESI): calcd for $\text{C}_{22}\text{H}_{17}\text{N}_2\text{O}_2$ ($\text{M}+\text{H}$) $^+$ 341.1290, found 341.1282

6-(4-methoxybenzyl)-6,6a-dihydroisindolo[2,1-a]quinazoline-5,11-dione (4i):



White color solid, Mp = 137-139 °C

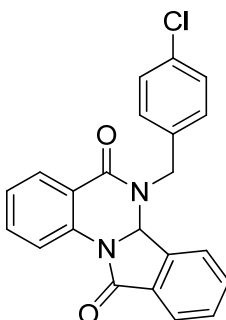
¹H NMR (DMSO-*d*₆, 400 MHz): 8.06-8.04 (m, 2H), 7.88-7.86 (m, 1H), 7.77-7.62 (m, 4H), 7.43-7.39 (m, 1H), 6.95 (d, J-8.8 Hz, 2H), 6.77 (d, J-8.8 Hz, 2H), 6.68 (s, 1H), 5.04-4.89 (m, 2H), 3.66 (s, 3H, CH₃)

¹³C NMR (DMSO-*d*₆, 100 MHz): 164.3, 163.2, 157.9, 138.1, 136.6, 133.4, 132.9, 131.7, 130.5, 129.0, 128.6, 127.3 (2C), 126.1, 125.1, 124.1, 120.1, 119.8, 113.8 (2C), 70.2, 54.9, 45.1

IR (KBr): 3428, 2833, 1724, 1660, 1488, 1466 cm⁻¹

HRMS (ESI): calcd for C₂₃H₁₉N₂O₃ (M+H)⁺ 371.1396, found 371.1398

6-(4-chlorobenzyl)-6,6a-dihydroisindolo[2,1-a]quinazoline-5,11-dione (4j):



White color solid, Mp = 178-181 °C

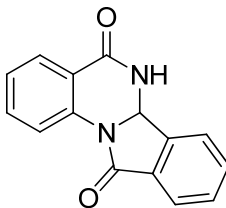
¹H NMR (DMSO-*d*₆, 400 MHz): 8.06-8.04 (m, 2H), 7.86-7.84 (m, 1H), 7.78-7.73 (m, 2H), 7.65-7.61 (m, 2H), 7.44-7.40 (m, 1H), 7.24 (d, J-8.8 Hz, 2H), 7.07 (d, J-8.8 Hz, 2H), 6.71 (s, 1H), 5.12-4.99 (m, 2H)

¹³C NMR (DMSO-*d*₆, 100 MHz): 164.3, 163.4, 138.0, 136.7, 136.6, 133.6, 133.0, 131.7, 131.1, 130.6, 128.7, 128.3 (2C), 127.9 (2C), 126.1, 125.2, 124.1, 120.0 (2C), 70.2, 45.1

IR (KBr): 3429, 2932, 1726, 1664, 1490, 1470 cm⁻¹

HRMS (ESI): calcd for $C_{22}H_{16}ClN_2O_2$ (M+H)⁺ 375.0900, found 375.0902

6,6a-dihydroisoindolo[2,1-a]quinazoline-5,11-dione (4k)



White color solid, Mp = 240-242 °C

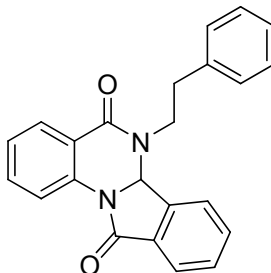
¹H NMR (DMSO-*d*₆, 400 MHz): 9.29 (s, 1H, NH), 8.13 (d, J=7.7 Hz, 1H), 8.05 (dd, J=7.9 Hz, 1.2 Hz, 1H), 7.90 (d, J=7.7 Hz, 2H), 7.75-7.61 (m, 3H), 7.33-7.29 (m, 1H), 6.33 (s, 1H)

¹³C NMR (DMSO-*d*₆, 100 MHz): 164.4, 163.6, 140.7, 137.1, 133.5, 133.2, 131.1, 130.1, 128.2, 124.7, 124.1, 123.8, 120.0, 119.5, 67.0

IR (KBr): 3058, 2930, 1731, 1676, 1495, 1475 cm⁻¹

HRMS (ESI): calcd for $C_{15}H_{11}N_2O_2$ (M+H)⁺ 251.0821, found 251.0830

6-phenethyl-6,6a-dihydroisoindolo[2,1-a]quinazoline-5,11-dione (4l):



White color solid, Mp = 123-125 °C

¹H NMR (DMSO-*d*₆, 400 MHz): 8.05 (d, J=7.4 Hz, 1H), 8.02-7.98 (m, 2H), 7.92 (d, J=7.4 Hz, 1H), 7.90-7.86 (m, 1H), 7.77 (d, J=7.4 Hz, 1H), 7.74-7.69 (m, 1H), 7.41-7.37 (m, 1H), 7.24-7.14 (m, 3H), 7.08-7.06 (m, 2H), 6.57 (s, 1H), 4.02-3.88 (m, 2H), 2.80-2.73 (m, 1H), 2.50-2.42 (m, 1H)

¹³C NMR (DMSO-*d*₆, 100 MHz): 164.3, 162.8, 138.3, 138.2, 136.5, 133.3, 133.2, 132.0, 130.7, 128.5, 128.4, 128.3, 128.2, 127.4, 126.3, 125.1, 124.9, 124.4, 120.3, 120.0, 70.0, 43.8, 34.1

IR (KBr): 3060, 2932, 1734, 1680, 1497, 1479 cm⁻¹

HRMS (ESI): calcd for $C_{23}H_{19}N_2O_2$ (M+H)⁺ 355.1447, found 355.1438

Single crystal X-ray data for compound **4j**

Single crystals suitable for X-ray diffraction of **4j** were grown from methanol. The crystals were carefully chosen using a stereo zoom microscope supported by a rotatable polarizing stage. The data was collected at room temperature on Bruker's KAPPA APEX II CCD Duo with graphite monochromated Mo-K α radiation (0.71073 Å). The crystals were glued to a thin glass fibre using FOMBLIN immersion oil and mounted on the diffractometer. The intensity data were processed using Bruker's suite of data processing programs (SAINT), and absorption corrections were applied using SADABS.¹ The crystal structure was solved by direct methods using SHELXS-97 and the data was refined by full matrix least-squares refinement on F^2 with anisotropic displacement parameters for non-H atoms, using SHELXL-97.²

Crystal data of **4j**: Molecular formula = C₂₂H₁₅ClN₂O₂, Formula weight = 374.1, Crystal system = Monoclinic, space group = $P2_1/c$, $a = 11.6223$ (6) Å, $b = 7.8255$ (4) Å, $c = 19.9457$ (10) Å, $V = 1772.91$ (16) Å³, $T = 296$ K, $Z = 4$, $D_c = 1.404$ Mg m⁻³, $\mu(\text{Mo-K}\alpha) = 0.71073$ mm⁻¹, 27323 reflections measured, 3833 independent reflections, 3311 observed reflections [$I > 2.0 \sigma(I)$], $R_{1_obs} = 0.047$, Goodness of fit = 1.030. Crystallographic data (excluding structure factors) for **4j** have been deposited with the Cambridge Crystallographic Data Centre as supplementary publication numbers CCDC 807140.

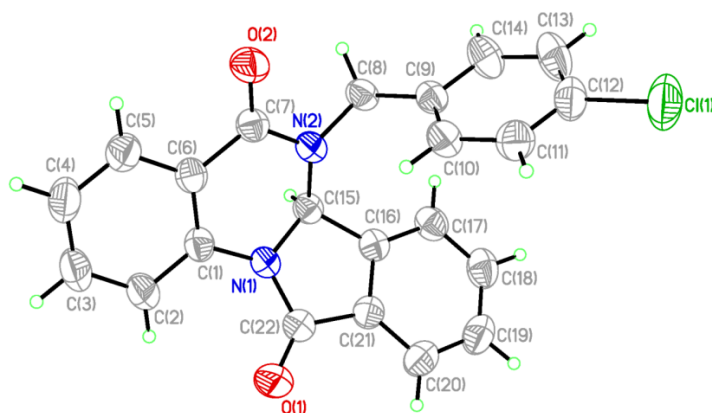


Figure 1. X-ray crystal structure of **4j** (ORTEP diagram). Thermal ellipsoids are drawn at 50% probability level.

Single crystal X-ray data for 6-(3-chlorophenyl)-6,6a-dihydroisindolo[2,1-a]quinazoline-5,11-dione

Single crystals suitable for X-ray diffraction of 6-(3-chlorophenyl)-6,6a-dihydroisindolo[2,1-a]quinazoline-5,11-dione [prepared by via the reaction of isatoic anhydride (1.0 g, 6.13 mmol), 3-chloroaniline (6.7 mmol), 2-formylbenzoic acid (1.01 g, 6.74 mmol) in the presence of catalytic amount of montmorillonite K10 (0.05 g, 5%) in ethanol (10 mL) at 80-85 °C for 15h] were grown from methanol and the X-ray analysis data were generated following the procedure as described above.

Crystal data of 6-(3-chlorophenyl)-6,6a-dihydroisindolo[2,1-a]quinazoline-5,11-dione: Molecular formula = $C_{22}H_{17}ClN_2O_3$, Formula weight = 392.83, Crystal system = Orthorhombic, space group = *Pbca*, $a = 21.631(5) \text{ \AA}$, $b = 7.564(12) \text{ \AA}$, $c = 22.783(4) \text{ \AA}$, $V = 3728.1(12) \text{ \AA}^3$, $T = 296 \text{ K}$, $Z = 8$, $D_c = 1.389 \text{ Mg m}^{-3}$, $\mu(\text{Mo-K}\alpha) = 0.71073 \text{ mm}^{-1}$, 8092 reflections measured, 1634 independent reflections, 1098 observed reflections [$I > 2.0 \sigma(I)$], $R_1_{\text{obs}} = 0.079$, Goodness of fit = 1.124.

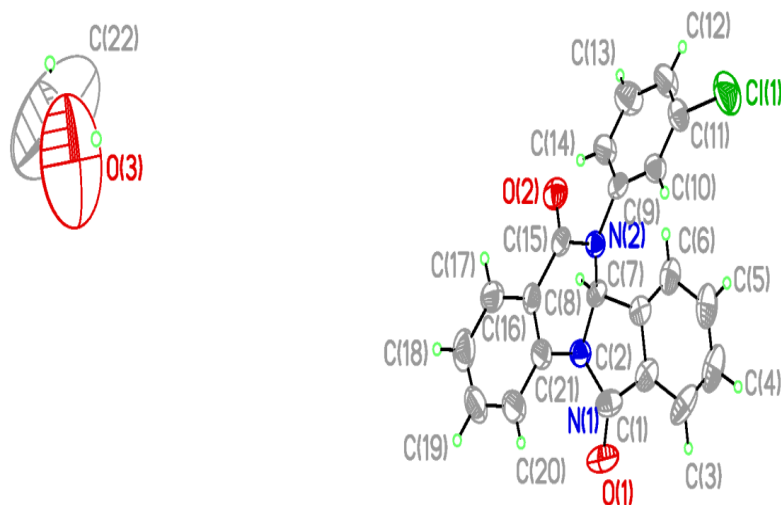


Figure 2. X-ray crystal structure of 6-(3-chlorophenyl)-6,6a-dihydroisindolo[2,1-a]quinazoline-5,11-dione (ORTEP diagram). Thermal ellipsoids are drawn at 50% probability level.

Reference

1. Bruker SADABS V2008-1, Bruker AXS.: Madison, WI, USA, **2008**.

2. Sheldrick, G. M.; SHELX-97, Program for Crystal Structure Determination, University of Göttingen, 1997.

Pharmacology

Materials and Methods

Cells and reagents

RAW 264.7 cells (murine macrophage cell line) were obtained from ATCC (Washington D.C., USA) and routinely maintained in RPMI 1640 medium with 10% fetal bovine serum (Invitrogen Inc., San Diego, CA, USA). Lipopolysaccharide (LPS) was from *Escherichia coli* strain 0127:B8 obtained from Sigma (St. Louis, MO, USA). Mouse TNF- α ELISA kit was procured from R&D Systems (Minneapolis, MN, USA).

TNF- α production assay

The production of TNF- α is measured following a procedure described previously after few modifications.¹ Briefly, RAW 264.7 cells were pre-incubated either with DMSO (vehicle control) or compound for 30 minutes and then stimulated with 1 μ g/ml of LPS overnight. Preliminary screening of the compounds was performed at 30 μ M and dose response studies were carried out at eight different concentrations (30, 10, 3, 1, 0.3, 0.1, 0.03, 0.01 μ M). Post-stimulation, cell supernatants were harvested, centrifuged to clear cell debris and the amount of TNF- α in the supernatants was measured using mouse TNF- α DuoSet ELISA kit from R&D Systems according to manufacturer's recommendations. The percentage of inhibition was calculated using the following formula:

$$\% \text{ inhibition} = 100 - \left| \frac{(\text{LPS stimulated}_{\text{compound}} - \text{unstimulated})}{(\text{LPS stimulated}_{\text{DMSO}} - \text{unstimulated})} \times 100 \right|$$

The IC₅₀ values were determined by a nonlinear regression analysis from dose response curve using Graphpad Prism software (San Diego, U.S.A). IC₅₀ values are expressed as mean $\pm$ SD.

Reference:

1. K. V. Parsa, L. P. Ganesan, M. V. Rajaram, M. A. Gavrilin, A. Balagopal, N. P. Mohapatra, M. D. Wewers, L. S. Schlesinger, J. S. Gunn and S. Tridandapani, *PLoS Pathog.* 2006, 2:e71.

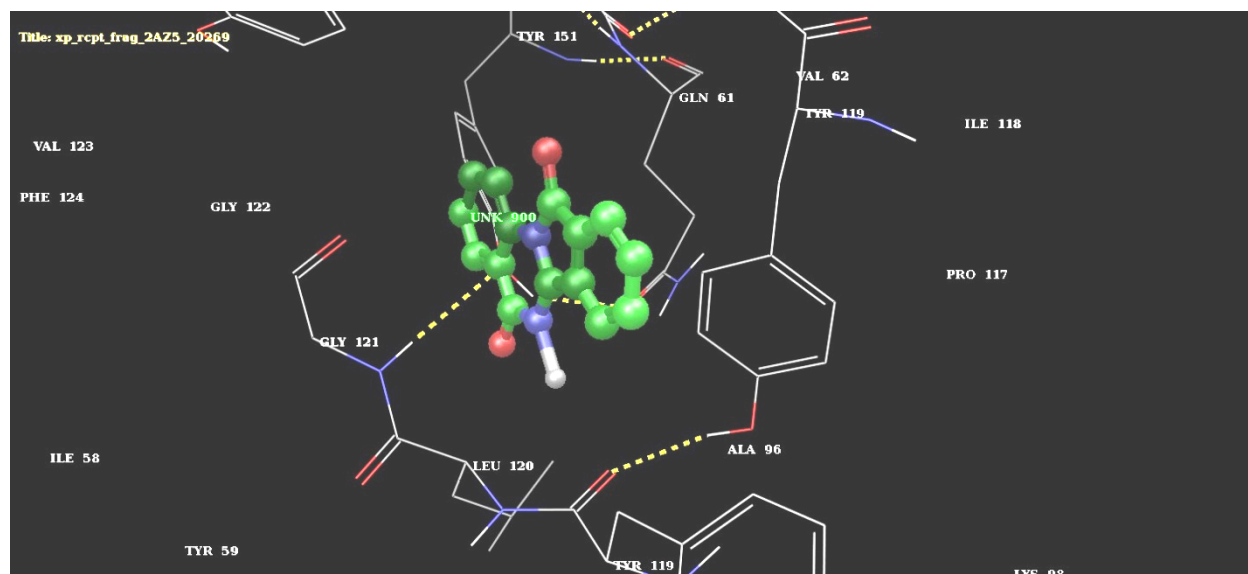
Docking studies

Docking Procedure: In the present study we have performed the energy minimization and conformational search with the MACROMODEL application in the Schrodinger package with MASTERO 9.1 version interface. The ligand molecules were energy minimized for flexibility followed by the conformational search. We used OPLS_2005 force field and water as implicit solvent. We have followed the PRCG (Polak-Ribier conjugate gradient) method of minimization with 500 iterations with a threshold gradient on 0.05kJ/mol. The conformational search was based on Montecarlo multiple minimum torsional sampling. The ligand molecules were then finally prepared with LIGPREP application.

The protein 2AZ5 (TNF- α) crystal structure was retrieved from the protein data bank and was refined with the PROTEIN PREPERATION WIZARD application in which the hydrogen atoms were added and missing side chains and loops were filled with PRIME application. The water molecules were observed within the 5A distance and other water molecules were deleted beyond 5A from the het(hetroatom) groups. Finally the protein was then optimized and minimized with impref using OPLS_2005 force filed.

GRID based Docking was carried out in the present study.

Docking of compound **4k** with TNF- α protein:



GLIDE SCORE = -8.57 Kcal/mol

Hydrophobic ensure reward = - 0.55 Kcal/mol

Total protein-ligand Van der Waals energy = - 4.50 K.cal/mol

Copies of spectra

A new three-component reaction: “Green” synthesis of novel isoindolo[2,1-*a*]quinazoline derivatives as potent inhibitors of TNF- α

K. Siva Kumar,^{a,b} P. Mahesh Kumar,^a K. Anil Kumar,^a M. Sreenivasulu,^a Ahamed A. Jafar,^b
D. Rambabu,^c G. Rama krishna,^d C. Malla Reddy,^d Ravikumar Kapavarapu,^c K. Shivakumar,^c
K. Krishna Priya,^c Kishore V. L. Parsa,^c Manojit Pal^{c,*}

^a*Custom Pharmaceutical Services, Dr. Reddy's Laboratories Limited, Bollaram Road Miyapur,
Hyderabad 500 049, India*

^b*PG and Research Department of Chemistry, Jamal Mohamed College, Tiruchirappalli 620020,
Tamil Nadu, India*

^c*Institute of Life Sciences, University of Hyderabad Campus, Gachibowli, Hyderabad 500 046,
India*

^d*Department of Chemical Sciences, Indian Institute of Science Education and Research, Kolkata,
West Bengal, 741252, India.*

495116

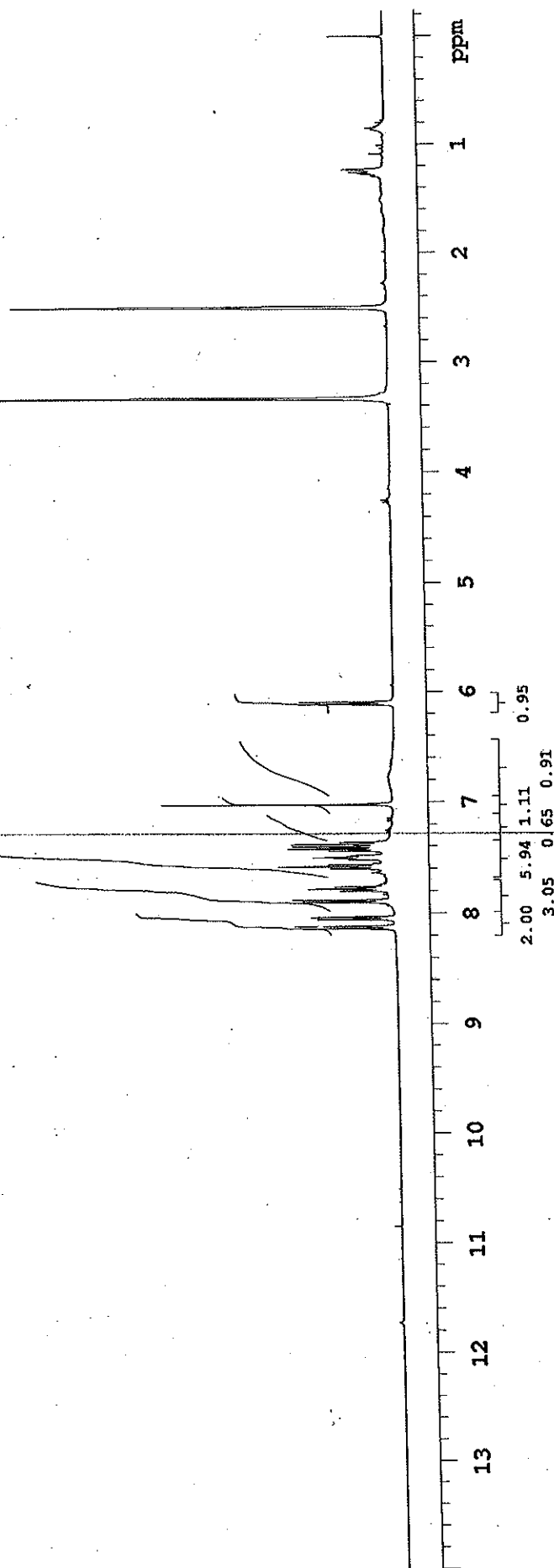
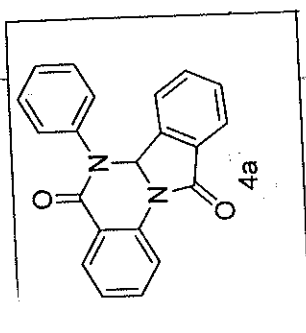
TDC-204 A558/QDR/05 in DMSO

NMR-400

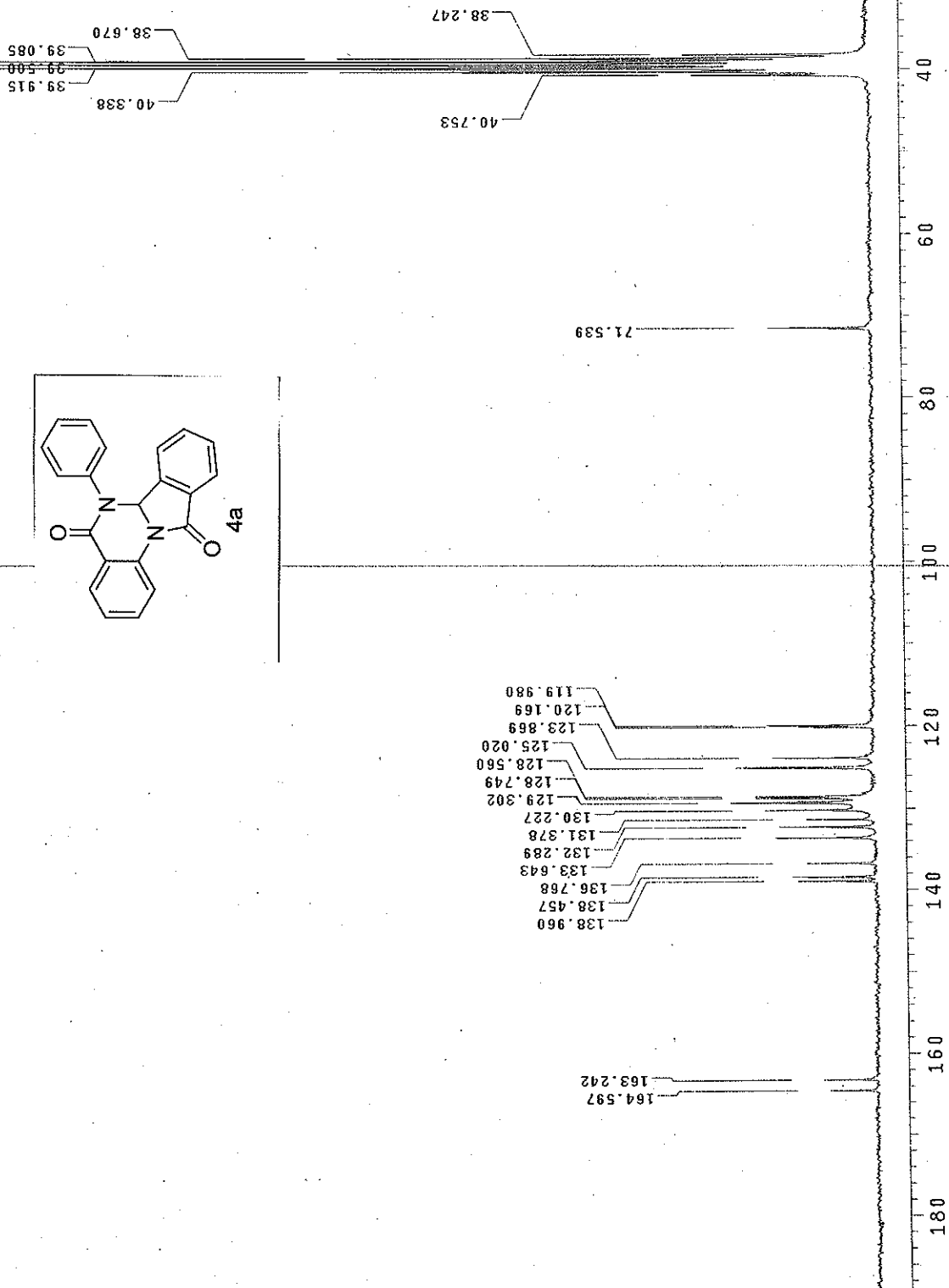
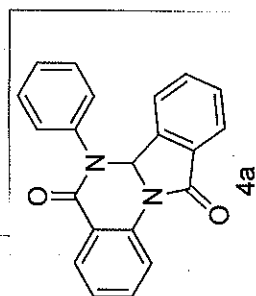
AR.No:ME1010/15

Analyst:Haribabu.R

Date: 01 st Oct 2010



AR&D, Aurigene Discovery Technologies Ltd, Hyderabad
Instrument : Gemini 2000 (Varian 200MHz)
Date & Time : Fri Dec 3 12:14:08 GMT 2010
Recorded By : Haribabu.R



A558-QDR-05 in DMSO
TDC-204

AR NO:GE1210/05
Analyst:Haribabu.R
Date: 2nd Dec. 2010

Elemental Composition Report

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.0, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

109 formula(e) evaluated with 1 results within limits (up to 4 best isotopic matches for each mass)

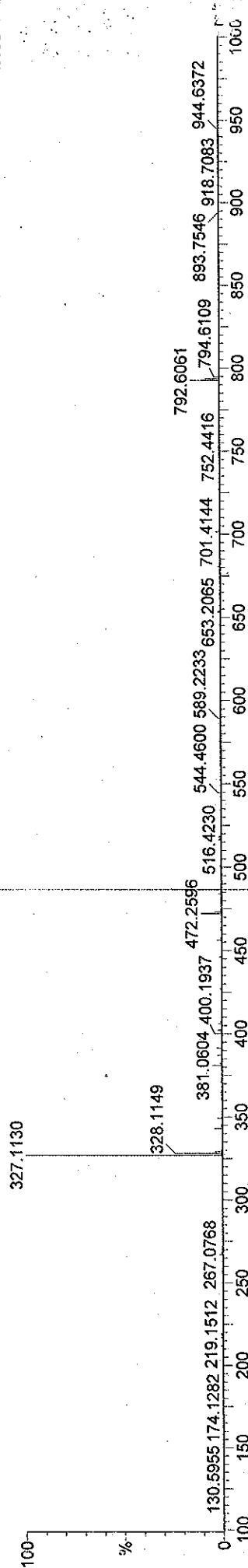
Elements Used:

C: 0-45 H: 0-55 N: 0-2 O: 0-8

A558-QDR-05

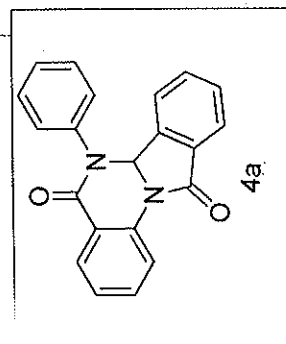
UT1110_66 26 (0.487) Cm (26:35-81:93)

1: TOF MS
4.13e+06



Minimum:
Maximum:

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Formula
327.1130	327.1134	-0.4	-1.2	15.5	1.5	C21 H15 N2 O2



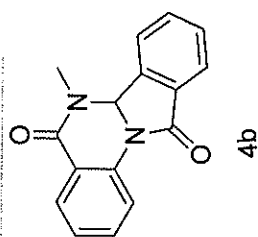
TDC-204-A558-QDR-26 in CDCl₃

NMR-400

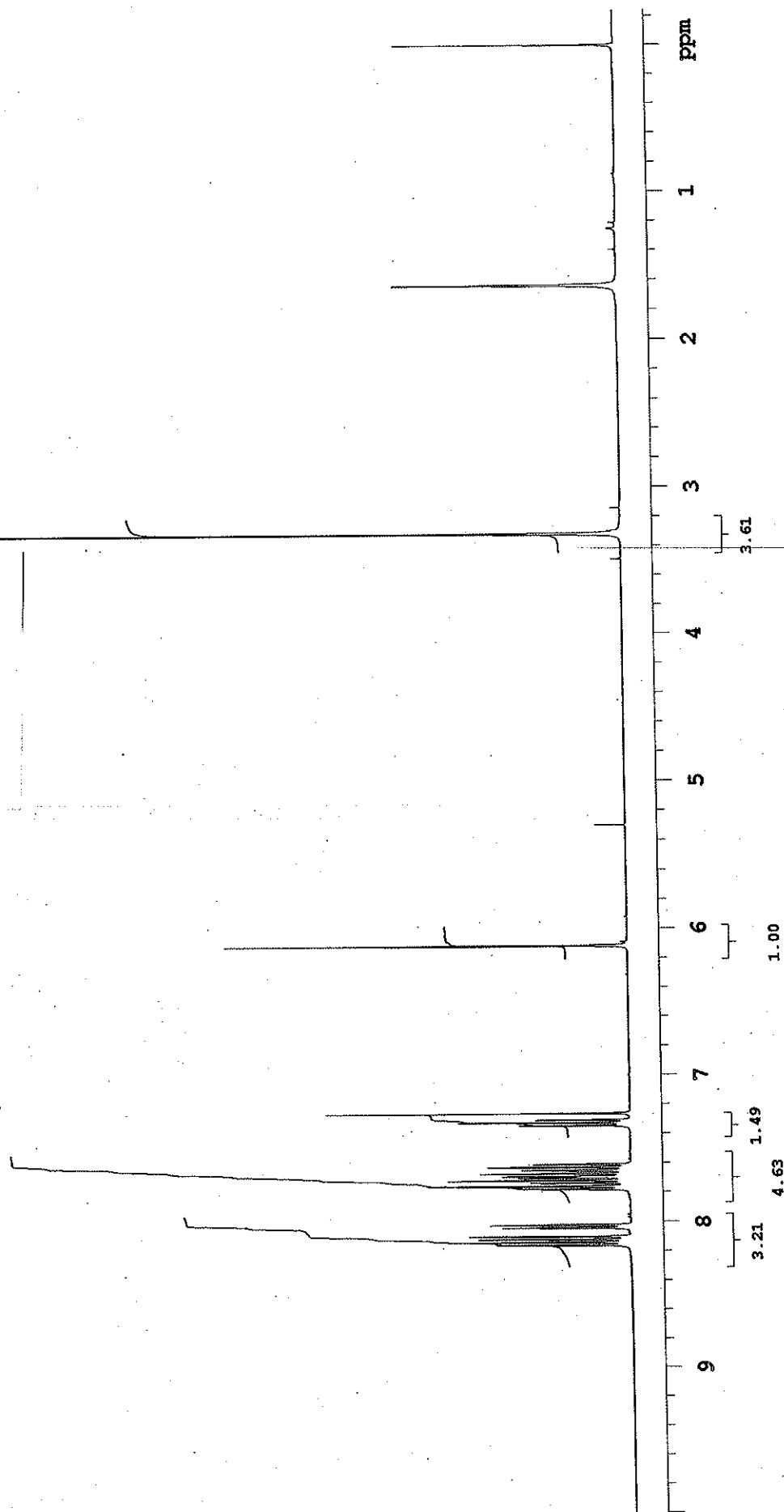
Ac.No:ME1010/1267

Analyst: Shruthi

Date: 15th Oct. 2010



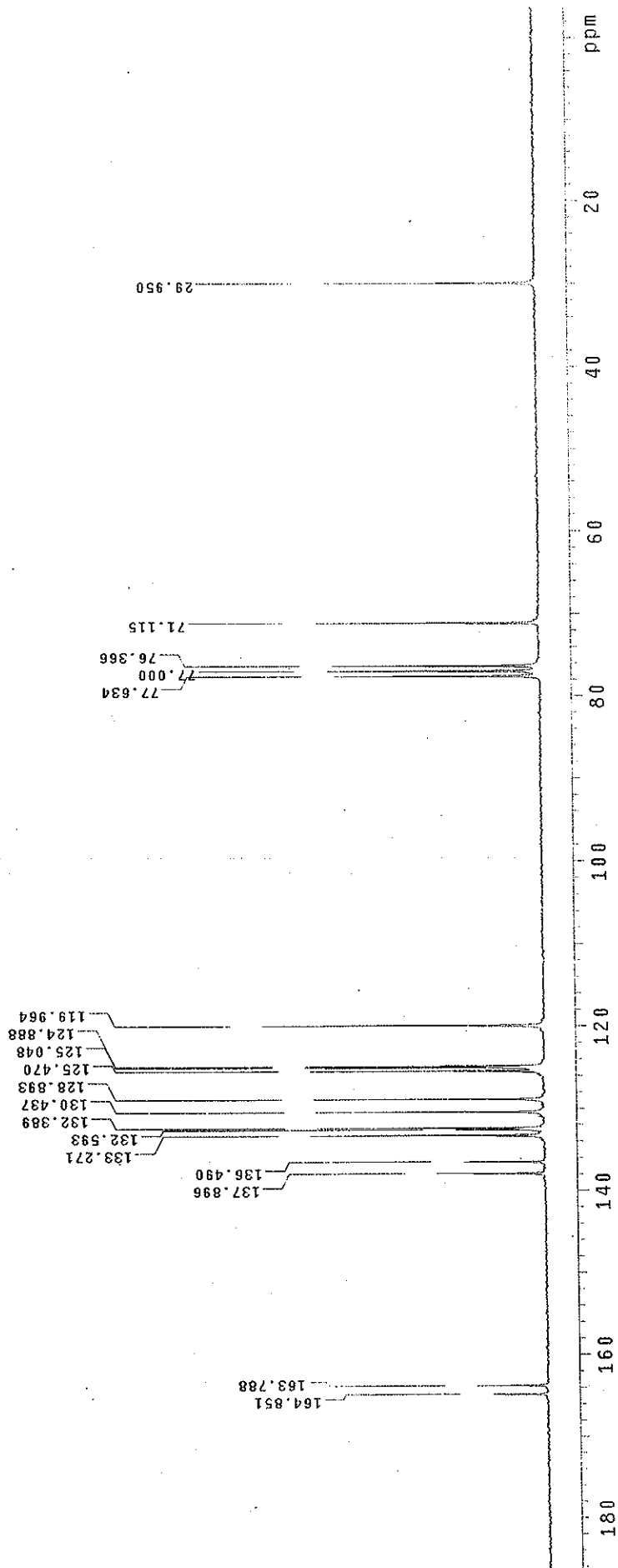
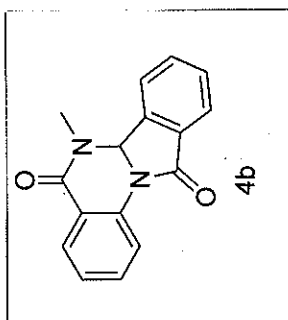
4b



7/11/10
S

AR&D. Aurigene Discovery Technologies Ltd, Hyderabad
Instrument : Gemini 2000 (Varian 200MHz)
Date & Time : Wed Oct 20 20:17:15 GMT 2010
Recorded By : Haribabu.R

ms
2d/10



A558/0DR/26 in CDC13
TDC-204

AR NO: GE1010/54
Analyst: Haribabu.R
Date: 20 th Oct 2010

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.0, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

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

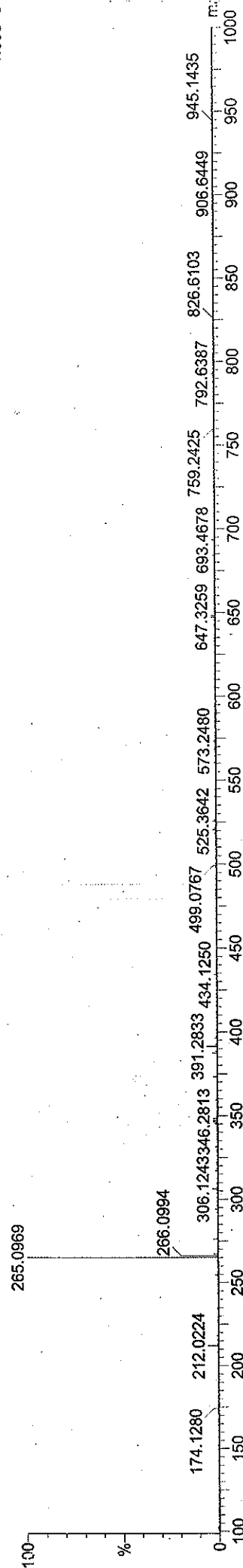
Elements Used:

C: 0-45 H: 0-70 N: 0-3 O: 0-3

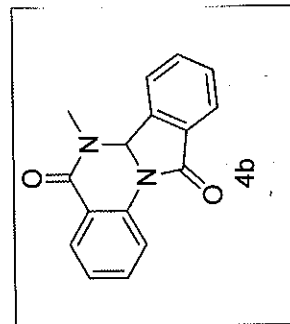
AE58QDR26

U11010_197 22 (0.413) Cm (21:24-85:94)

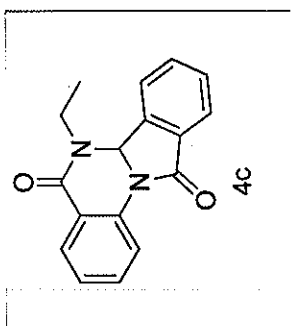
1: TOF MS ES
1.05e+0



Minimum:	5.0	5.0	-1.0
Maximum:			80.0
Mass	Calc. Mass	mDa	DBE
265.0969	265.0977	-0.8	11.5
		-3.0	0.4
Formula	C16 H13 N2 O2		



VARIAN



TDC-204 A558_QDR/32 in DMSO

NMR-400

AR.No:ME1110/901

Analyst:Haribabu.R

Date: 15 th Nov 2010

Sample Name:

901-TDC-204_A558_QDR_32

Data Collected on:

mercury-j-mercury400

Archive directory:

/home/sms50/routine/Nov2010/d15

Sample directory:

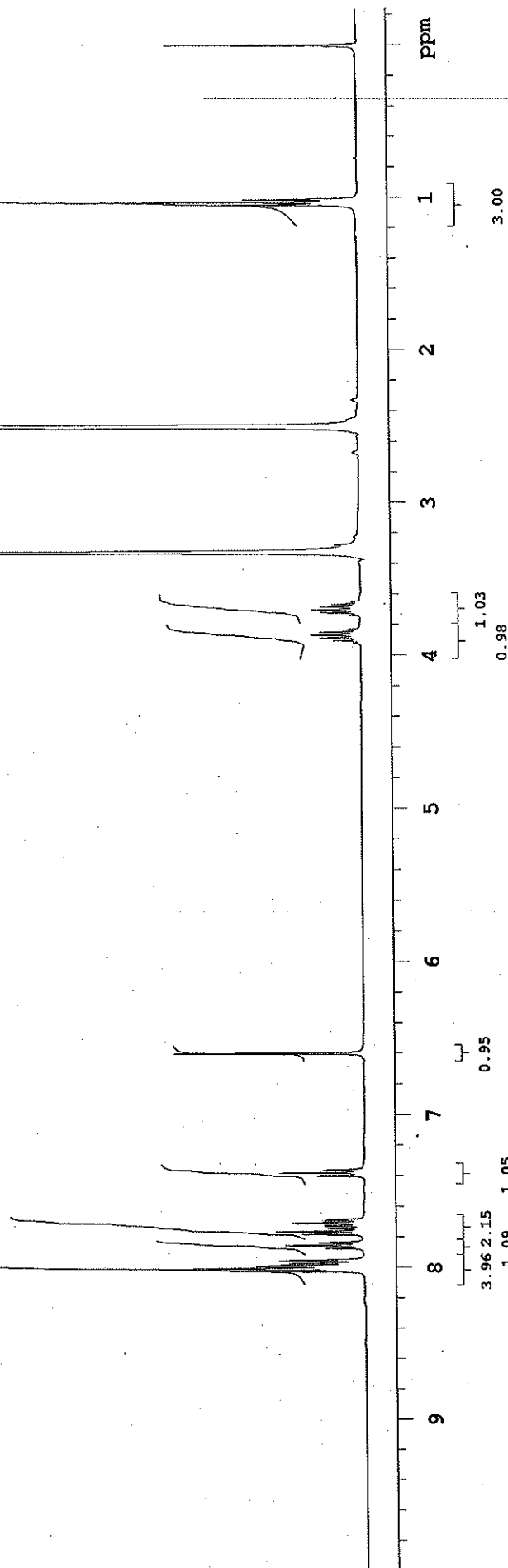
901-TDC-204_A558_QDR_32_20101115_01

FidFile: 901-TDC-204_A558_QDR_32_20101115_01

Pulse Sequence: PROTON (s2pul)

Solvent: dmsd

Data collected on: Nov 15 2010



Plotname: 901-TDC-204_A558_QDR_32_20101115_01_plot01

VARIAN

1714

TDC-204 A558-QDR-32 in DMSO

NMR-400

AR.No:ME1110/1252.

Analyst:Haribabu.R

Date: 17 th Nov 2010

Sample Name:

1252-TDC-204_A558-QDR-32

Data Collected on:

mercuryj-mercury400

Archive directory:

/home/sms50/routine/Nov2010/d_17

Sample directory:

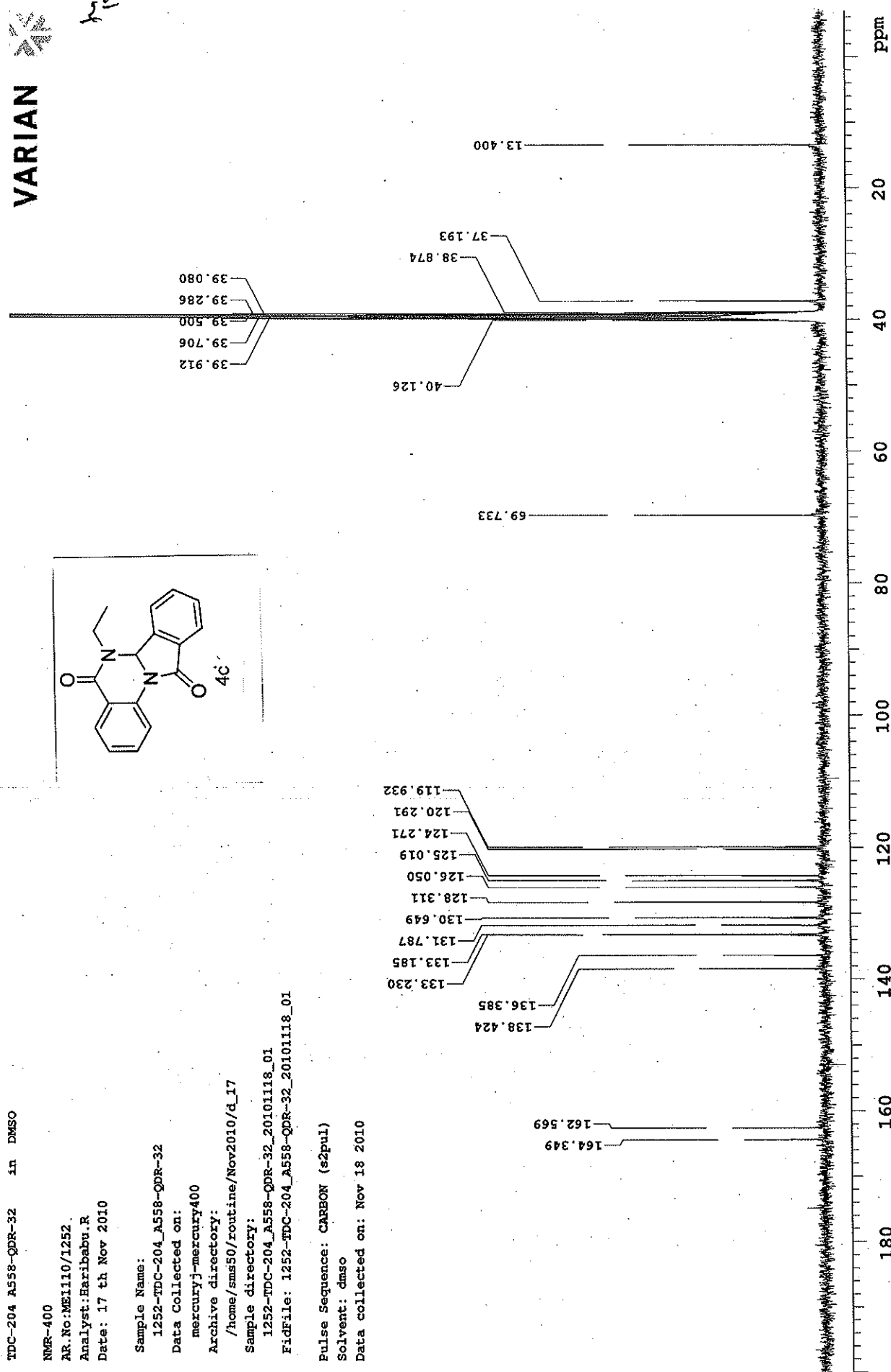
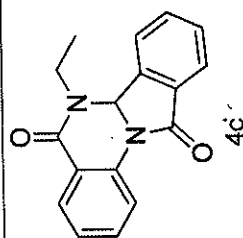
1252-TDC-204_A558-QDR-32_20101118_01

FidFile: 1252-TDC-204_A558-QDR-32_20101118_01

Pulse Sequence: CARBON (s2pul)

Solvent: dmsO

Data collected on: Nov 18 2010



Plotname: 1252-TDC-204_A558-QDR-32_20101118_01_Plot01

Elemental Composition Report

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.0, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

159 formula(e) evaluated with 1 results within limits (up to 4 best isotopic matches for each mass)

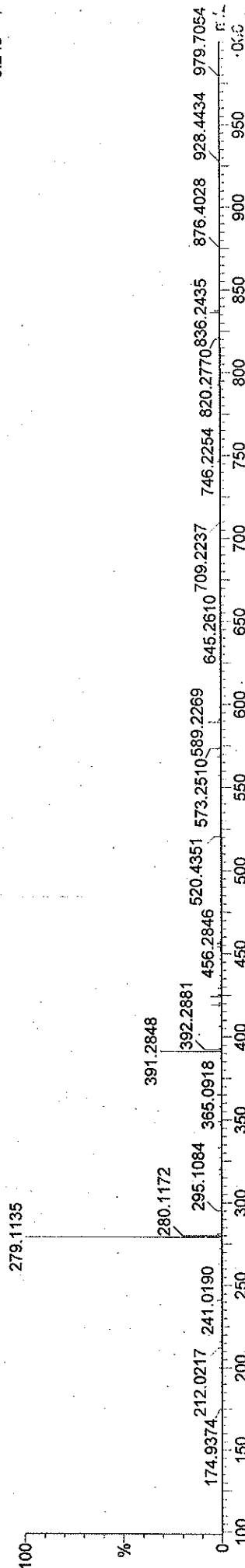
Elements Used:

C: 0-45 H: 0-55 N: 0-4 O: 0-4 Br: 0-1

A558-QDR-32

UT1110_74 17 (0.326) Cm (17:24-81:90)

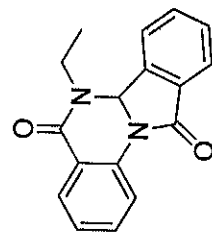
1: TOF MS
8.24e



Minimum:
Maximum:

5.0 5.0
-1.0 80.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Formula
279.1135	279.1134	0.1	0.4	11.5	0.1	C17 H15 N2 O2

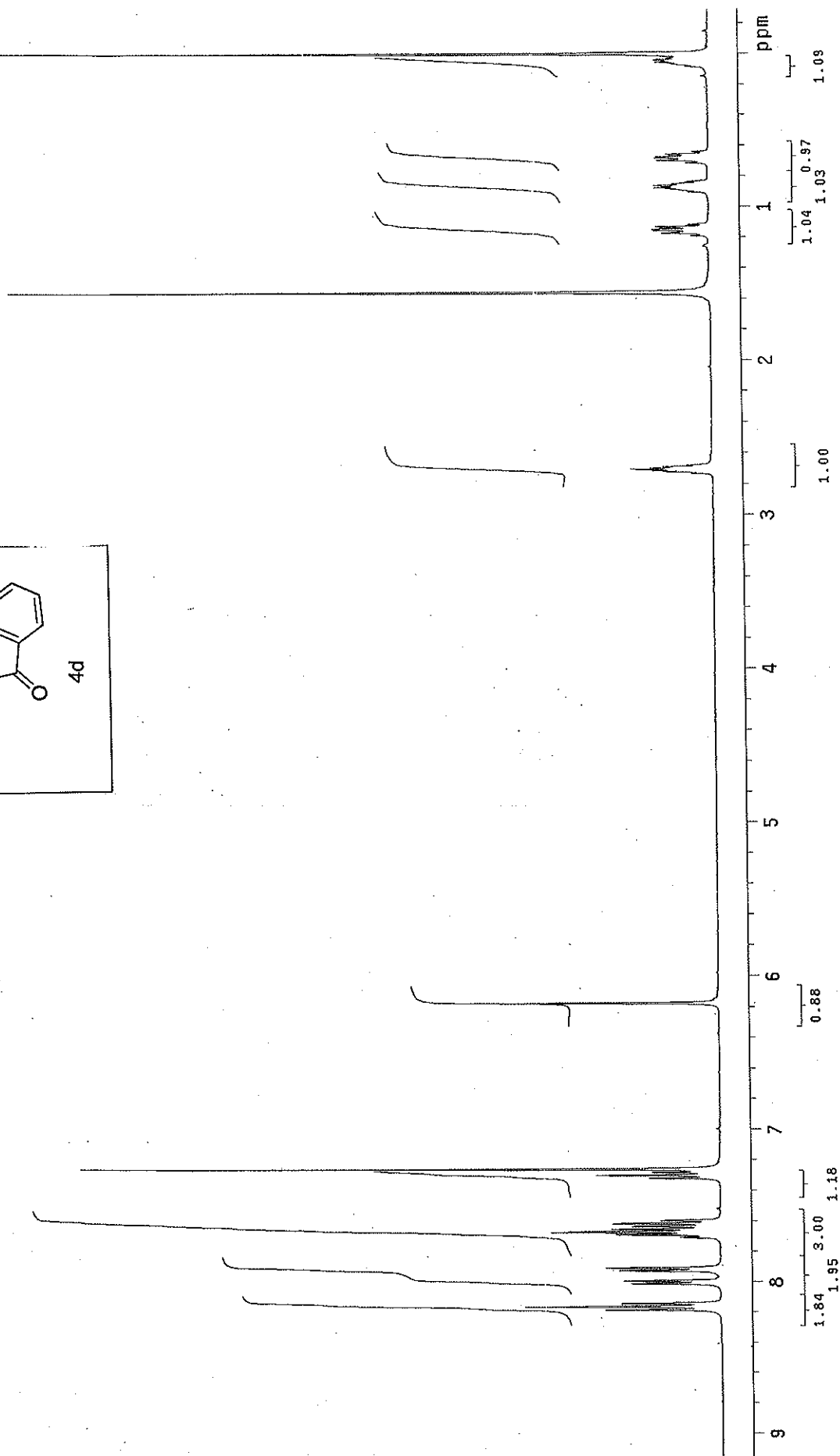
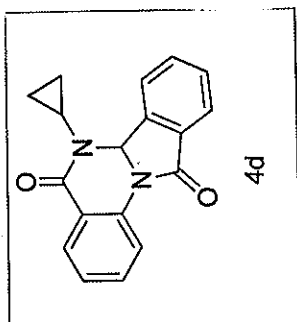


4c

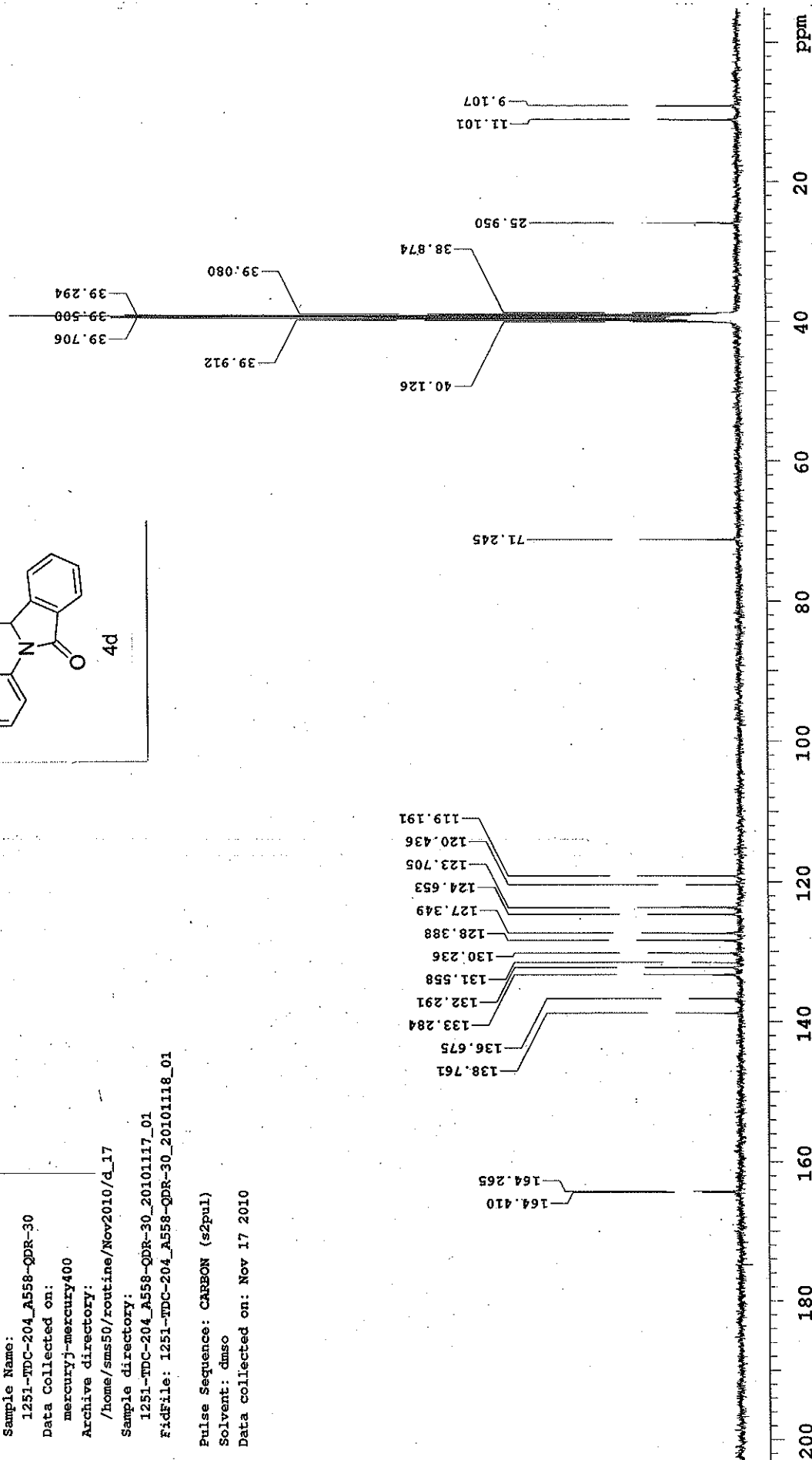
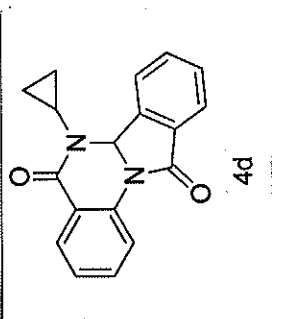
Cont

TDC -204 A558-QDR-30 in CDCl₃

NMR-400
AR.No:ME0111/199
Analyst:Haribabu.R
Date: 04 th Jan. 2011



VARIAN



Plotname: 1251-TDC-204_A558-QDR-30_20101118_01_plot01

TDC-204 A558-QDR-30 in DMSO

NMR-400

AR.No:ME1110/1251

Analyst:Haribabu.R

Date: 17 th Nov 2010

Sample Name:

1251-TDC-204_A558-QDR-30

Data Collected on:

mercury-j-mercury400

Archive directory:

/home/sms50/routine/Nov2010/d_17

Sample directory:

1251-TDC-204_A558-QDR-30_20101117_01

FidFile: 1251-TDC-204_A558-QDR-30_20101118_01

Pulse Sequence: CARBON (s2pul)

Solvent: dmsd

Data collected on: Nov 17 2010

Elemental Composition Report

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.0, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

168 formula(e) evaluated with 1 results within limits (up to 4 best isotopic matches for each mass)

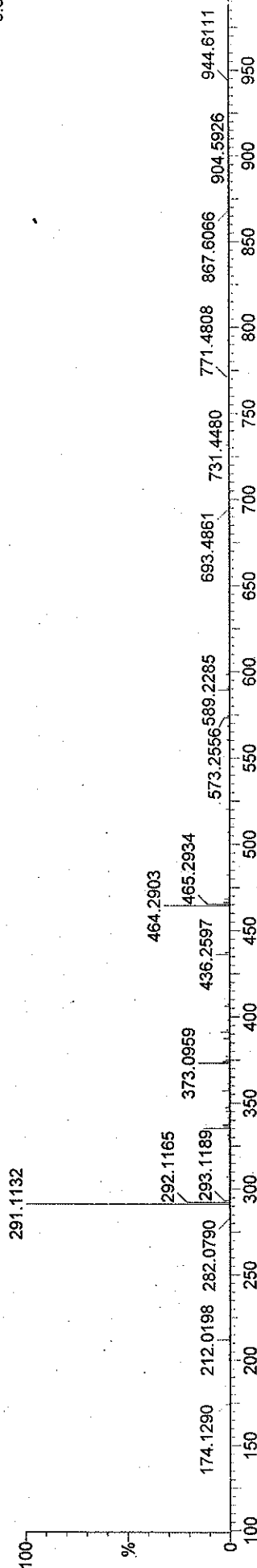
Elements Used:

C: 0-45 H: 0-55 N: 0-4 O: 0-4 Br: 0-1

A558-QDR-30

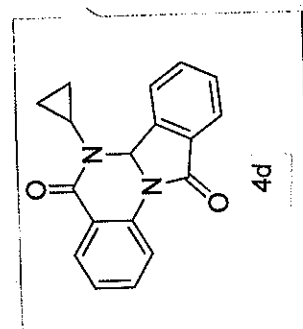
UT1110_72 20 (0.367) Cm (20:26-80:99)

1: TOF MS E⁺
6.67e+003



Minimum: -1.0
Maximum: 80.0

Mass	Calc. Mass	mDa	DBE	i-FIT	Formula
291.1132	291.1134	-0.2	12.5	0.1	C18 H15 N2 O2



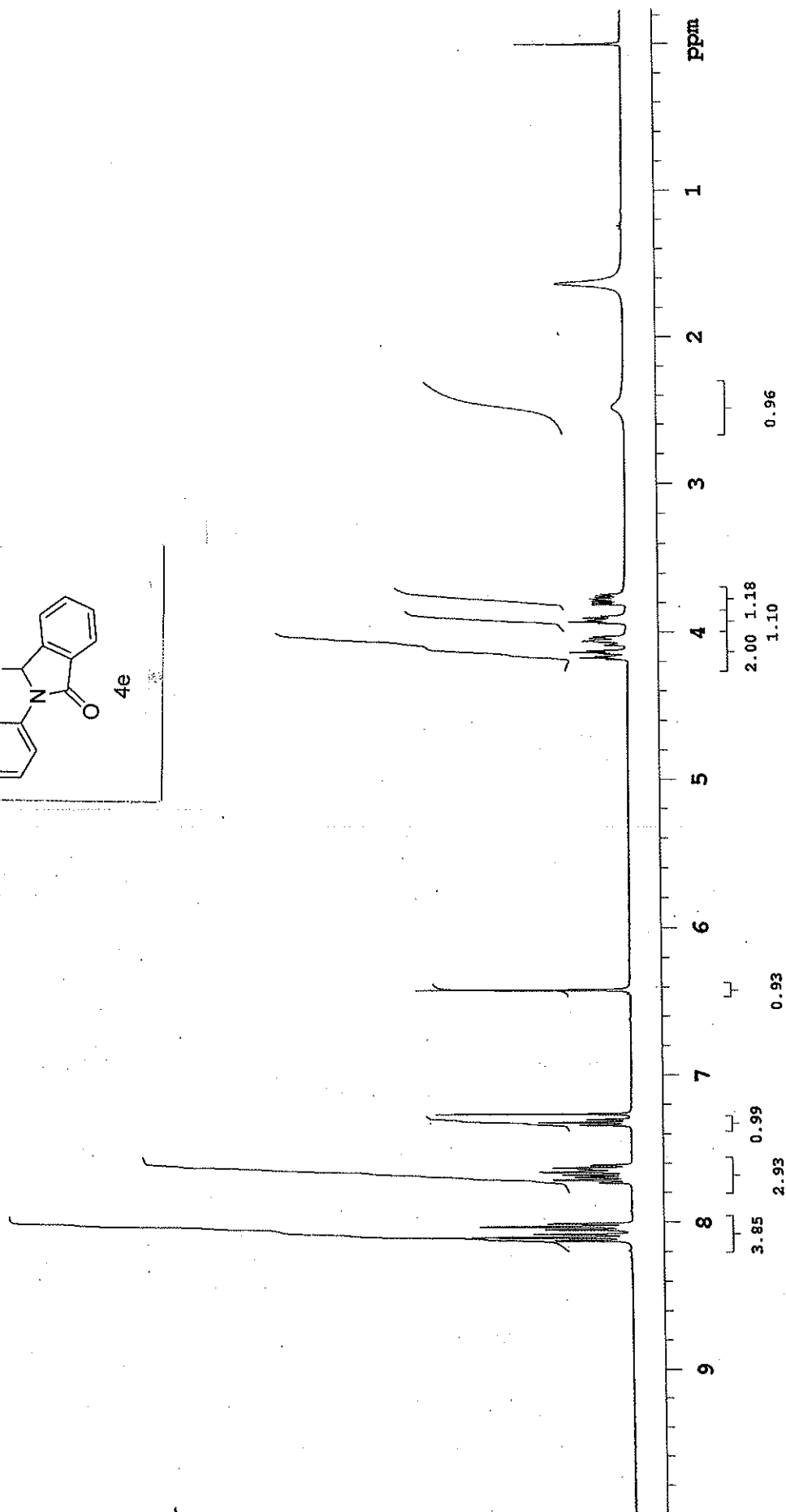
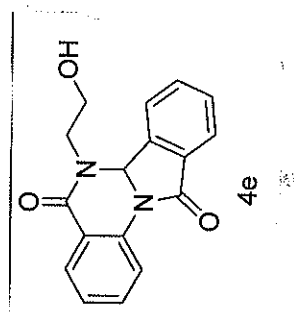
TDC-204 A558/QDR/28 in CDCl3

NMR-400

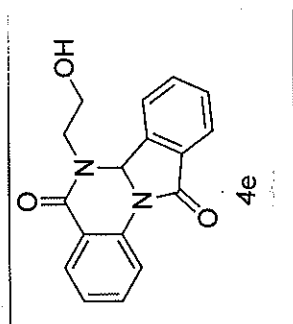
AR.No:ME1010/1409

Analyst:Haribabu.R

Date: 18 th Oct 2010



2116



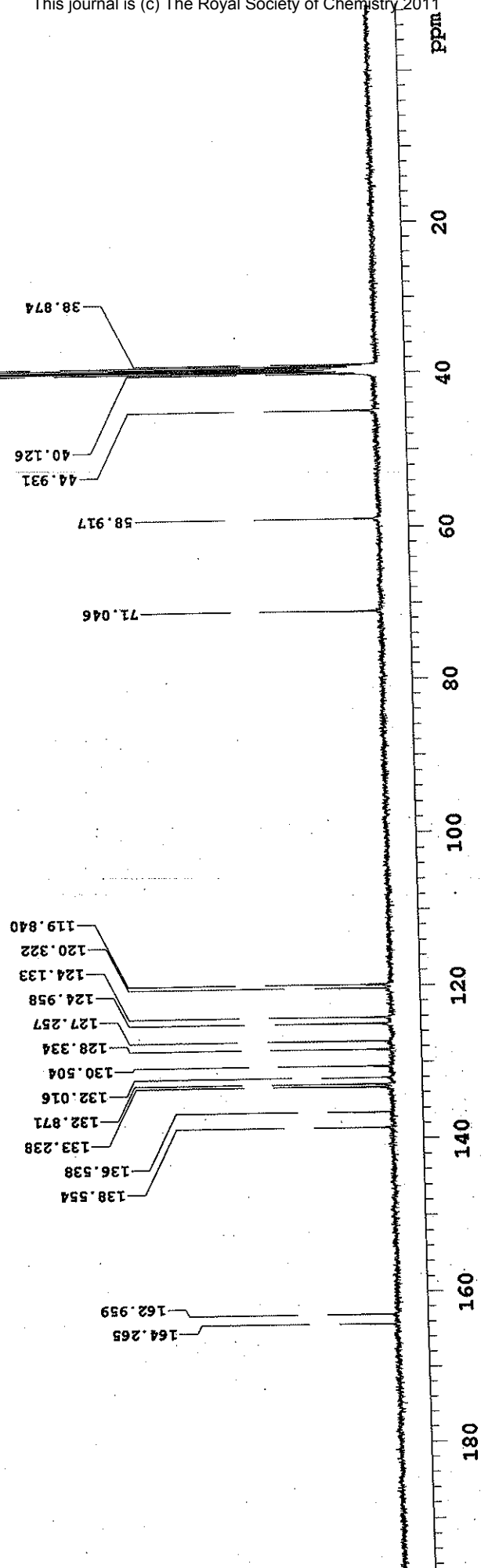
TDC-204 A558/QDR/28 in DMSO

NMR-400

AR.No:ME1010/1709

Analyst:Shruthi

Date: 20 th Oct 2010



Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.0, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

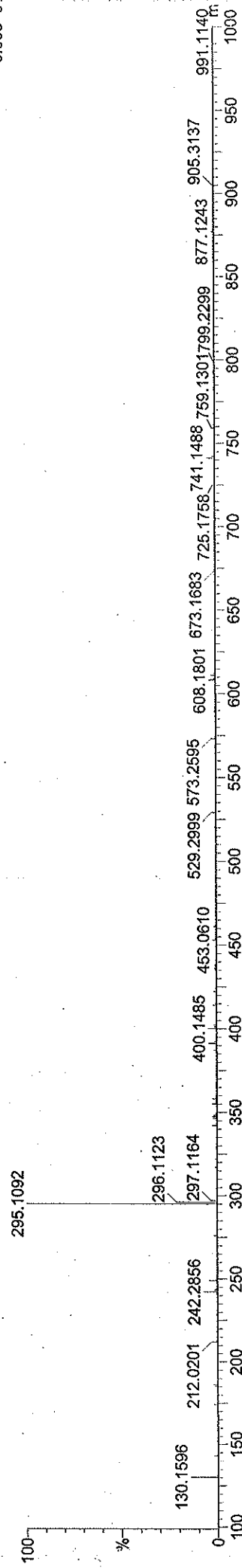
66 formula(e) evaluated with 1 results within limits (up to 4 closest results for each mass)

Elements Used:

C: 0-45 H: 0-70 N: 0-3 O: 0-3

A558/QDR128

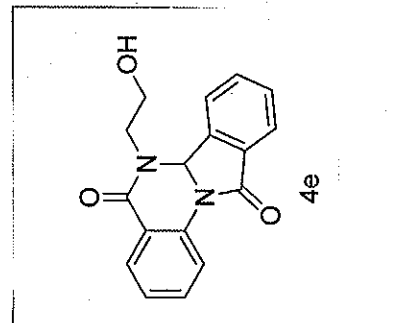
UT1010_194 16 (0.293) Cm (16:22-77:88)

1: TOF MS ES
6.06e+0.1

Minimum:

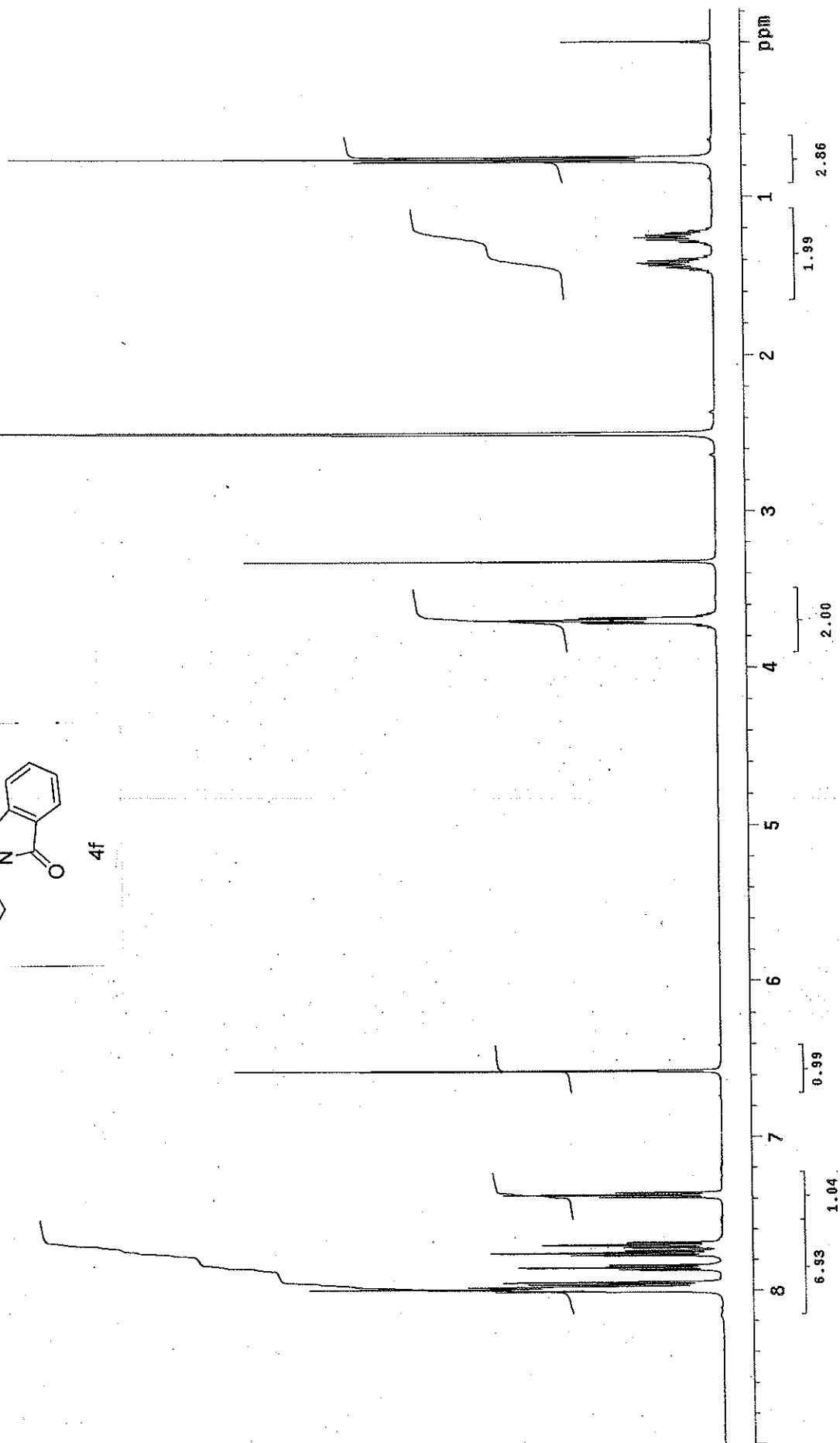
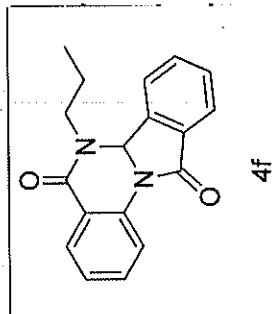
Maximum:

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Formula
295.1092	295.1083	0.9	3.0	11.5	2.1	C17 H15 N2 O3



A558/QDR/031 in DMSO
TDC-204

NMR: 500MHz
AR_No: IN1010/242
Date: 25th Oct. 2010
Analyst: Haribabu. R



VARIAN



7/5/2010

TDC-204 A558-QDR-31 in DMSO

NMR-400

AR.No:ME1110/1656

Analyst:Haribabu.R

Date: 22 th Nov 2010

Sample Name:

1656-TDC-204_A558-QDR-31

Data Collected on:

mercuryj-mercury400

Archive directory:

/home/sms50/routine/Nov2010/d_22

Sample directory:

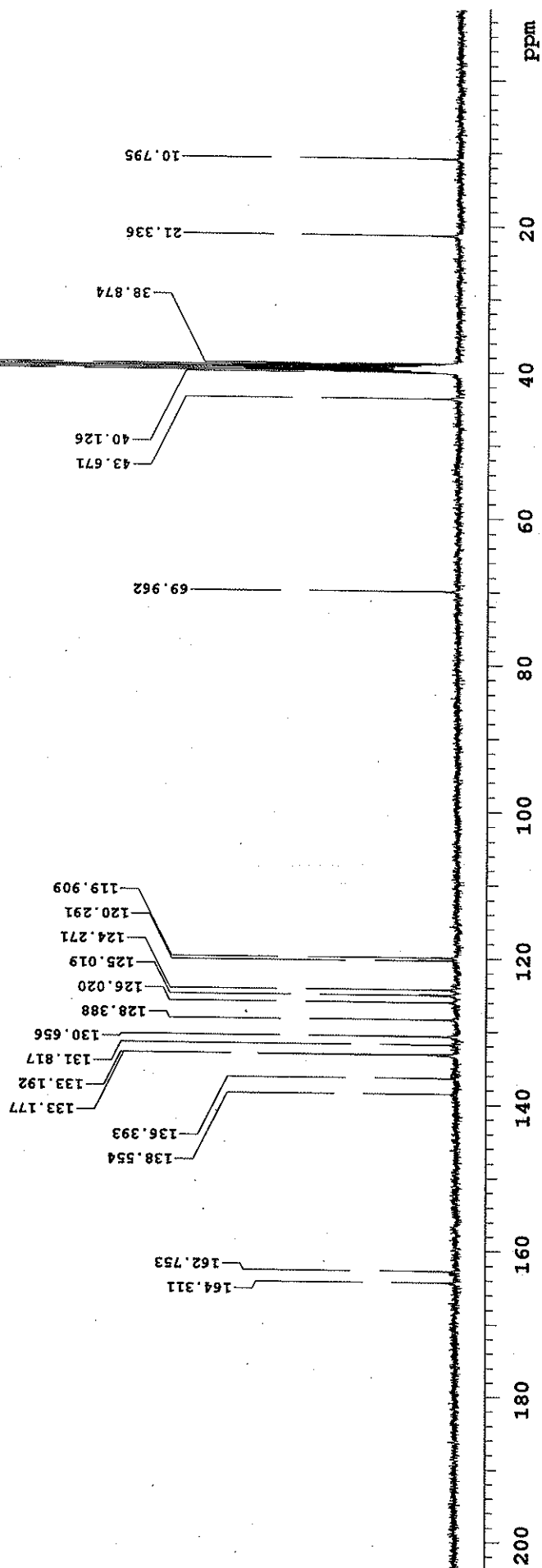
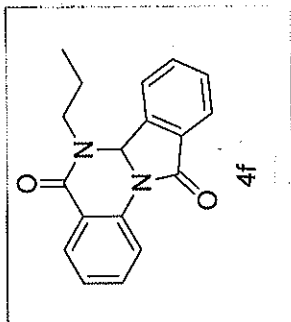
1656-TDC-204_A558-QDR-31_20101122_01

FidFile: 1656-TDC-204_A558-QDR-31_20101123_01

Pulse Sequence: CARBON (s2pul)

Solvent: dmsc

Data collected on: Nov 22 2010



Plotname: 1656-TDC-204_A558-QDR-31_20101123_01_plot01

Elemental Composition Report

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.0, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

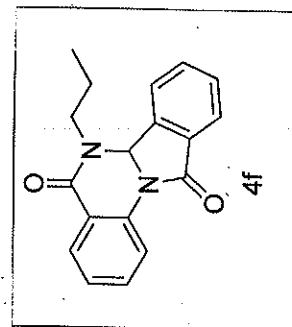
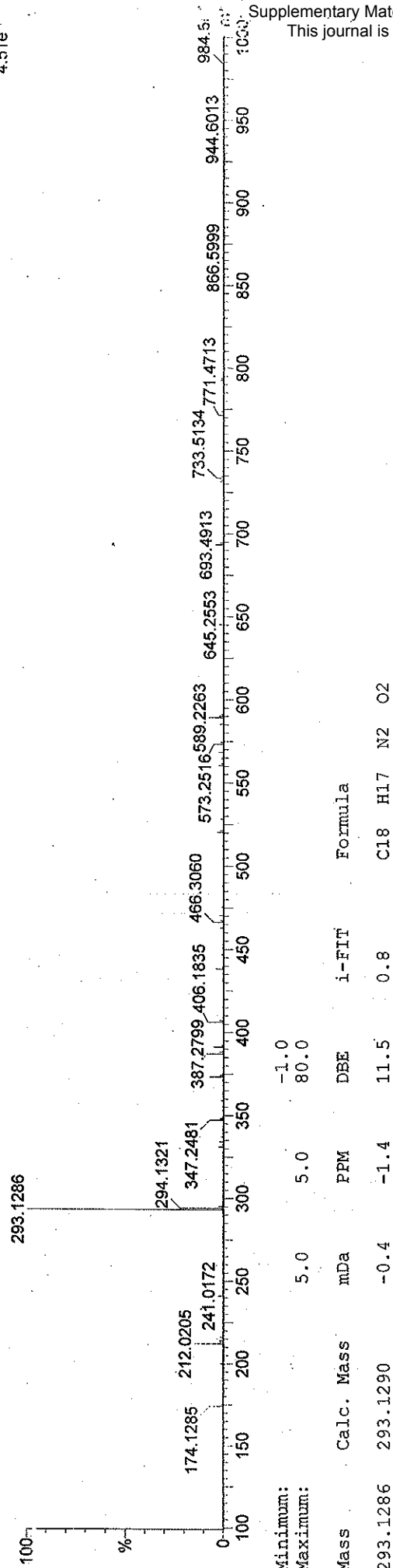
166 formula(e) evaluated with 1 results within limits (up to 4 best isotopic matches for each mass)

Elements Used:

C: 0.45 H: 0.55 N: 0.4 O: 0.4 Br: 0.1

A558-QDR-31

UT1110_73 20 (0.367) Cm (20:27-73:81)



A558/QDR/18 1n DMSO

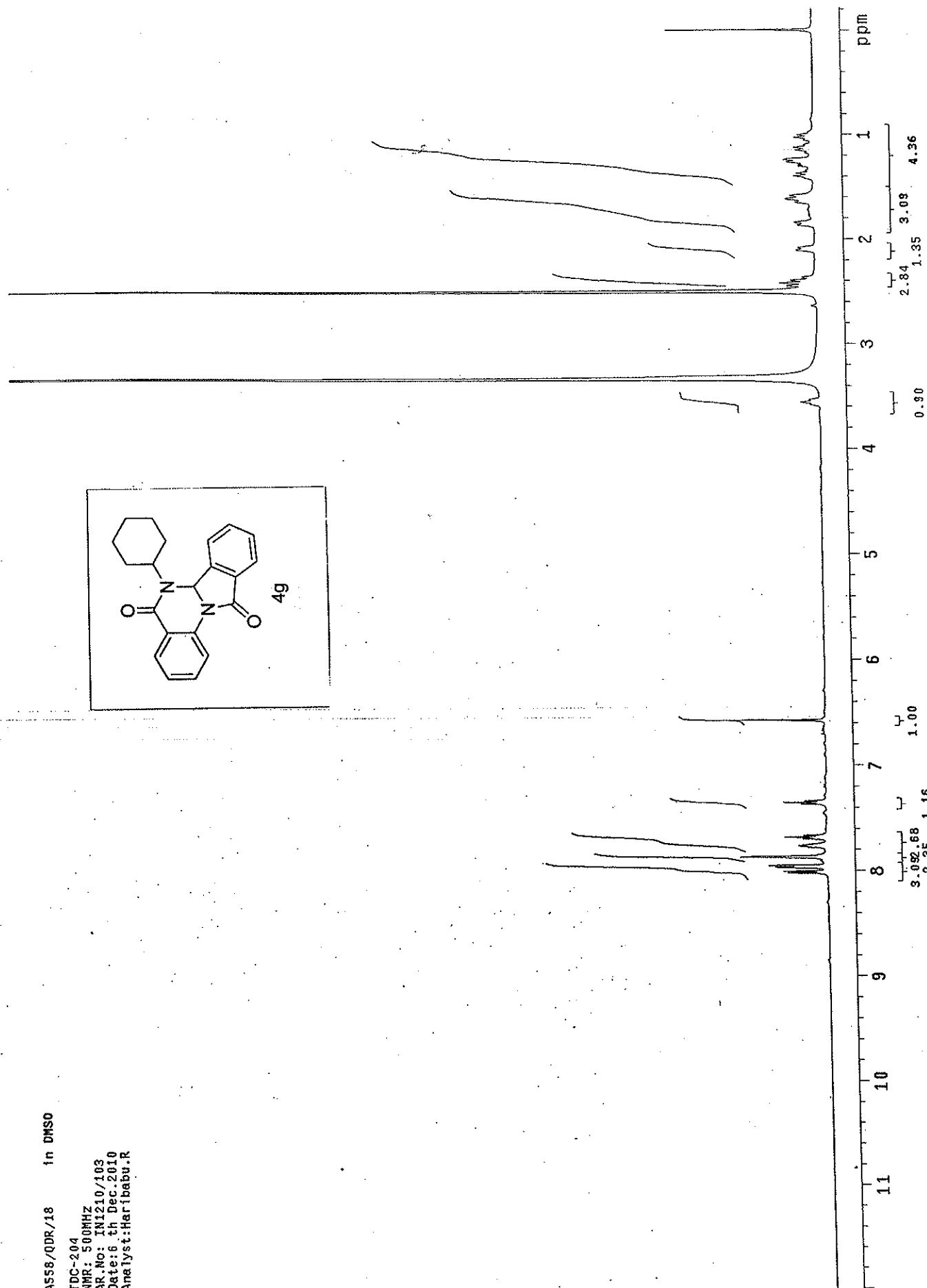
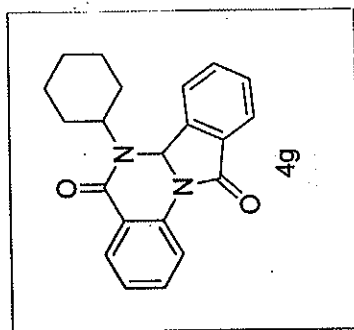
TDC-204

NMR: 500MHZ

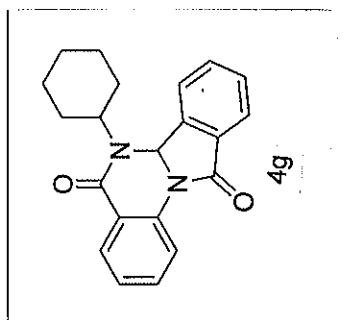
AR.No: IN1210/103

Date: 6 th Dec. 2010

Analyst: Haribabu.R



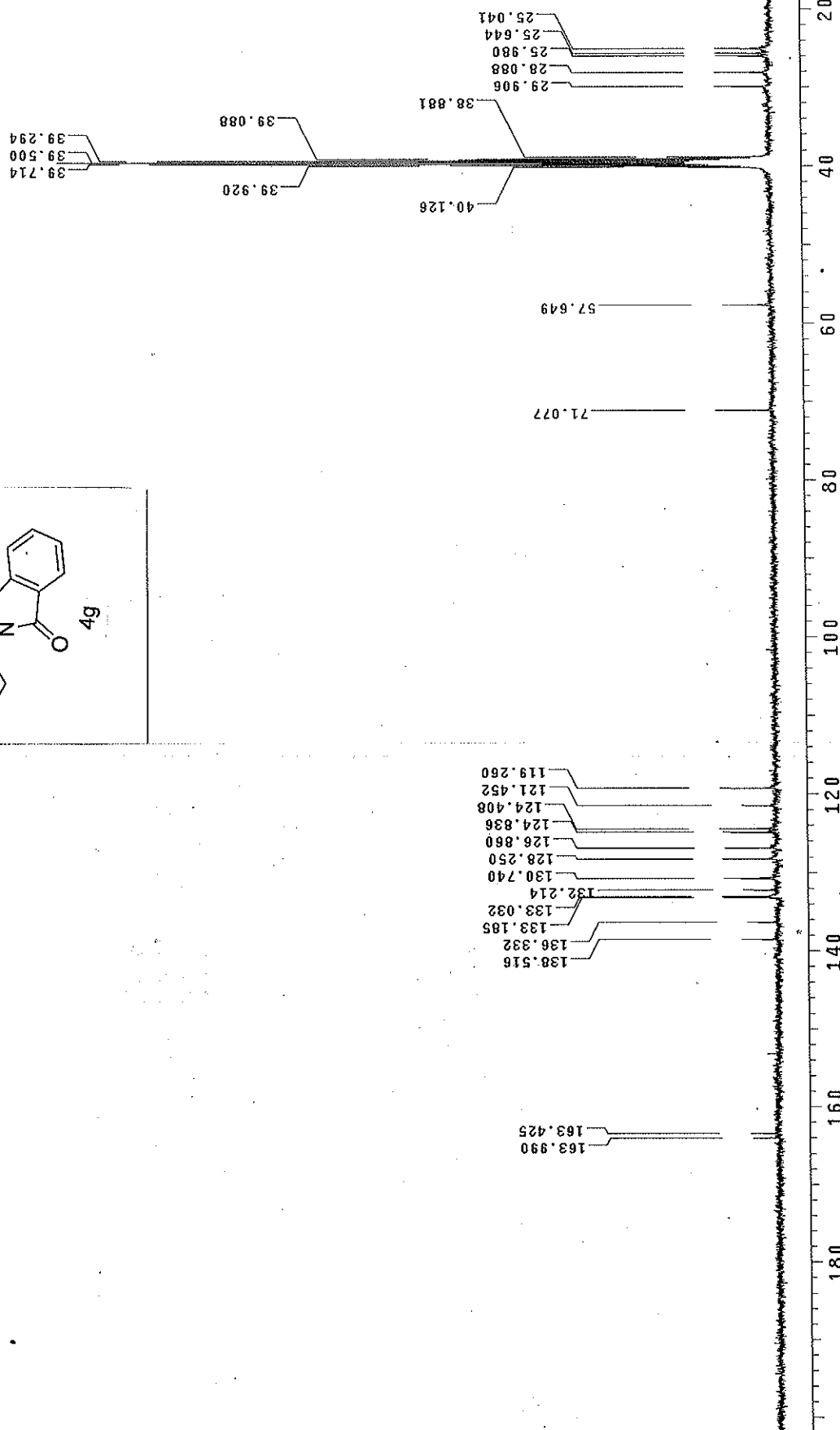
21/10/11



in DMSO

TDC-204 A558-QDR-18

NMR-400
AR.No:ME1210/1009
Analyst:Haribabu.R
Date: 10 th Dec. 2010



Elemental Composition Report

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.0, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

181 formula(e) evaluated with 1 results within limits (up to 4 best isotopic matches for each mass)

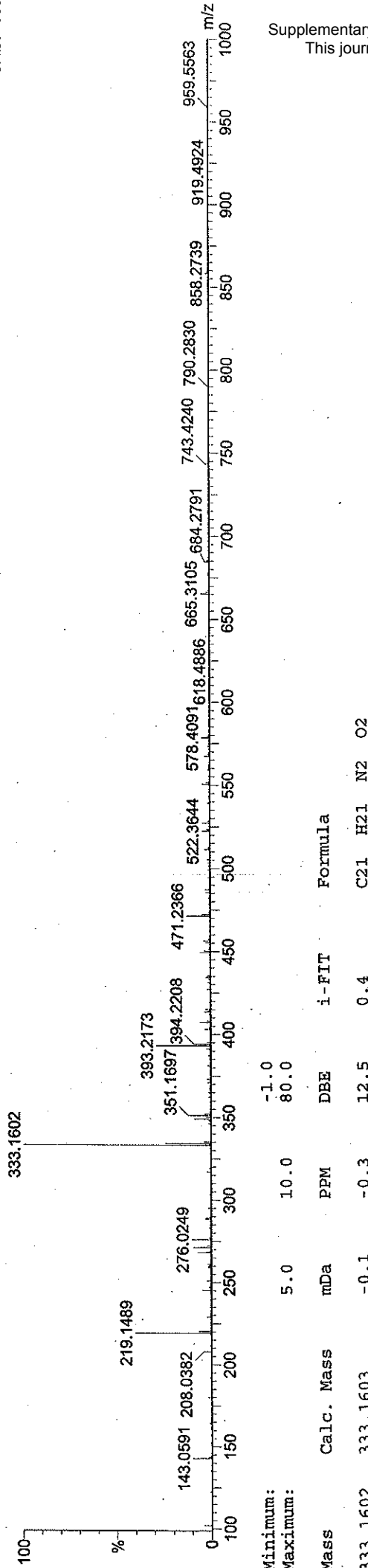
Elements Used:

C: 0-40 H: 0-55 N: 0-4 O: 0-8

A558/QDR/18

UT1210_148.16 (0.292) Cm (16:18-61:70)

1: TOF MS ES+
5.42e+003



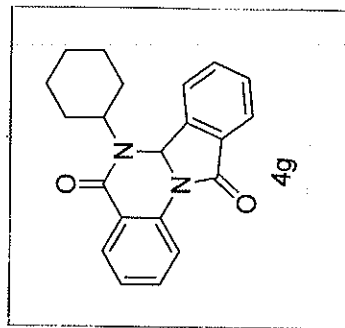
Minimum:
Maximum:

5.0 10.0
mDa PPM

-1.0
80.0

DBE i-FIT Formula

333.1602 333.1603 -0.1 -0.3 12.5 0.4 C21 H21 N2 O2



25/10

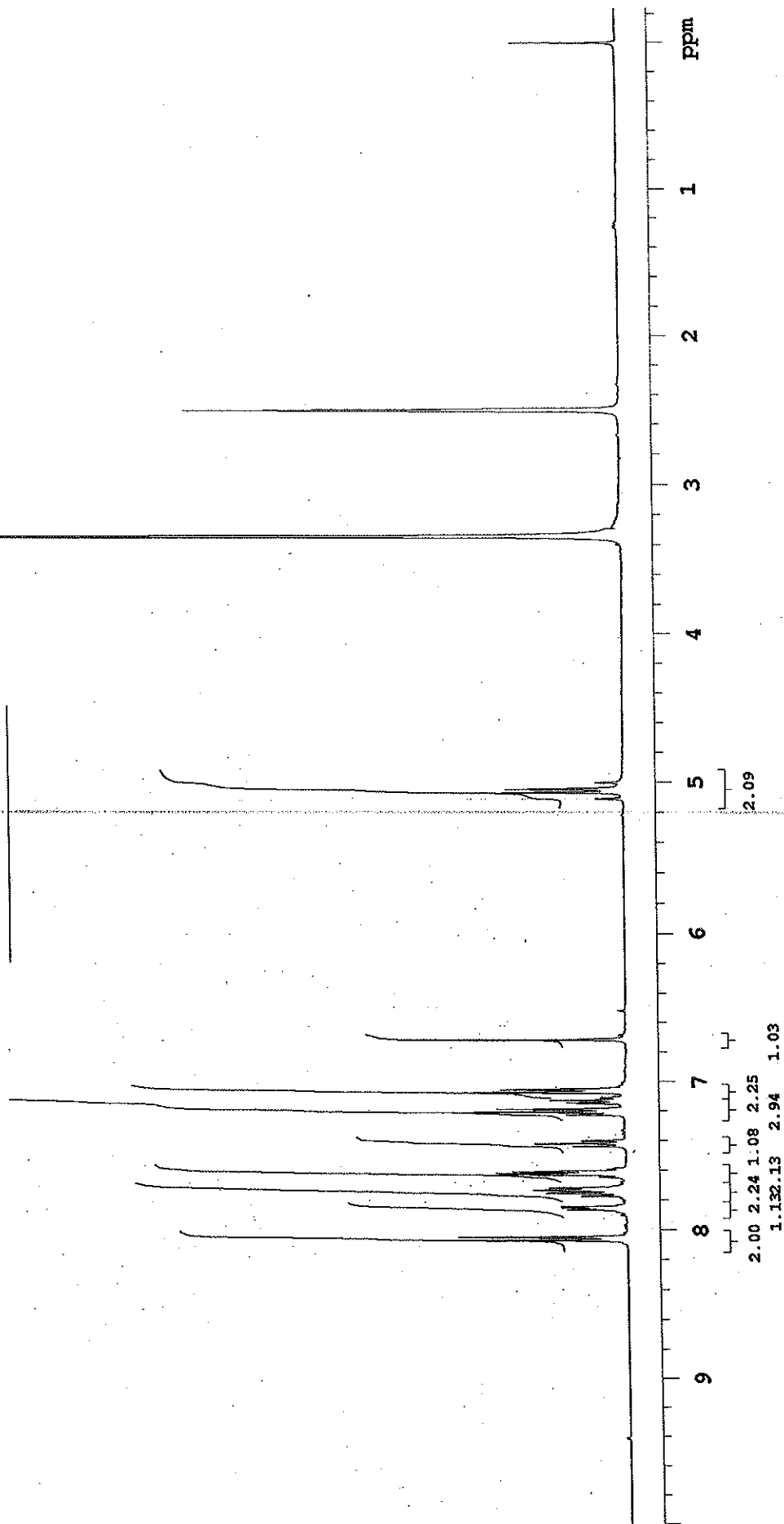
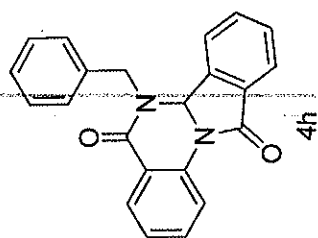
TDC-204 A558/QDR/29-2 in DMSO

NMR-400

AR.No:ME1010/2023

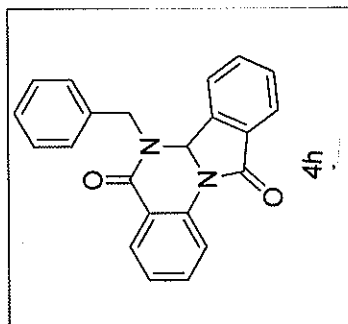
Analyst:Haribabu.R

Date: 25 th Oct 2010



VARIAN

20.11.10



TDC-204 A558-QDR-29 in DMSO

NMR-400

AR.No:ME1110/1655

Analyst:Haribabu.R

Date: 22 th Nov 2010

Sample Name:

1655-TDC-204_A558-QDR-29

Data Collected on:

mercuryj-mercury400

Archive directory:

/home/sms50/routine/Nov2010/d_22

Sample directory:

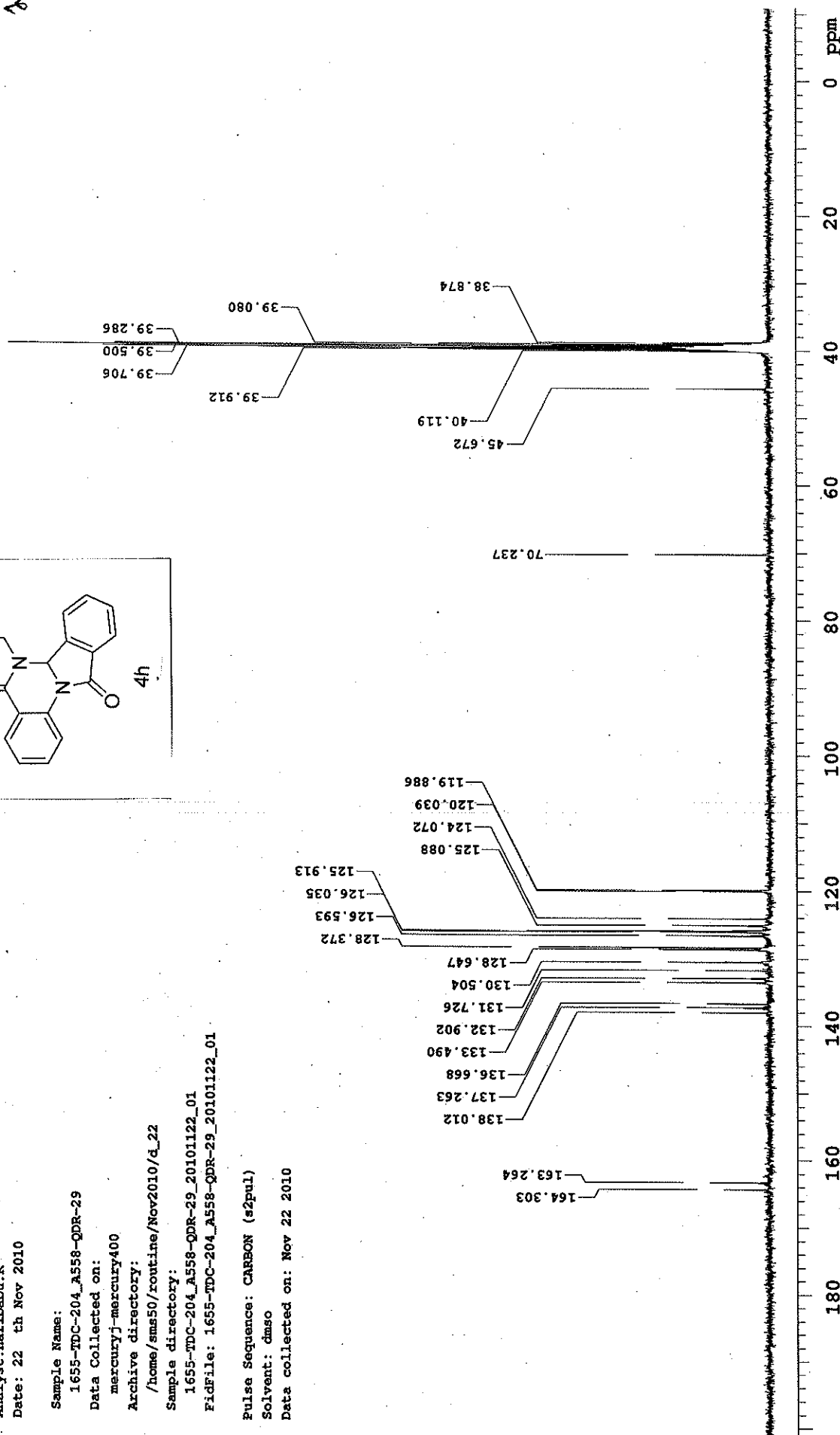
1655-TDC-204_A558-QDR-29_20101122_01

FidFile: 1655-TDC-204_A558-QDR-29_20101122_01

Pulse Sequence: CARBON (s2pul)

Solvent: dmsc

Data collected on: Nov 22 2010



Plotname: 1655-TDC-204_A558-QDR-29_20101122_01_plot01

Elemental Composition Report

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.0, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

196 formula(e) evaluated with 1 results within limits (up to 4 best isotopic matches for each mass)

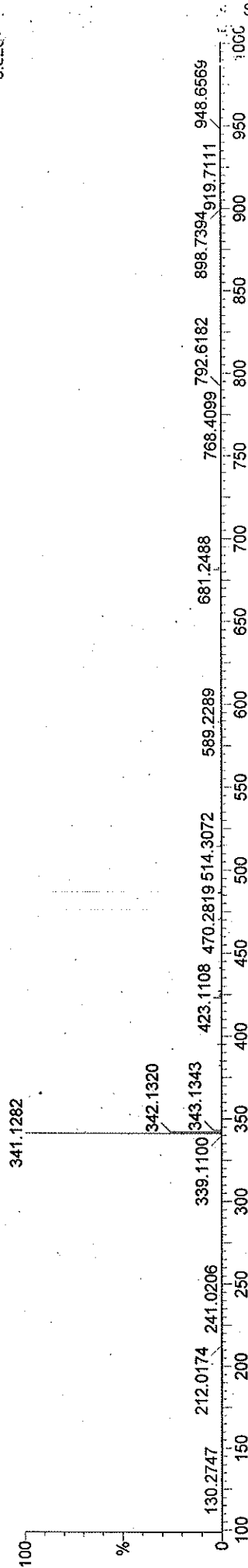
Elements Used:

C: 0-45 H: 0-55 N: 0-4 O: 0-4 Br: 0-1

A558-QDR-29

UT1110_76 25 (0.472) Cm (25:33-78:89)

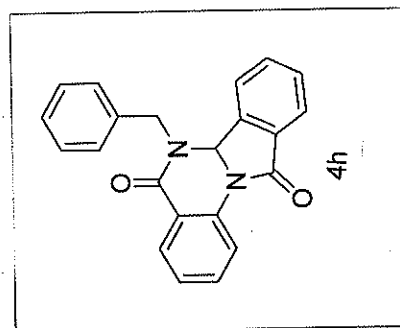
1: TOF MS
8.22e4



Minimum:
Maximum:

-1.0
80.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Formula
341.1282	341.1290	-0.8	-2.3	15.5	0.9	C22 H17 N2 O2



VARIAN

TDC-204 A558/QDR/33 in DMSO

NMR-400

AR.No:ME1110/900

Analyst:Haribabu.R

Date: 15 th Nov 2010

Sample Name:

900-TDC-204_A558_QDR_33

Data Collected on:

mercuryj-mercury400

Archive directory:

/home/sms50/routine/Nov2010/d_15

Sample directory:

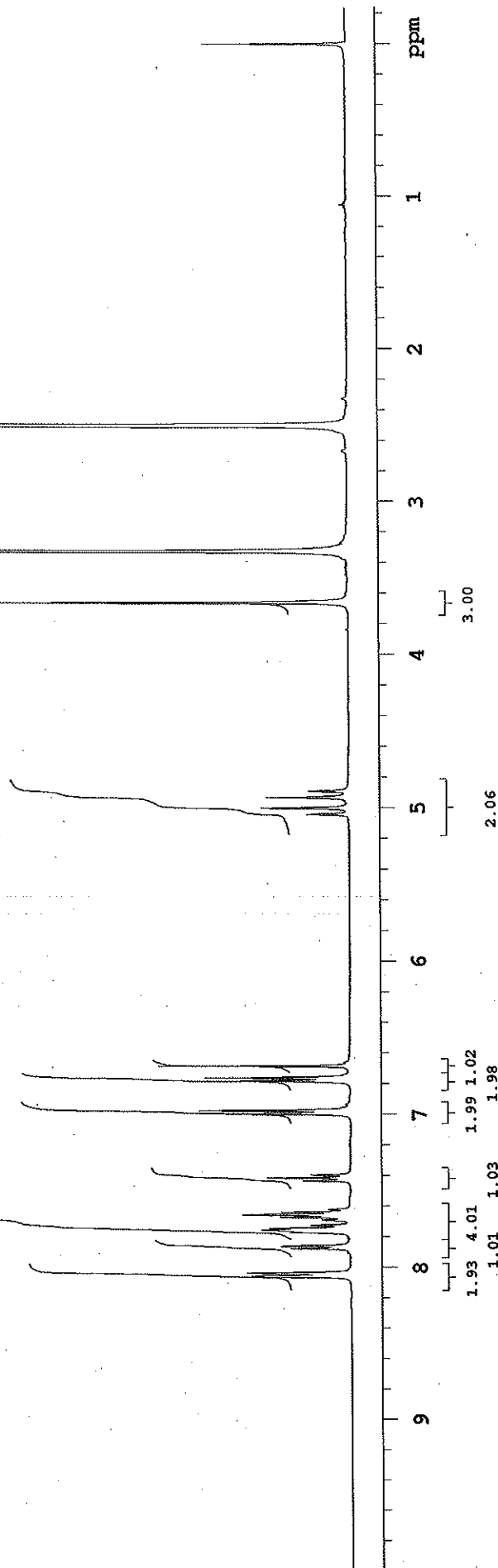
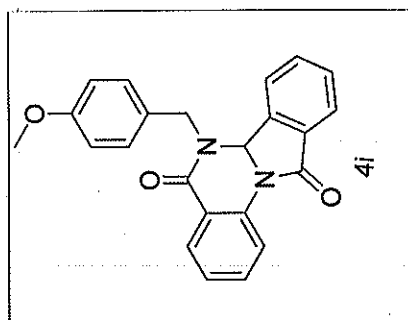
900-TDC-204_A558_QDR_33_20101115_01

FidFile: 900-TDC-204_A558_QDR_33_20101115_01

Pulse Sequence: PROTON (s2pul)

Solvent: dmsc

Data collected on: Nov 15 2010



Plotname: 900-TDC-204_A558_QDR_33_20101115_01_plot01

VARIAN

TDC-204 A558-QDR-33 in DMSO

NMR-400

AR.No:ME1110/1657

Analyst:Haribabu.R

Date: 22 th Nov 2010

Sample Name:

1657-TDC-204_A558-QDR-33

Data Collected on:

mercuryj-mercury400

Archive directory:

/home/sms50/routine/Nov2010/d_22

Sample directory:

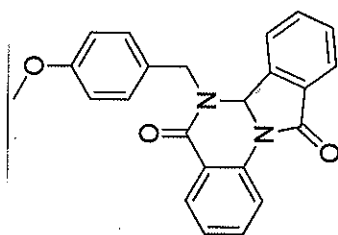
1657-TDC-204_A558-QDR-33_20101123_01

FidFile: 1657-TDC-204_A558-QDR-33_20101123_01

Pulse Sequence: CARBON (s2pul)

Solvent: dmsO

Data collected on: Nov 23 2010



39.920
39.714
39.500
39.294
39.088

54.930
45.107
40.126
38.874

70.206

128.640
127.303
126.111
125.073
124.103
120.092
119.848
113.814

138.088
136.637
133.452
132.933
131.749
130.534
128.968

164.280
163.234
157.925

180 160 140 120 100 80 60 40 20 ppm

Plotname: 1657-TDC-204_A558-QDR-33_20101123_01_plot01

Elemental Composition Report

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.0, max = 80.0
 Element prediction: Off
 Number of isotope peaks used for i-FIT = 3

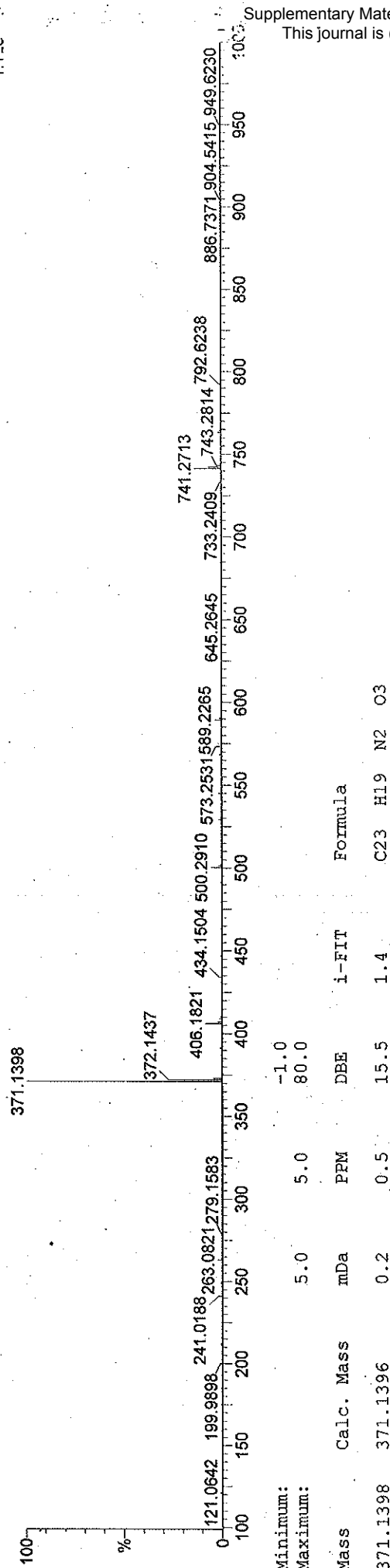
Monoisotopic Mass, Even Electron Ions
 213 formula(e) evaluated with 1 results within limits (up to 4 best isotopic matches for each mass)
 Elements Used:

C: 0-45 H: 0-55 N: 0-4 O: 0-4 Br: 0-1

A558-QDR-33

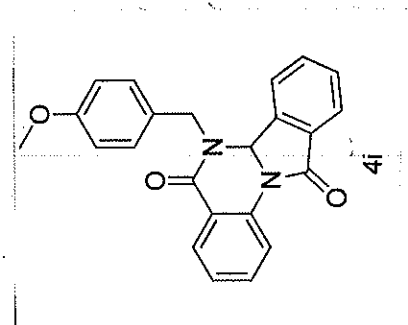
UT1110_75 17 (0.325) Cm (17:23-77:88)

1: TOF MS E
 1.12e4



Minimum:
 Maximum:

Mass Calc. Mass mDa PPM DBE i-FIT Formula
 371.1398 371.1396 0.2 0.5 15.5 1.4 C23 H19 N2 O3



TDC-204 A558-QDR-35 in DMSO

NMR-400

AR.No:ME1110/1197

Analyst:Haribabu.R

Date: 17 th Nov 2010

Sample Name:

1197-TDC-204_A558-QDR-35

Data Collected on:

mercuryj-mercury400

Archive directory:

/home/sms50/routine/Nov2010/d_17

Sample directory:

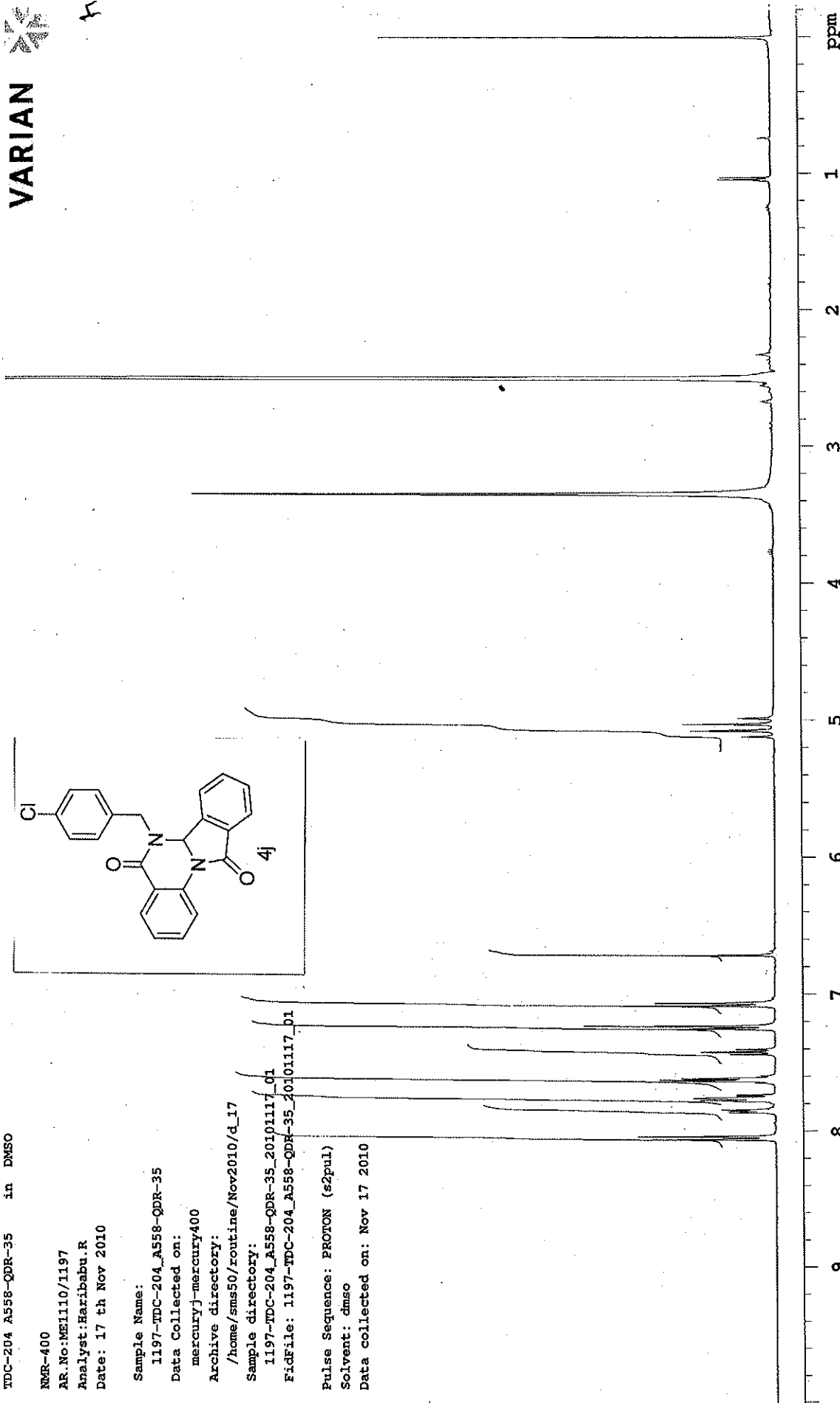
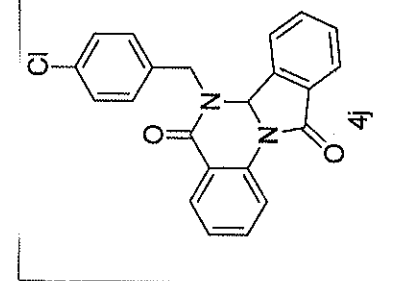
1197-TDC-204_A558-QDR-35_20101117_01

FidFile: 1197-TDC-204_A558-QDR-35_20101117_01

Pulse Sequence: PROTON (s2pul)

Solvent: dmsc

Data collected on: Nov 17 2010

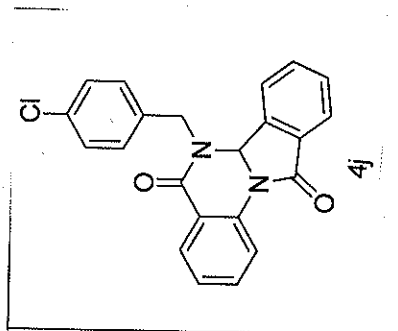


Plotname: 1197-TDC-204_A558-QDR-35_20101117_01_plot01

VARIAN

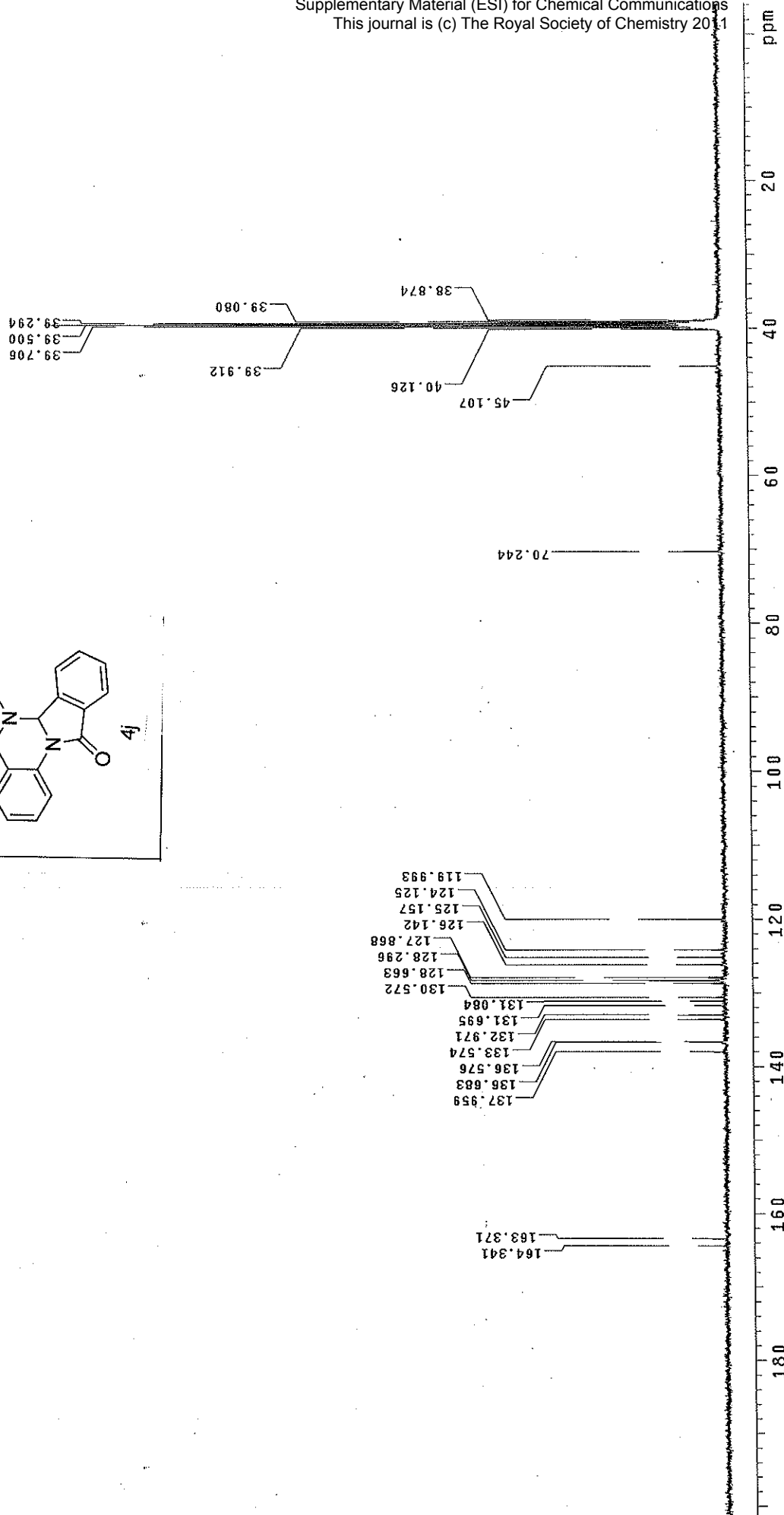
1197/10

20101110



TDC-204 A558-QDR-35 in DMSO

NMR-400
AR.NO:ME1210/1011
Analyst:Haribabu.R
Date: 10 th Dec. 2010



Elemental Composition Report

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.0, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

376 formula(e) evaluated with 4 results within limits (up to 4 best isotopic matches for each mass)

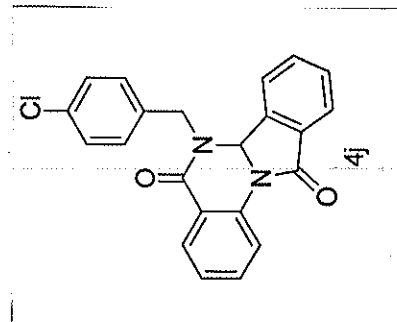
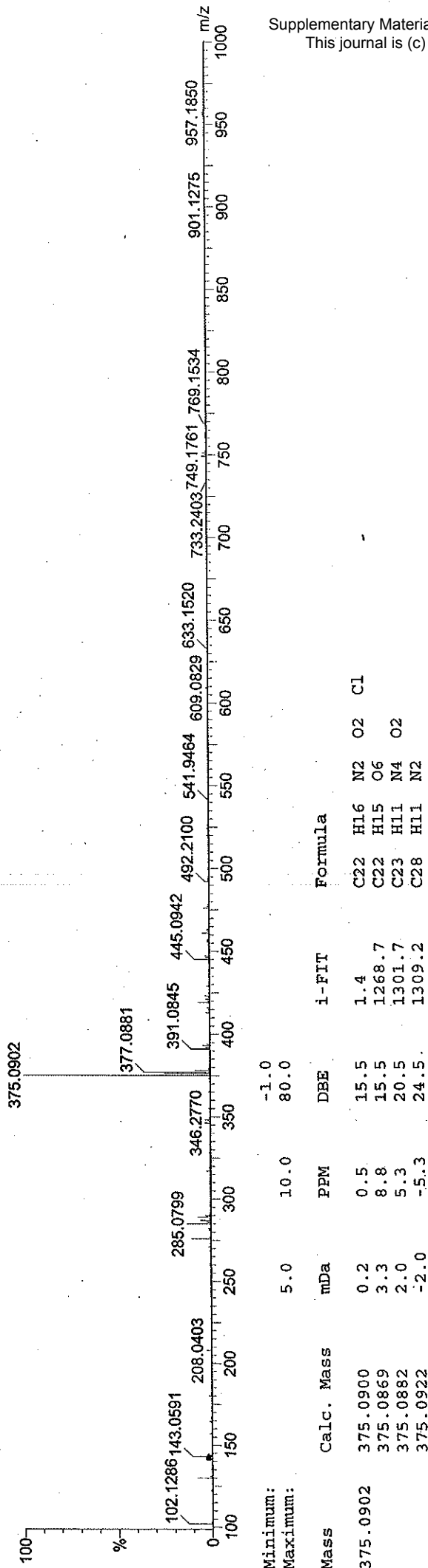
Elements Used:

C: 0-40 H: 0-55 N: 0-4 O: 0-8 Cl: 0-1

A558/QDR/35

UT1210_150 17 (0.325) Cm (17:20-64:70)

1: TOF MS ES+
9.30e+003



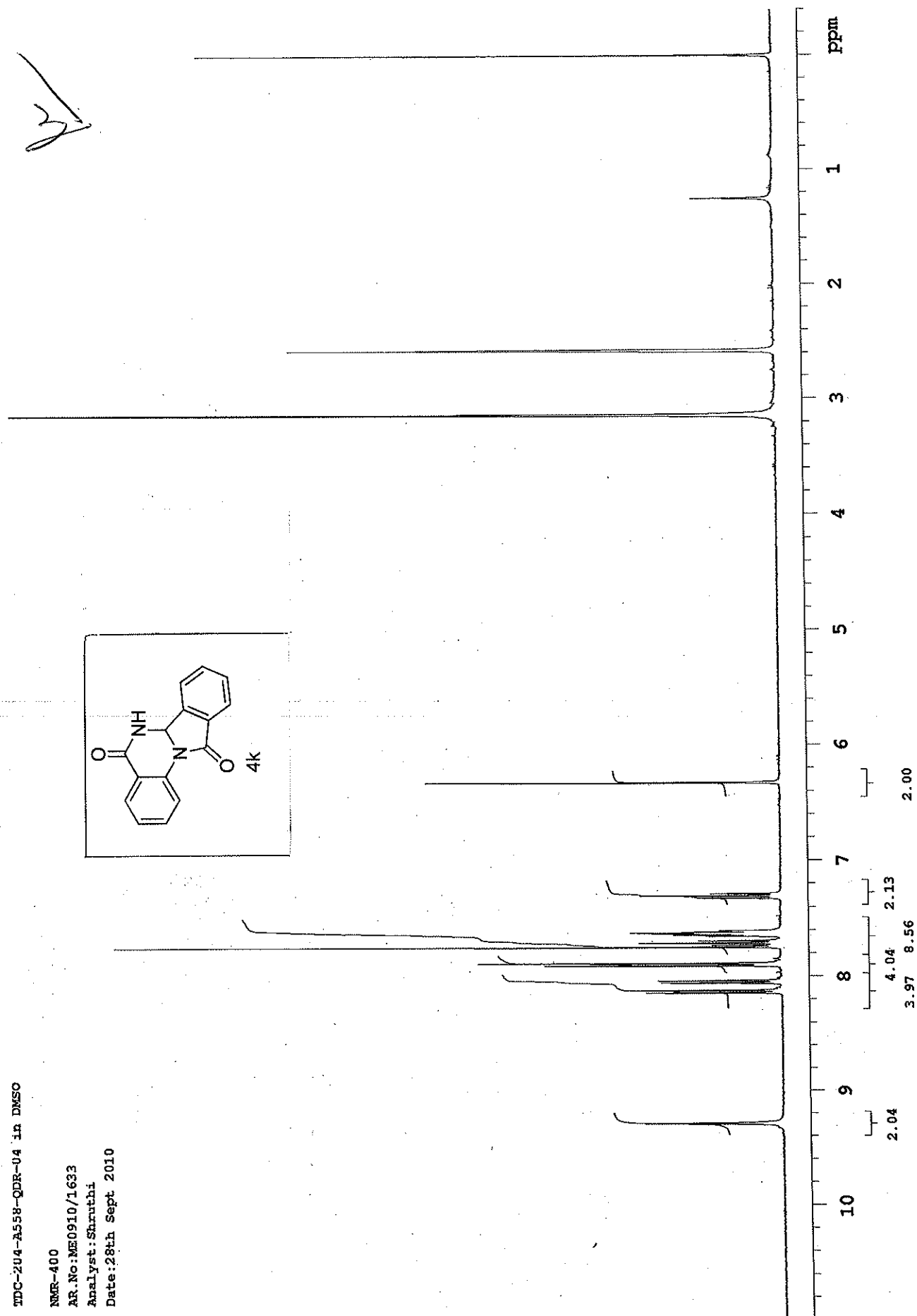
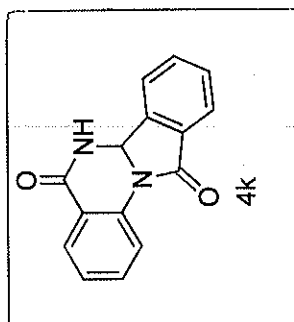
TDC-204-A558-QDR-04 in DMSO

NMR-400

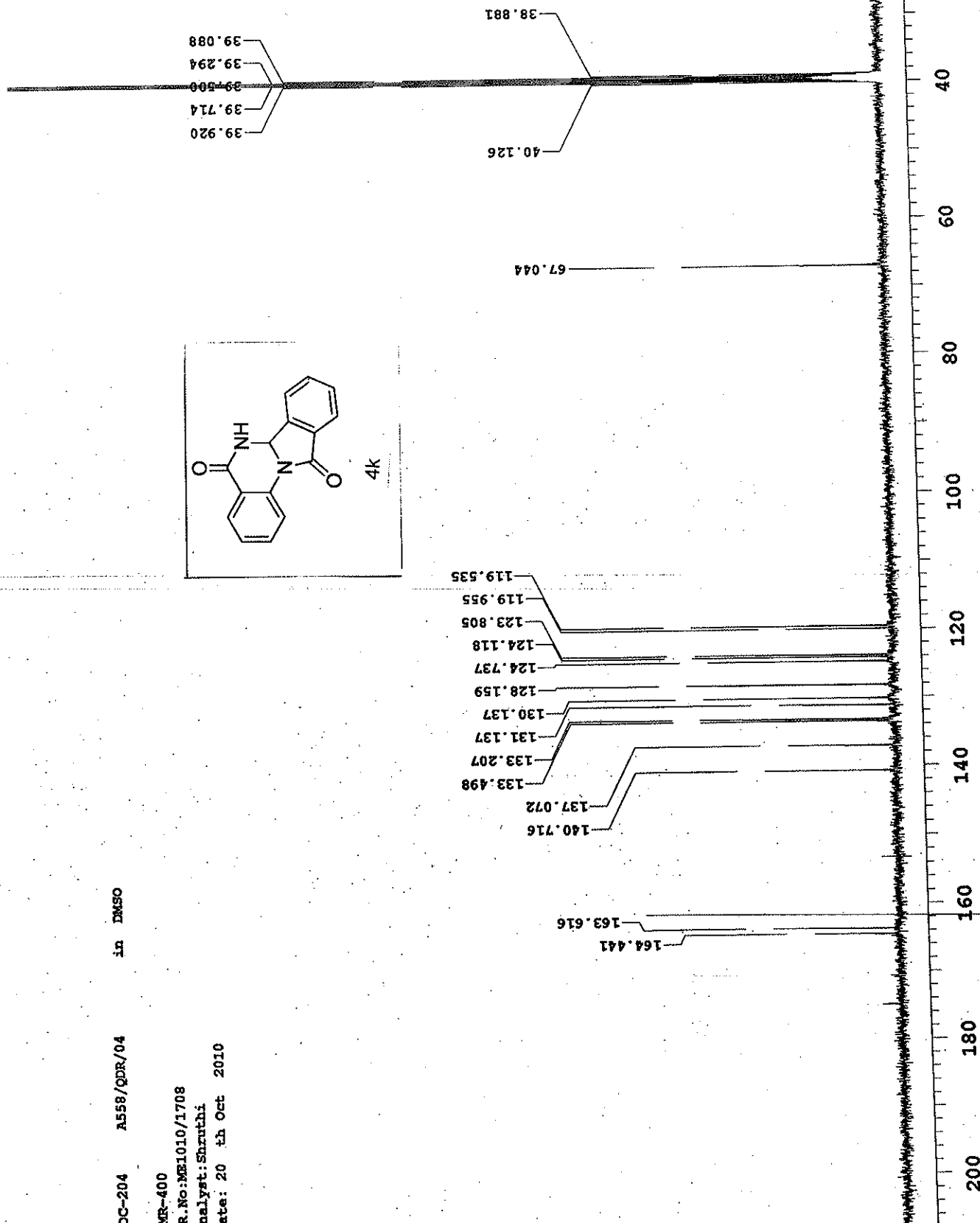
AR.No:ME0910/1633

Analyst: Shruthi

Date: 28th Sept 2010



MS
24/10



Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.0, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

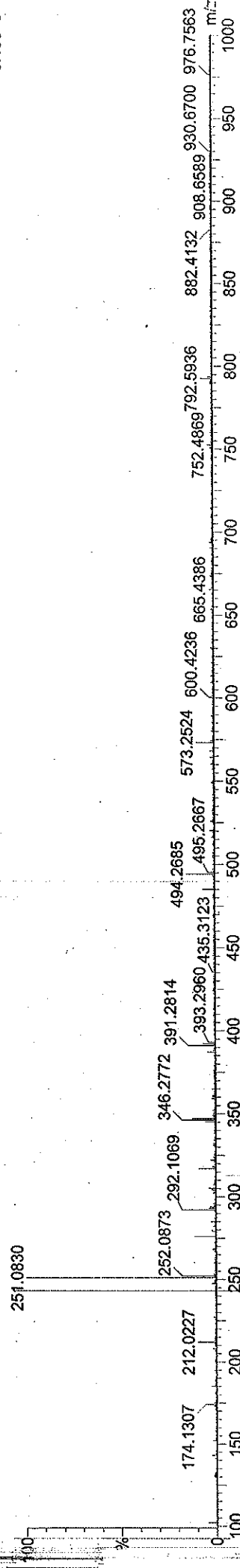
4C5 formula(e) evaluated with 1 results within limits (up to 4 closest results for each mass)

Elements Used:

C: 0-45 H: 0-70 N: 0-6 O: 0-6 S: 0-2

A558QDR104

U71010_191 22 (0.412) Cm (21:24-73:80)

1: TOF MS ES
3.15e+0

Minimum:

Maximum:

Mass

Calc. Mass

mDa

PPM

DBE

i-FIT

Formula

251.0830 251.0821

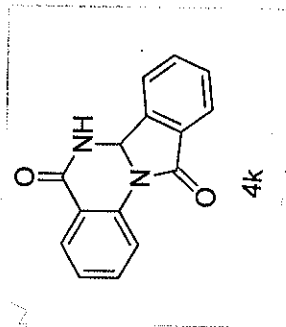
0.9

3.6

11.5

3.5

C15 H11 N2 O2



TDC-204 A558/QDR/34 in DMSO

NMR-400

AR.No:ME1110/1509

Analyst:Haribabu.R

Date: 19 th Nov 2010

Sample Name:

1509-TDC-204_A558_QDR_34

Data Collected on:

mercuryj-mercury400

Archive directory:

/home/sms50/routine/Nov2010/d_19

Sample directory:

1509-TDC-204_A558_QDR_34_20101120_01

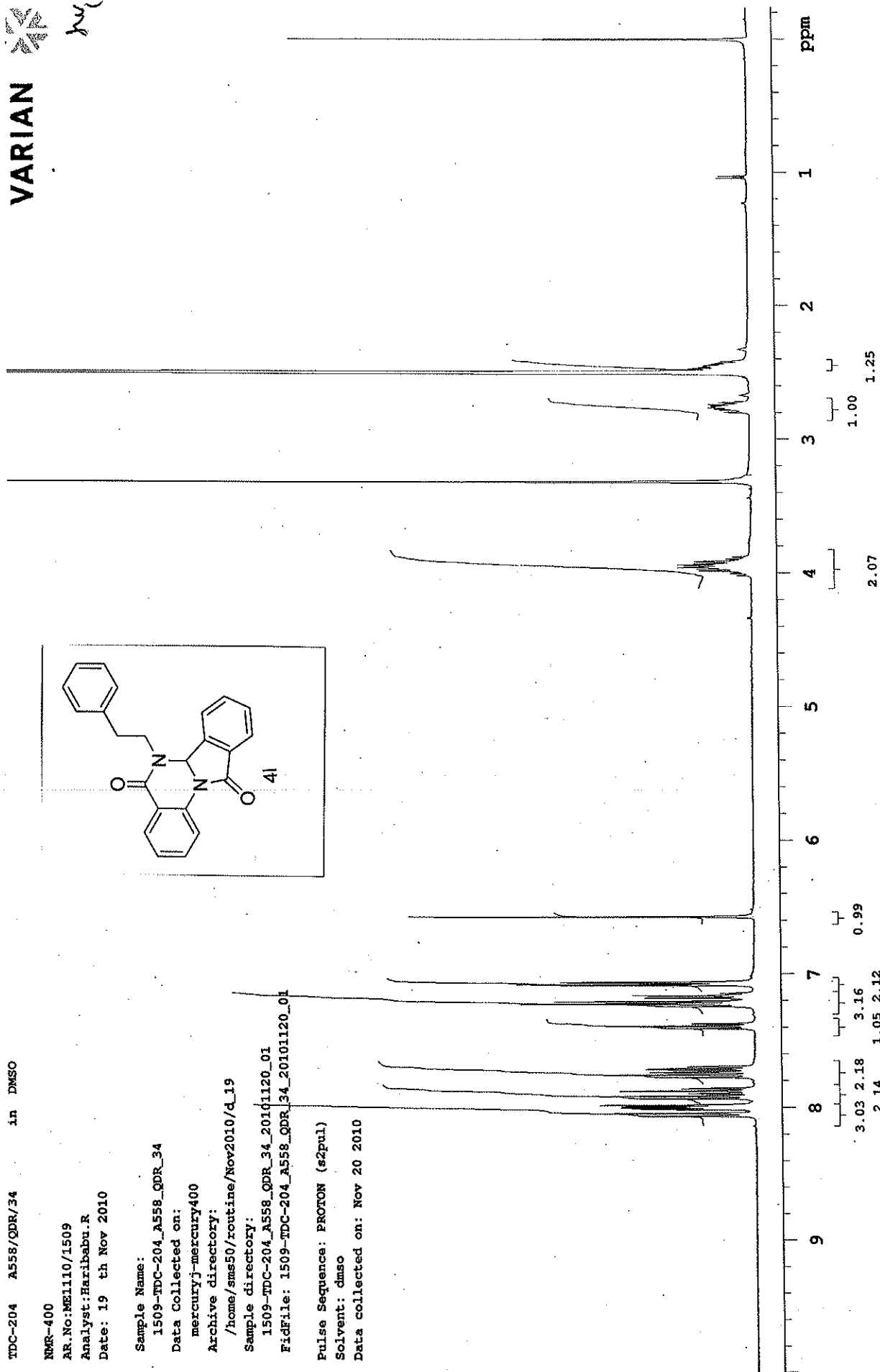
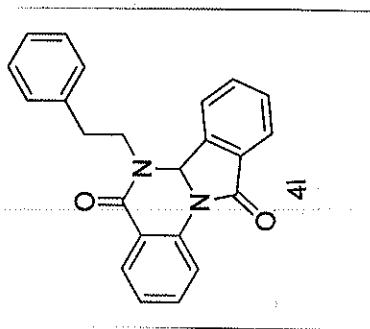
FidFile: 1509-TDC-204_A558_QDR_34_20101120_01

Pulse Sequence: PROTON (s2pul)

Solvent: dmsO

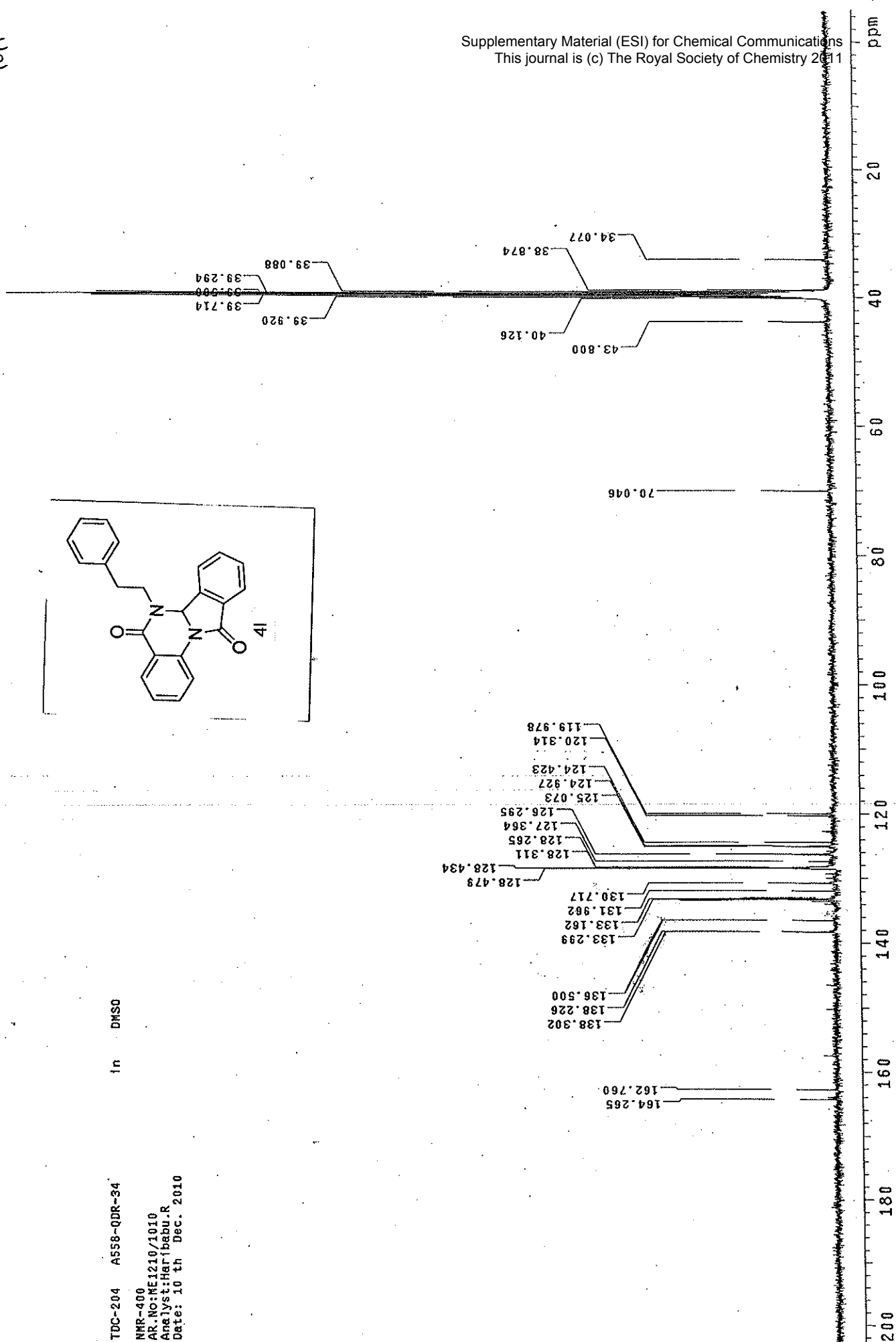
Data collected on: Nov 20 2010

VARIAN



Plotname: 1509-TDC-204_A558_QDR_34_20101120_01_Plot01

20101010



in DMSO

TDC-204 A558-QDR-34

NMR-400

AR No: ME1210/1010

Analyst: Haribabu.R

Date: 10 th Dec. 2010

Elemental Composition Report

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.0, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

191 formula(e) evaluated with 2 results within limits (up to 4 best isotopic matches for each mass)

Elements Used:

C: 0-40 H: 0-55 N: 0-4 O: 0-8

A558/QDR/34

UT1210_149 17 (0.325) Cm (17:20-66:72)

1: TOF MS ES+
7.45e+003

