Palladium-catalyzed C–N cross coupling of sulfinamides and aryl halides

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Supporting Information

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General Reagent Information

All reactions were carried out under an argon atmosphere. All glassware used was dried in electric oven at 120 °C. Racemic, (R)- and (S)-tert-butanesulfinamides are purchased from Jingzhou Winchem Pharm. & Chem. Co., Ltd. All other chemicals were purchased and used as received.

General Analytical Information

All compounds were characterized by ¹H NMR, ¹³C NMR, ESI-MS and IR spectroscopy, and elemental analysis. Copies of the ¹H and ¹³C spectra can be found at the end of the Supporting Information. Nuclear Magnetic Resonance spectra were recorded on a 300MHz instrument or a 400 MHz instrument. All ¹H NMR experiments are reported in δ units, parts per million (ppm), and were measured relative to the signals for residual chloroform (7.26 ppm) in the deuterated solvent, unless otherwise stated. All ¹³C NMR spectra are reported in ppm relative to deuteron-chloroform (77.23 ppm), unless otherwise stated, and all were obtained with ¹H decoupling. All IR spectra were taken on an infrared spectrometer. All chiral HPLC analyses were performed on a liquid chromatography with a Chiralcel OD-H chiral column (4.6 mm × 250 mm × 5 µm). All rotation data are recorded on an auto rotation (Na D line, cell long 10cm, λ = 589 nm). Electron-spraying ionization mass spectra are recorded on an LC/MS instrument. Elemental analyses of these compounds are performed on an elemental analyzer.

X-ray structure determination

A single crystal of (*R*)-*N*-(3-methoxyphenyl) *tert*-butanesulfinamide was cultured from a solution of petroleum ether and ethyl acetate in a test tube. The single-crystal data were collected on an CCD diffractometer. X-ray generator was operated at 50 kV and 1 mA using an Enhance Mo*Ka* radiation. Data were collected with a ω scan width of 1°. The data reduction, empirical absorption correction, and space group determination were done using CrysAlisPro RED (Oxford Diffraction, 2009). The crystal structure was solved by direct methods and refined by full-matrix leasts quares method using SHELXL97, present in the program suite WinGx (Version 1.63.04a). The molecular diagrams were generated using ORTEP 3 and the packing diagrams were generated using CAMERON. Geometrical calculations were done using PARST95 and PLATON. The positions of all H atoms were fixed geometrically and refined isotropically using the riding atom model.

Crystal Data for (*R*)-*N*-(3-methoxyphenyl) *tert*-butanesulfinamide 3g. $C_{11}H_{18}NO_2S$, *M* =228.32, Orthorhombic, *a* = 7.4418(9) Å, *b* = 9.7027(12) Å, *c* = 16.862(2) Å, *U* = 1217.5(3) Å³, *T* = 293.0, space group P2₁2₁2₁ (no. 19), *Z* = 4, μ (Mo K α) = 0.248, 7010 reflections measured, 2481 unique (*R*_{int} = 0.0309) which were used in all calculations. The final *wR*(*F*₂) was 0.0916 (all data).

Description about the single crystal (*R*)-*N*-(3-methoxyphenyl) *tert*-butanesulfinamide 3g.

All of the *N*-aryl sulfinamides are crystalline solid and readily to crystalize. In order to probe the internal reasons of these crystalline *N*-aryl sulfinamides, three-dimensional molecular structures and crystal-packing modes of *N*-aryl *tert*-butanesulfinamide, a crystal of (*R*)-*N*-(3-methoxyphenyl) *tert*-butanesulfinamide was cultured and analyzed.¹ The ORTEP style plot (Figure 1) shows the three dimension structure of (*R*)-*N*-(3-methoxyphenyl) *tert*-butanesulfinamide **3g**, and its stereo-structure has uniform configuration (Figure 2) and is consistent with (*R*)-*N*-(3-methoxyphenyl) *tert*-butanesulfinamide (Figure 1). The crystal stacking plot (Figure 2) clearly demonstrates the strong N–H^{...}O=S hydrogen bonds in a head-to-tail fashion throughout the single crystal, which is

different from the racemic *N*-aryl tert-butanesulfinamides with a couple of hydrogen bonds between (*R*)- and (*S*)-isomers.² The intermolecular hydrogen bonds throughout the single crystal should be the main intermolecular force to keep *N*-aryl *tert*-butanesulfinamides in crystalline solid and make the melting points of our single enantiomer products commonly higher than that of the corresponding racemic ones.²

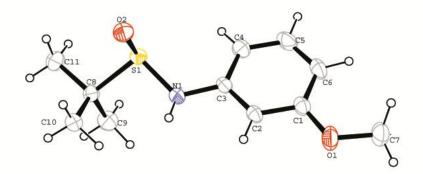


Figure 1. ORTEP style plot of (R)-N-(3-methoxyphenyl) *tert*-butanesulfinamide **3g** in the solid state. Thermal ellipsoids are drawn at the 50% probability level.

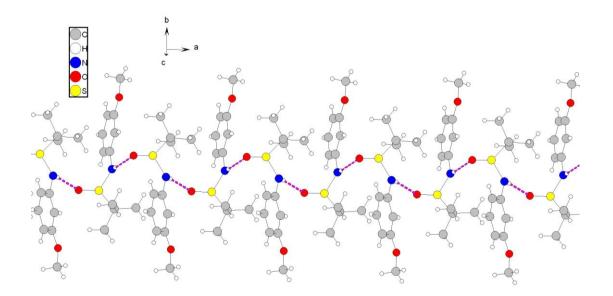
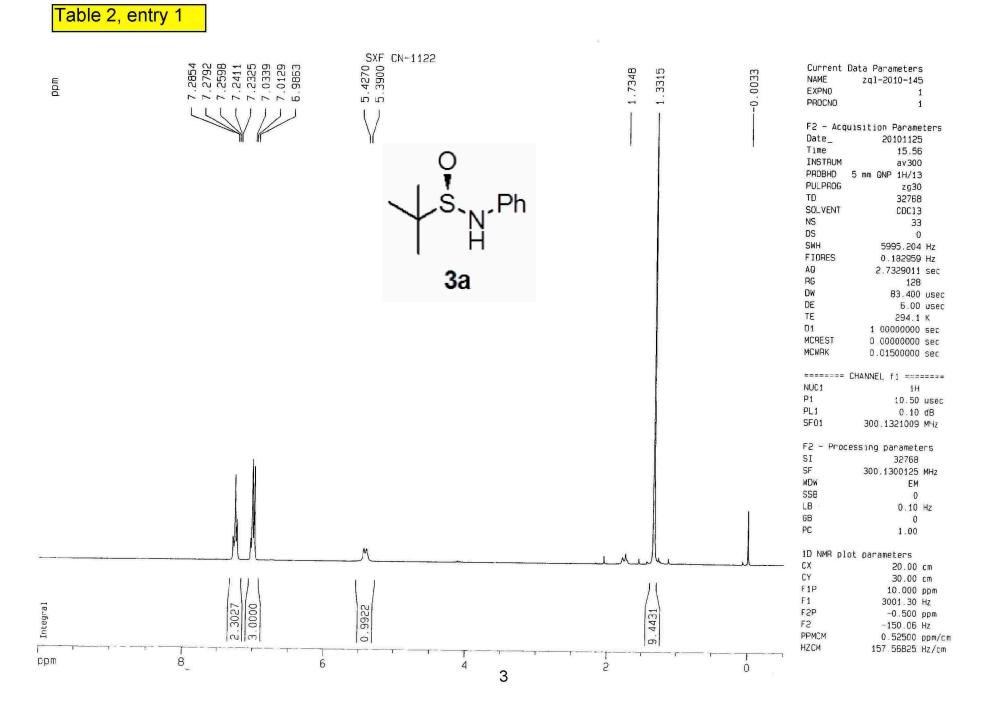
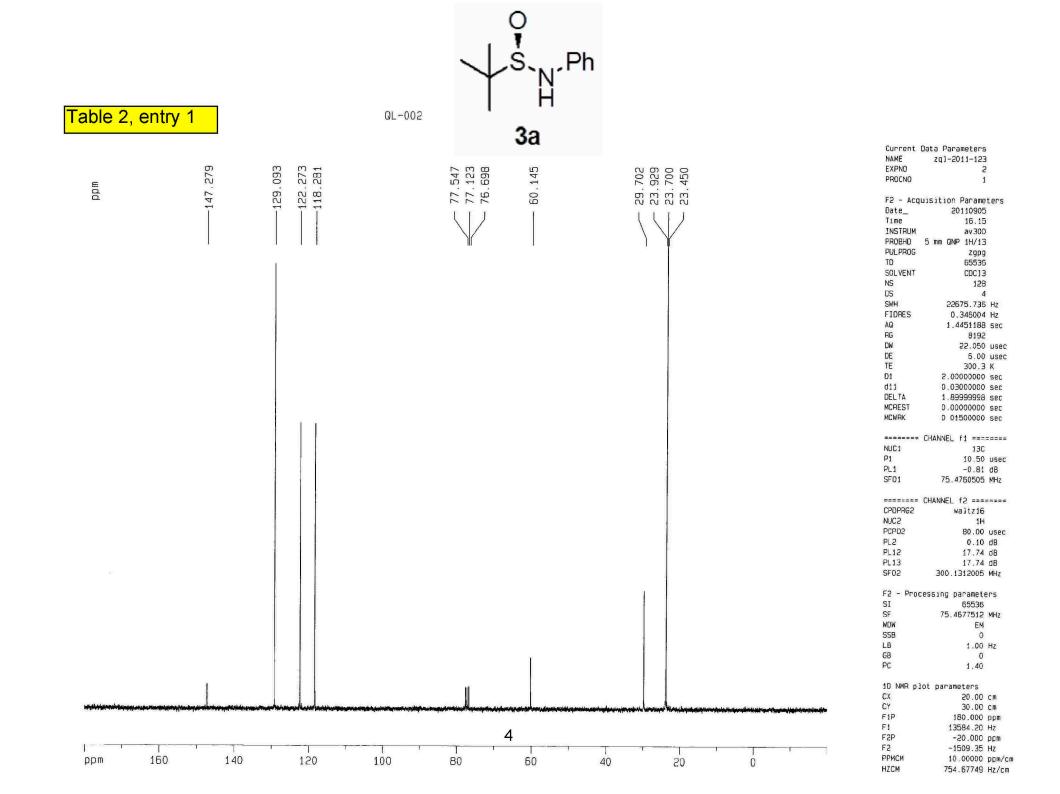


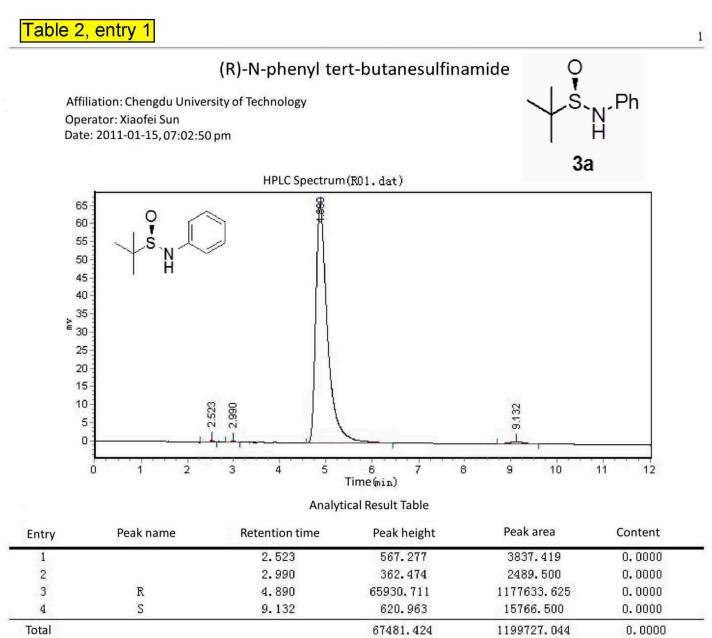
Figure 2. Crystal stacking plot of (*R*)-*N*-(3-methoxyphenyl) *tert*-butanesulfinamide 3g.

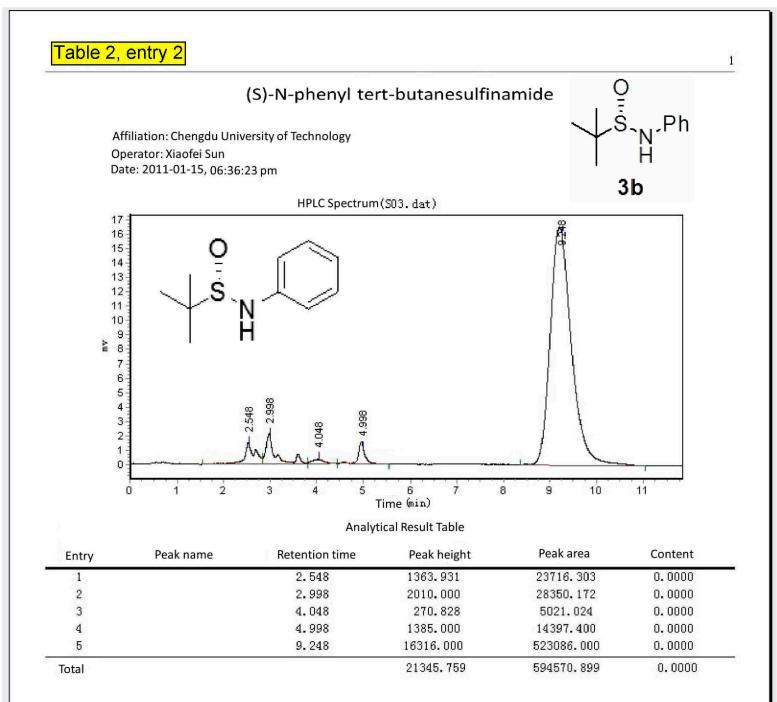
References

- (1) CCDC 862954 contains the supplementary crystallographic data for this paper. These data can be obtained free of charge from the Cambrige Crystallographic Data Center via <u>www.ccdc.cam.ac.uk/data_request/cif</u>.
- (2) (a) Datta, M.; Buglass, A. J.; Elsegood, M. R. J. Acta Cryst. 2009, E65, o2034. (b) Datta, M.; Buglass, A. J.; Elsegood, M. R. J. Acta Cryst. 2010, E66, o109. (c) Datta, M.; Buglass, A. J.; Elsegood, M. R. J. Acta Cryst. 2009, E65, o2823.

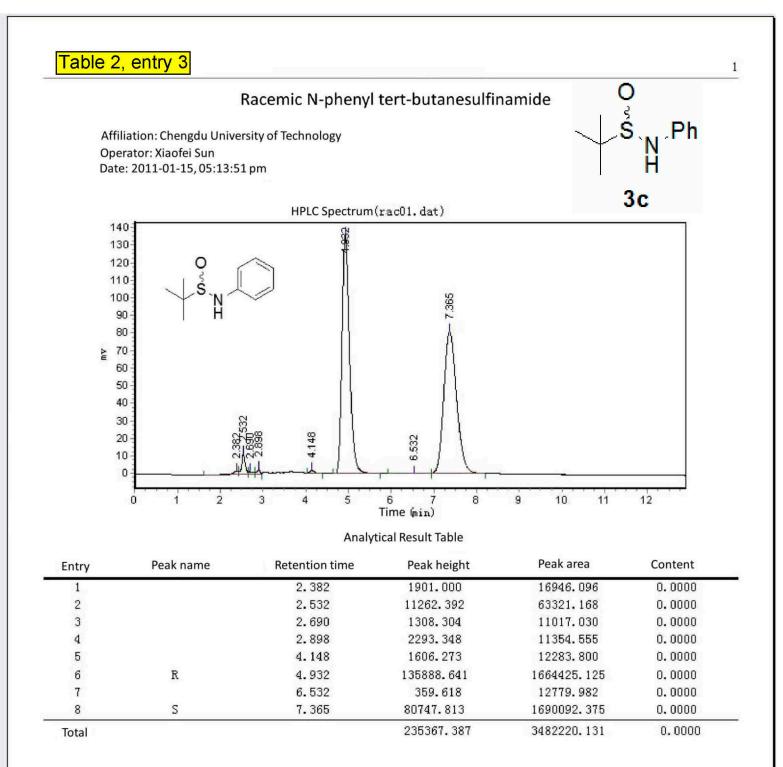








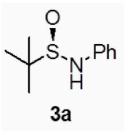
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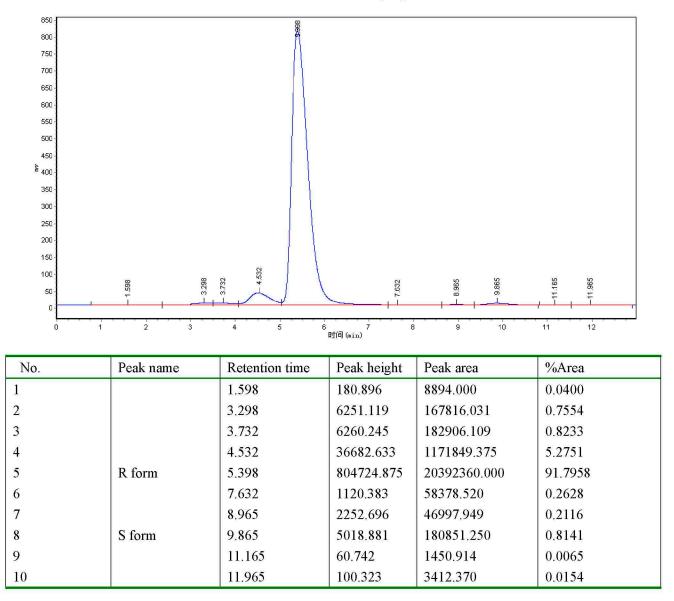
(R)-N-Phenyl tert-butanesulfinamide

(Re-analyzed on March 3, 2012)



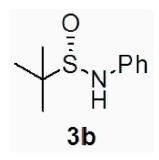
Affiliation: Chengdu University of Technology Date: 2012-03-03 Operator: Xingzhao Tu

HPLC, Diacel Chiralcel OD-H column, 90:10 hexanes/2-propanol, 1 mL/min, 254 nm.



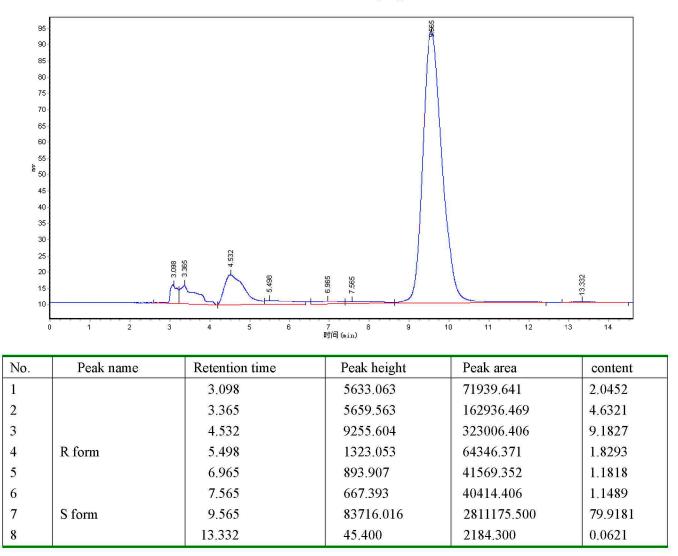
(S)-N-Phenyl tert-butanesulfinamide

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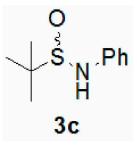
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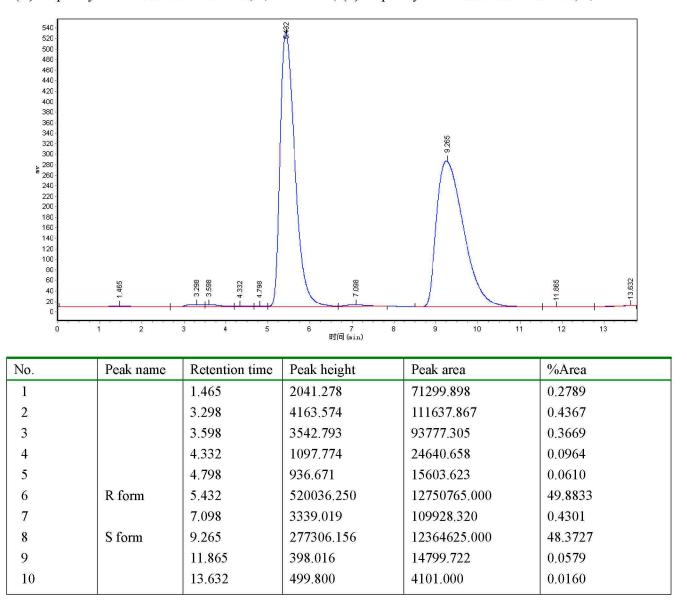
Racemic N-Phenyl tert-butanesulfinamide

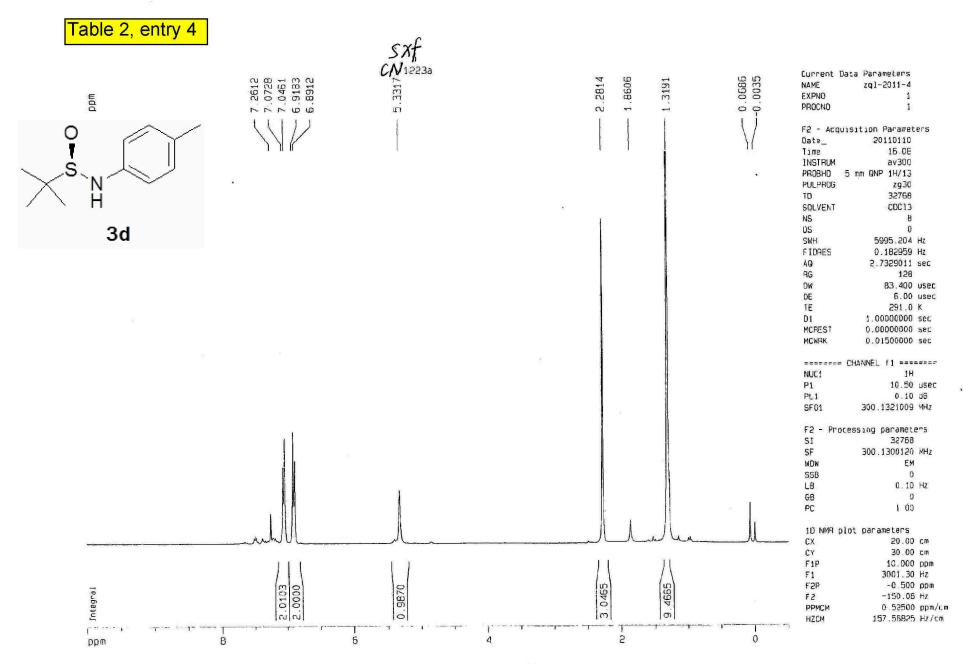
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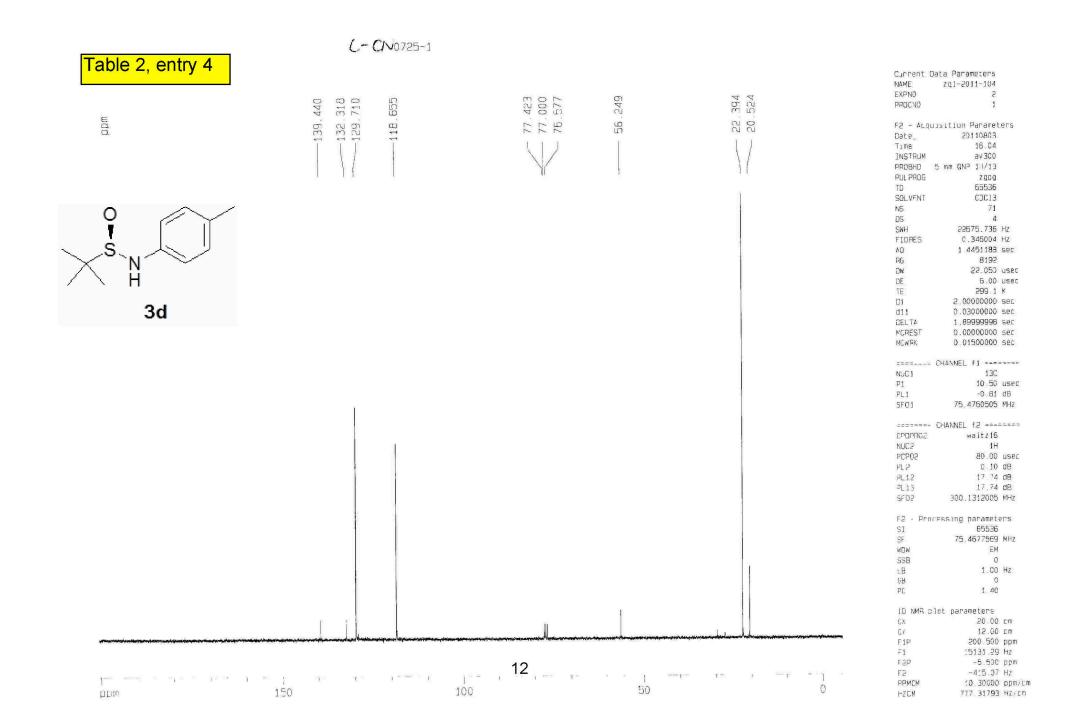


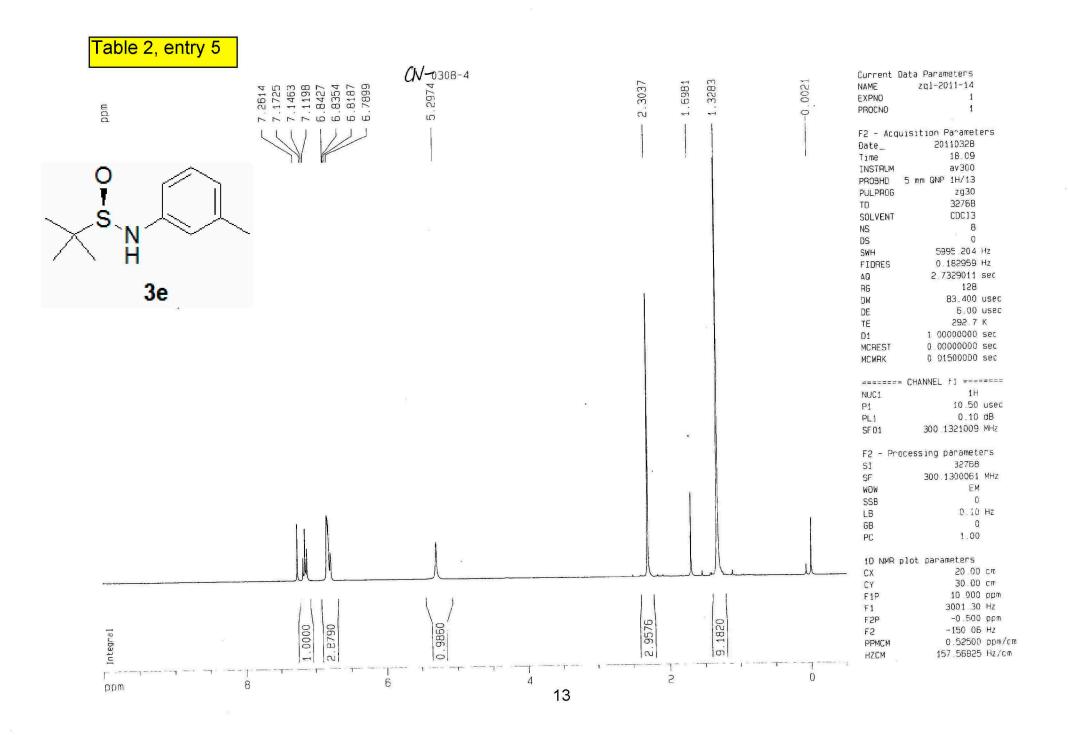
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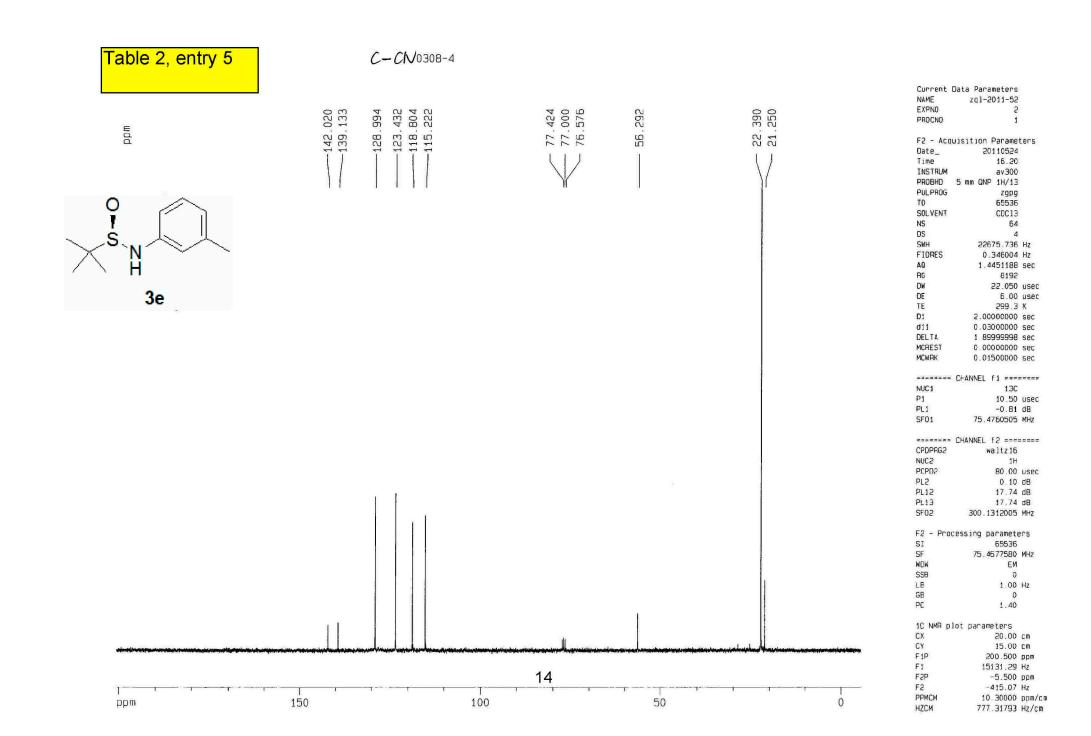
HPLC, Diacel Chiralcel OD-H column, 90:10 hexanes/2-propanol, 1 mL/min, 254 nm. (*R*)-N-phenyl *tert*-butanesulfinamide, $r_1 = 5.4$ min; (*S*)-*N*-phenyl *tert*-butanesulfinamide, $r_1 = 9.2$ min.

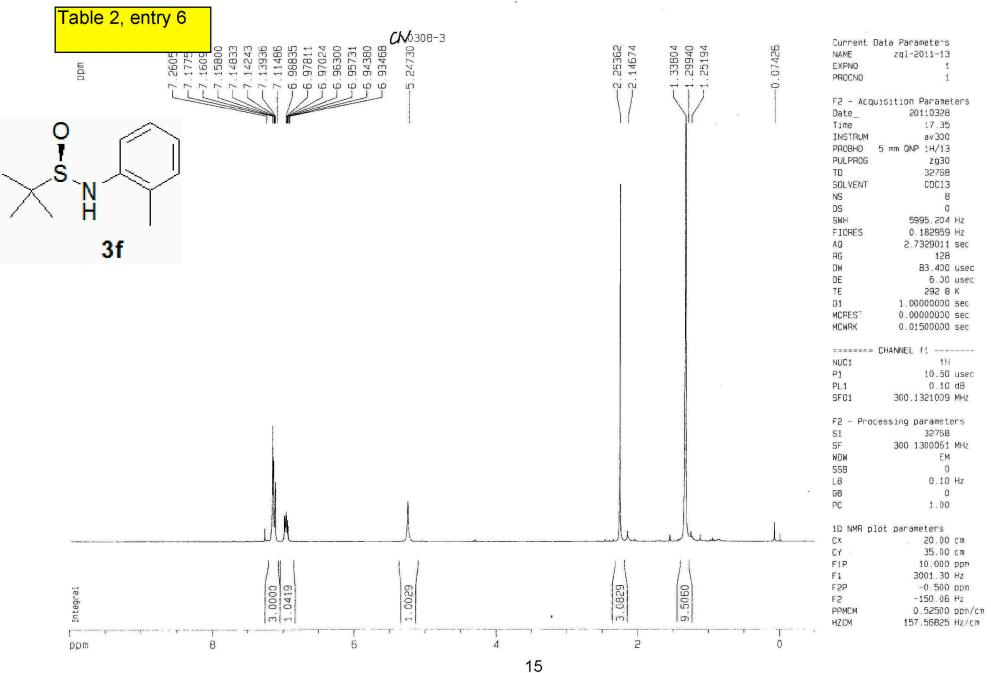


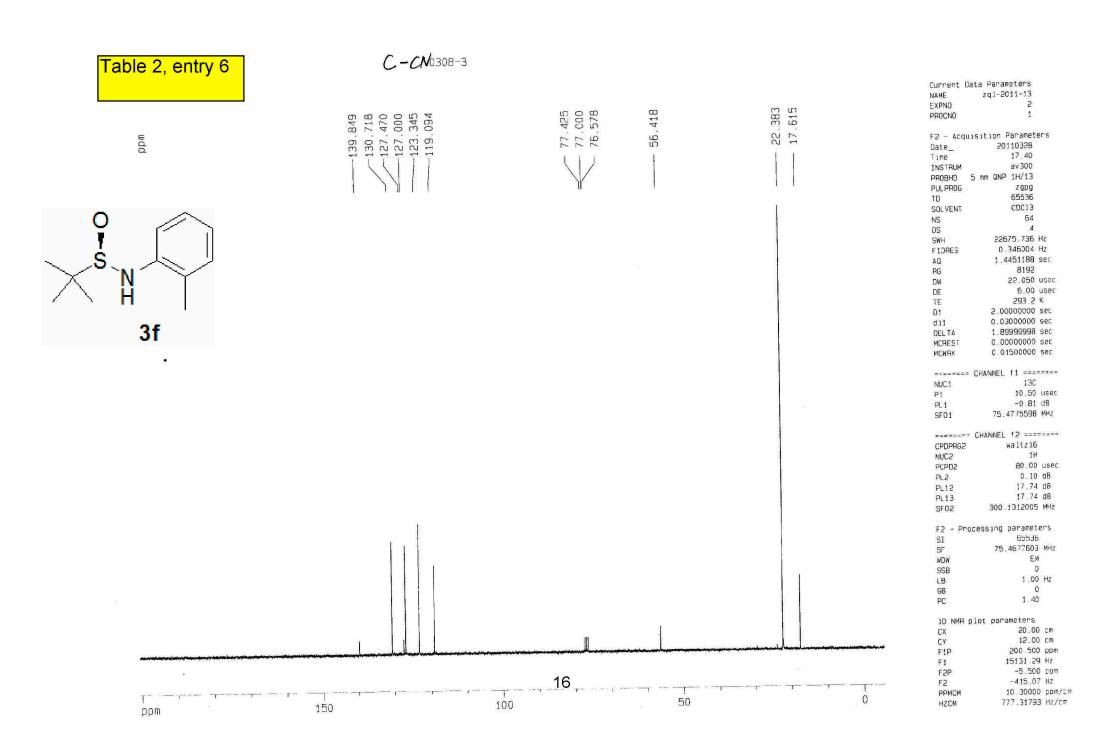


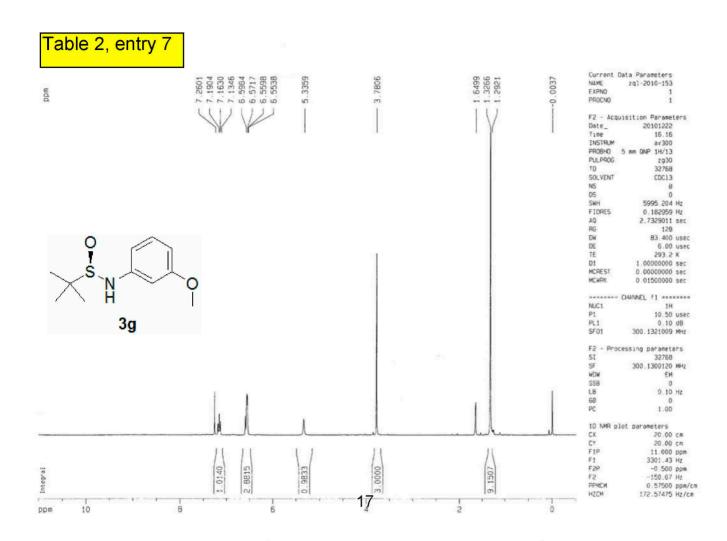


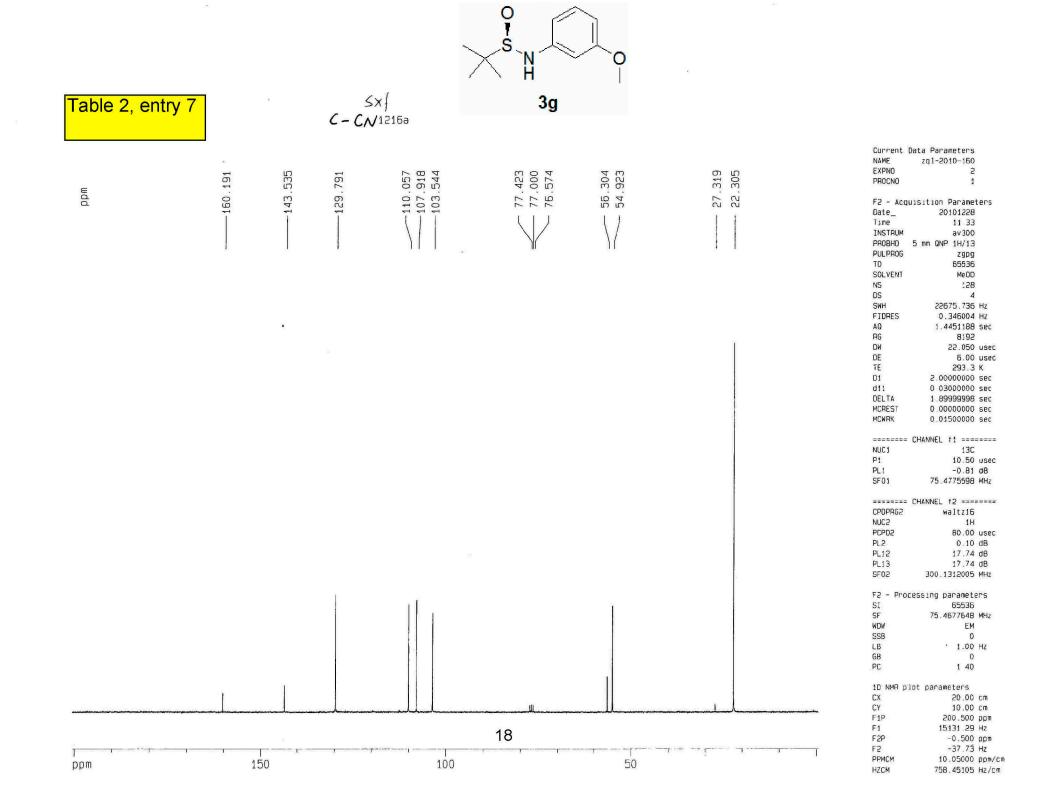


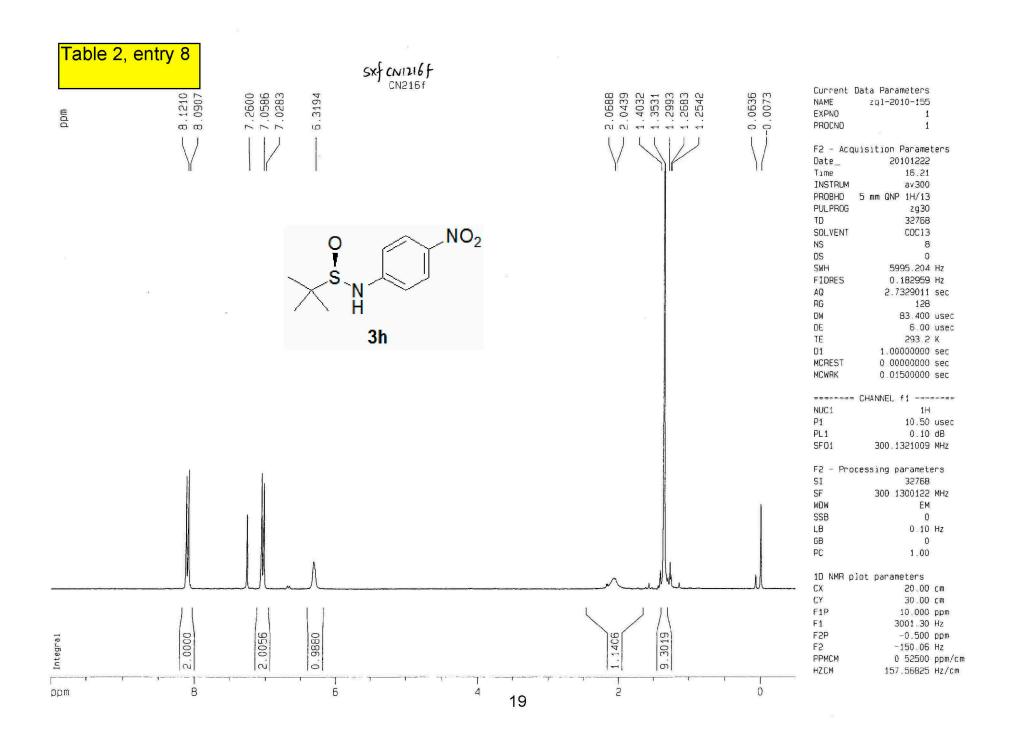








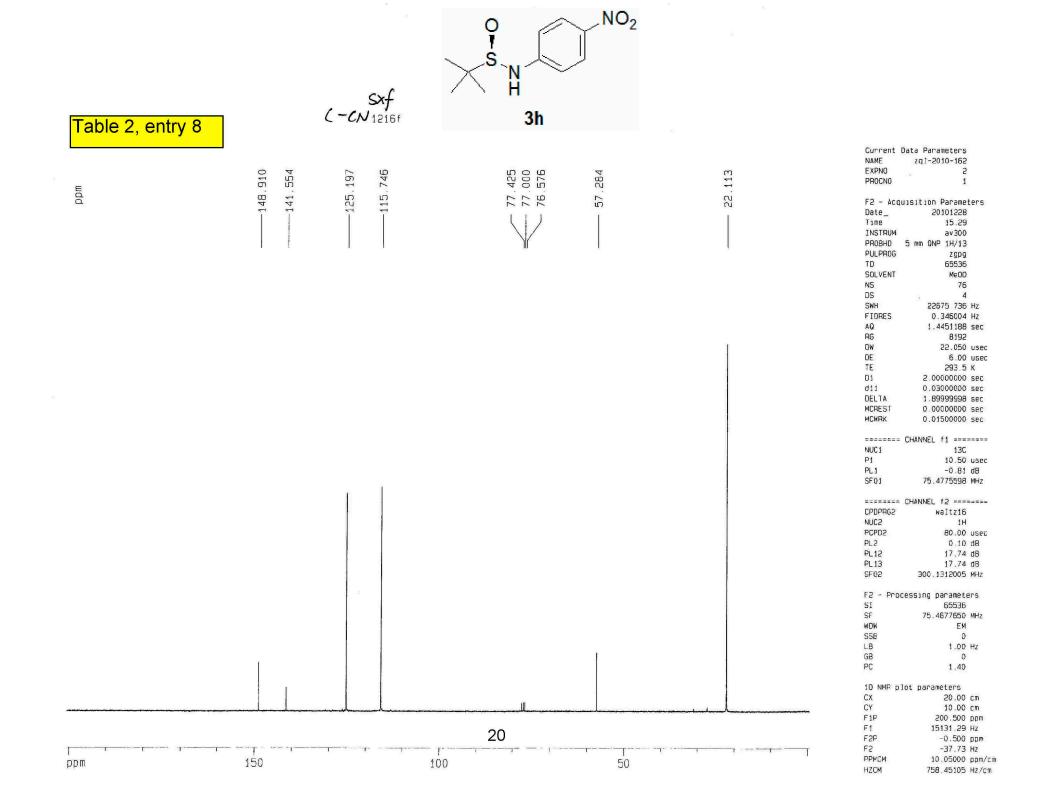


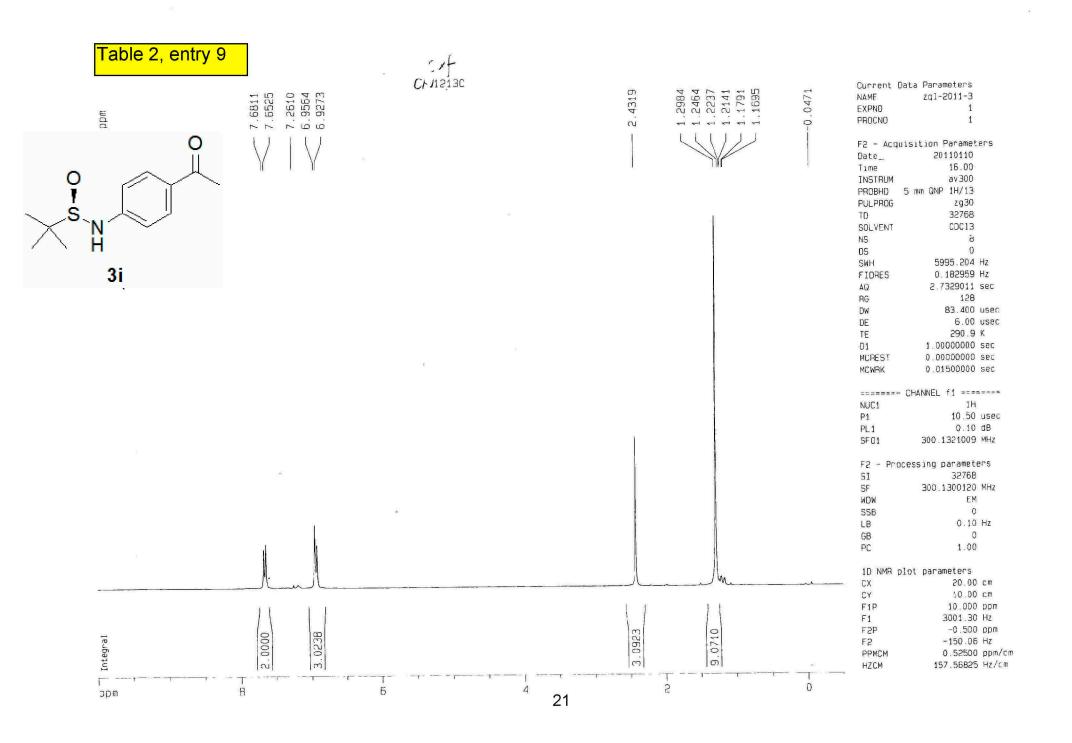


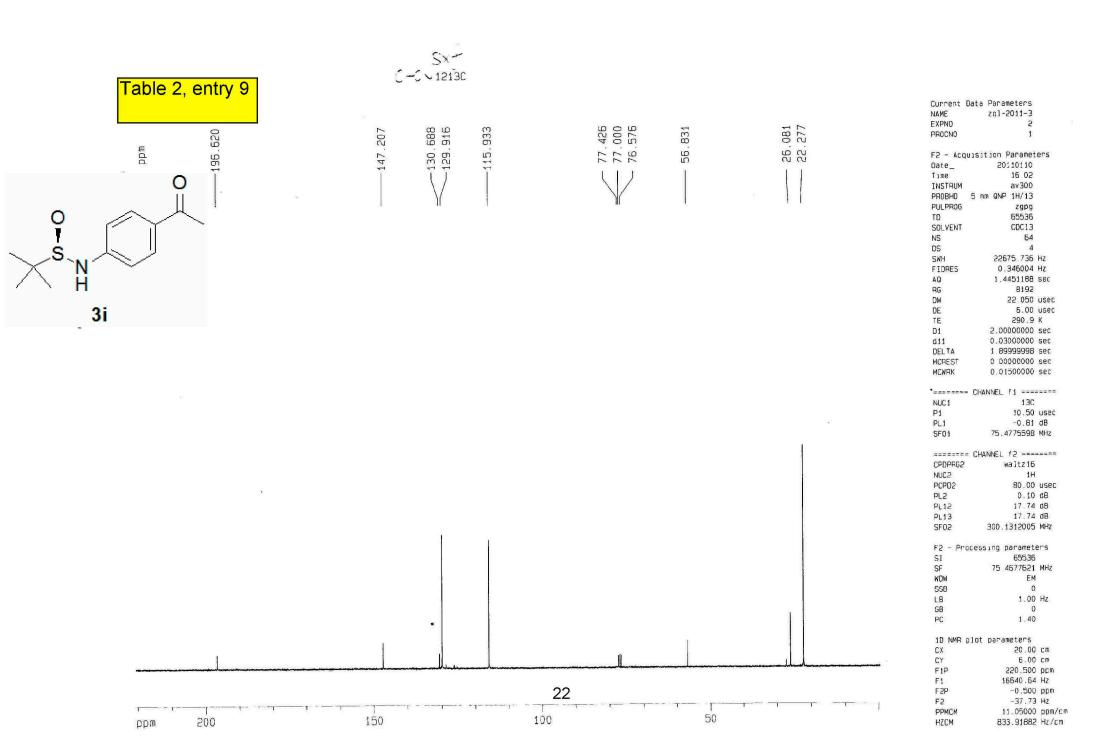
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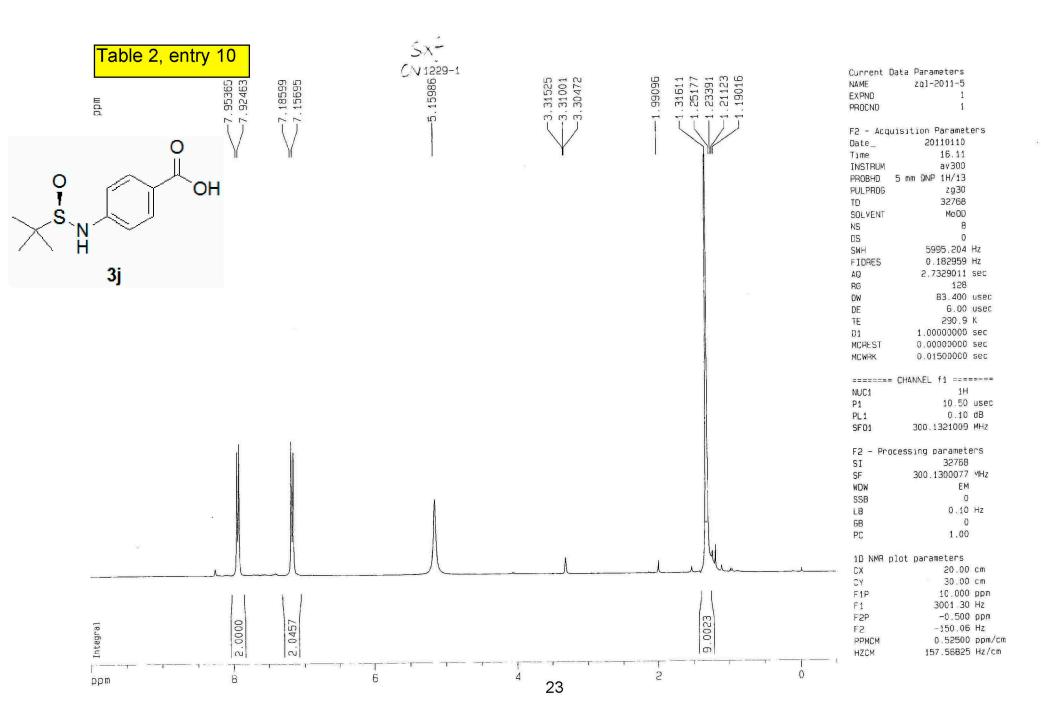
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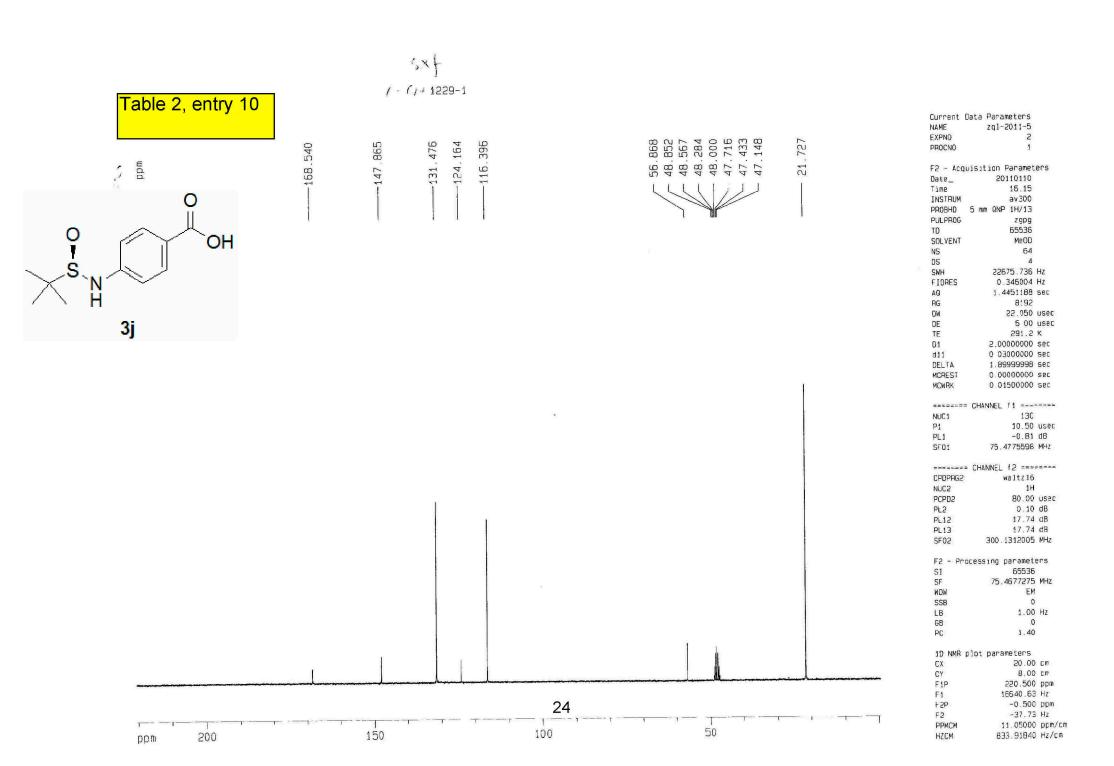


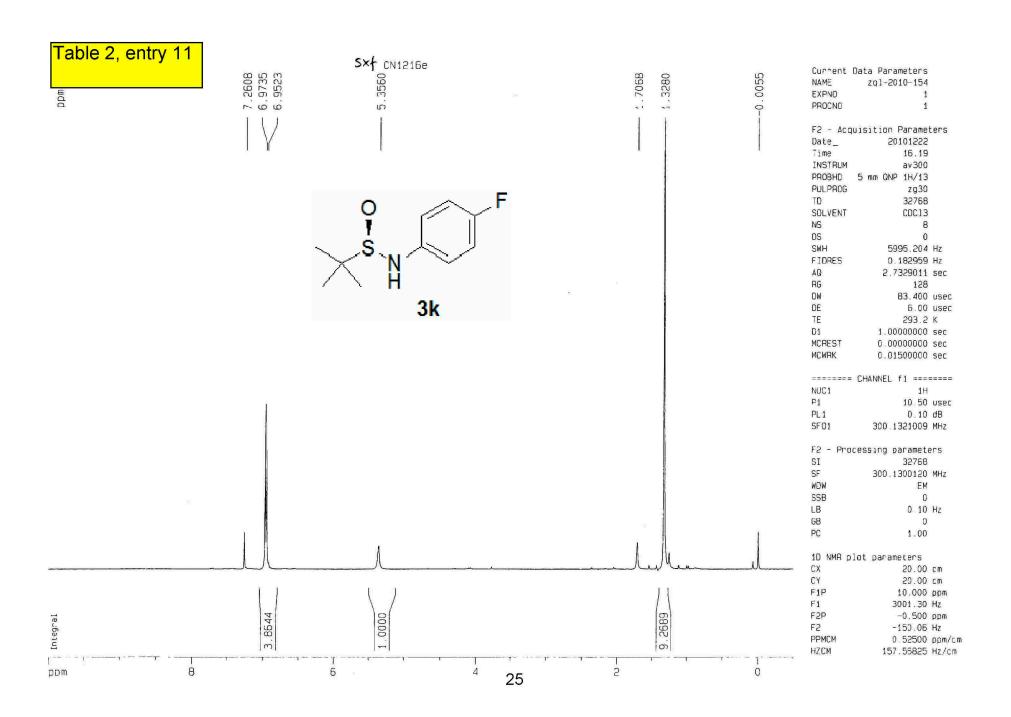




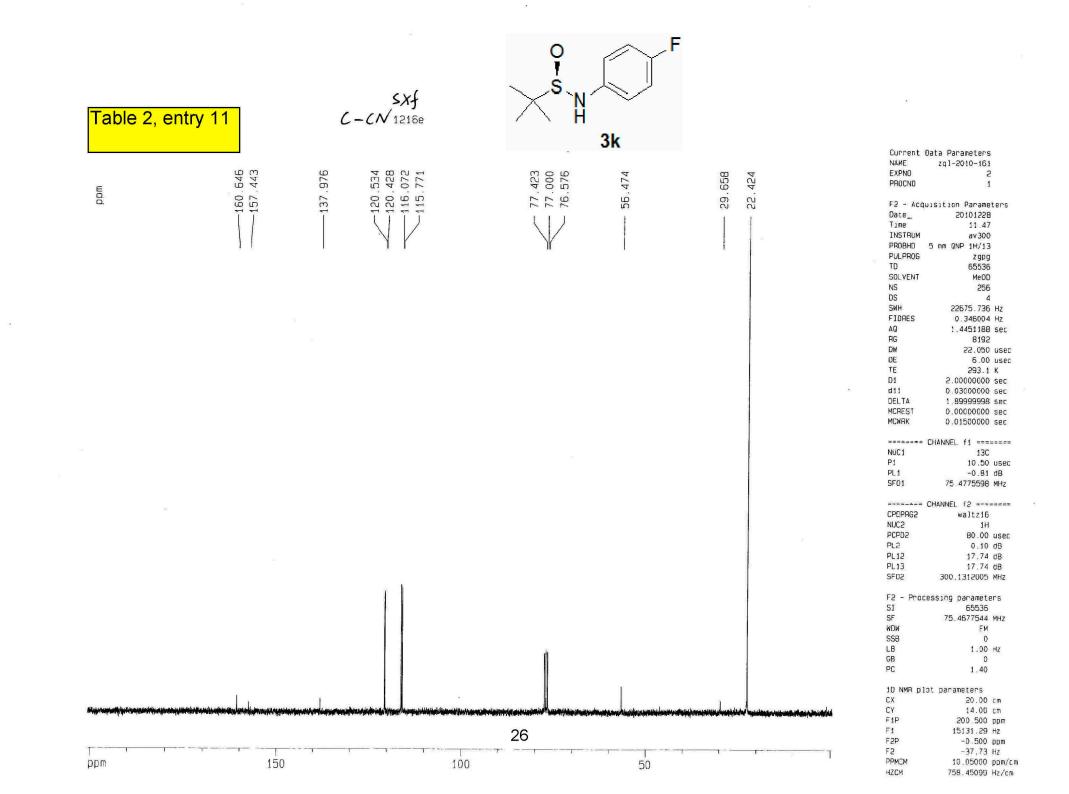
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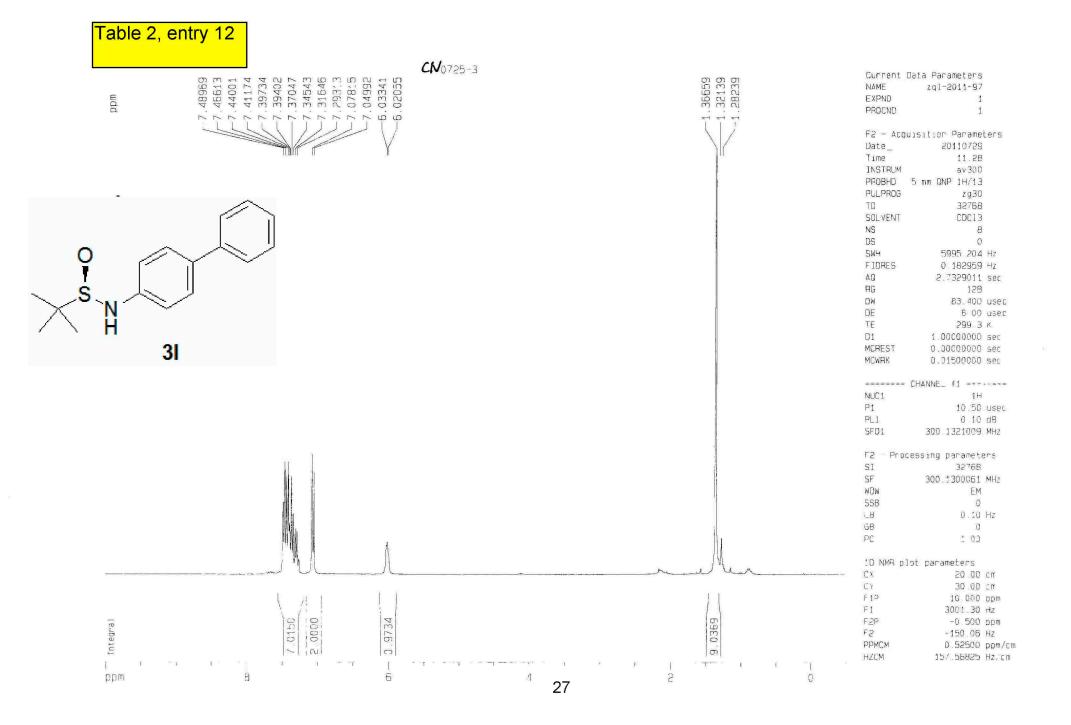
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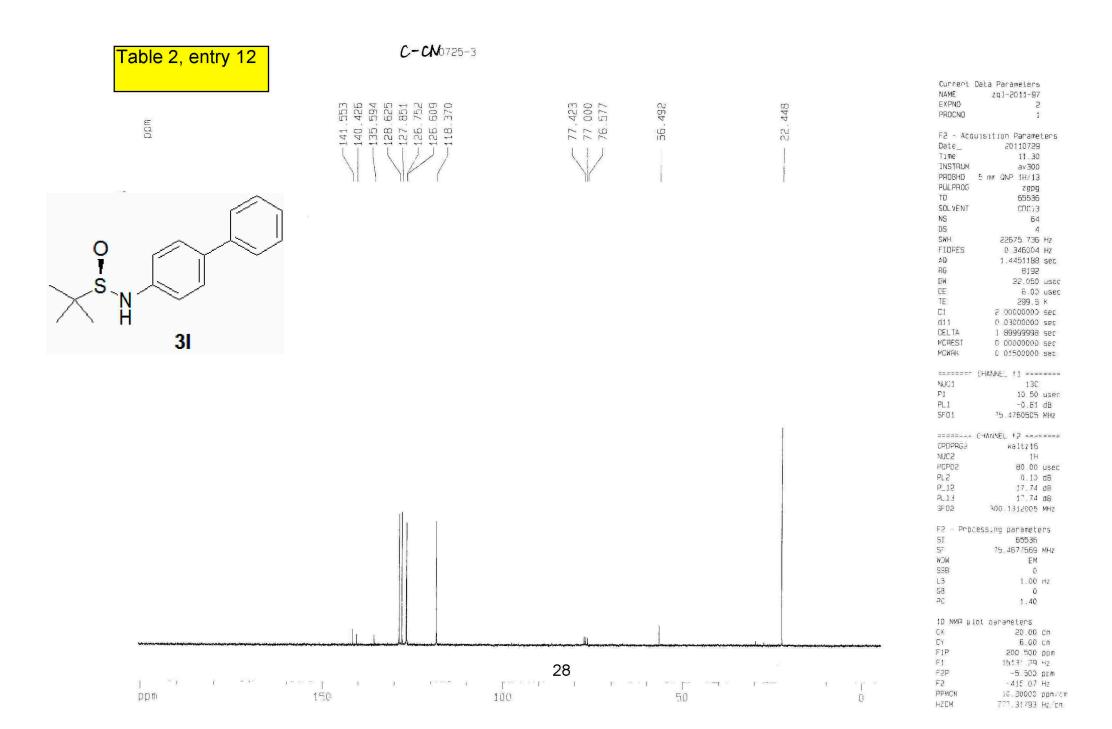


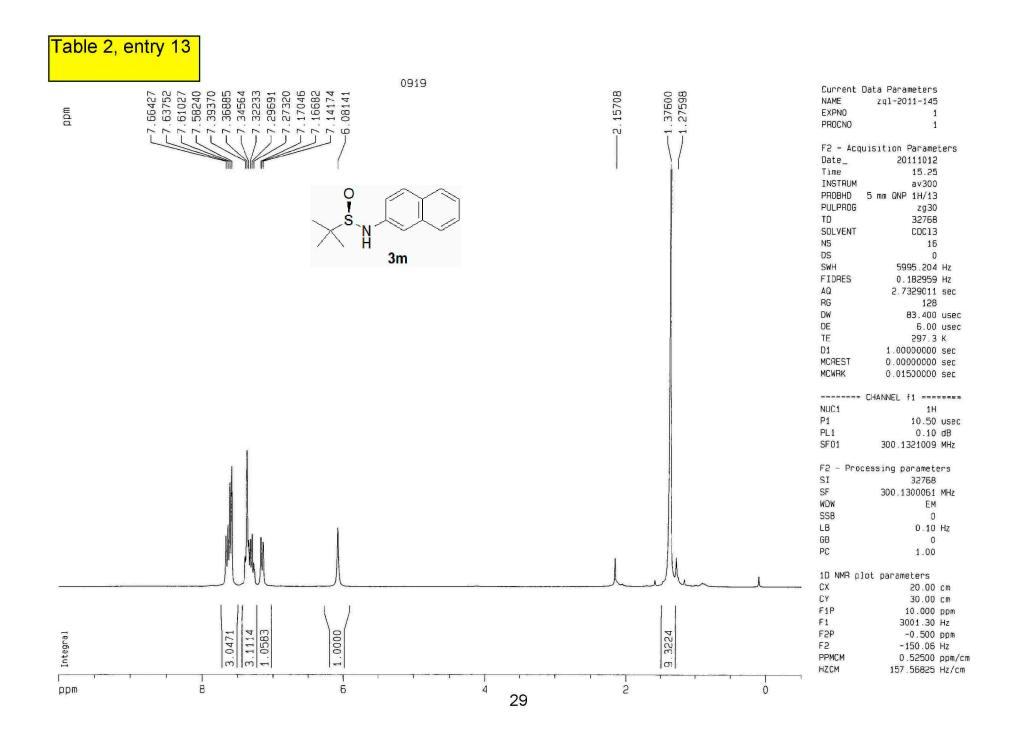


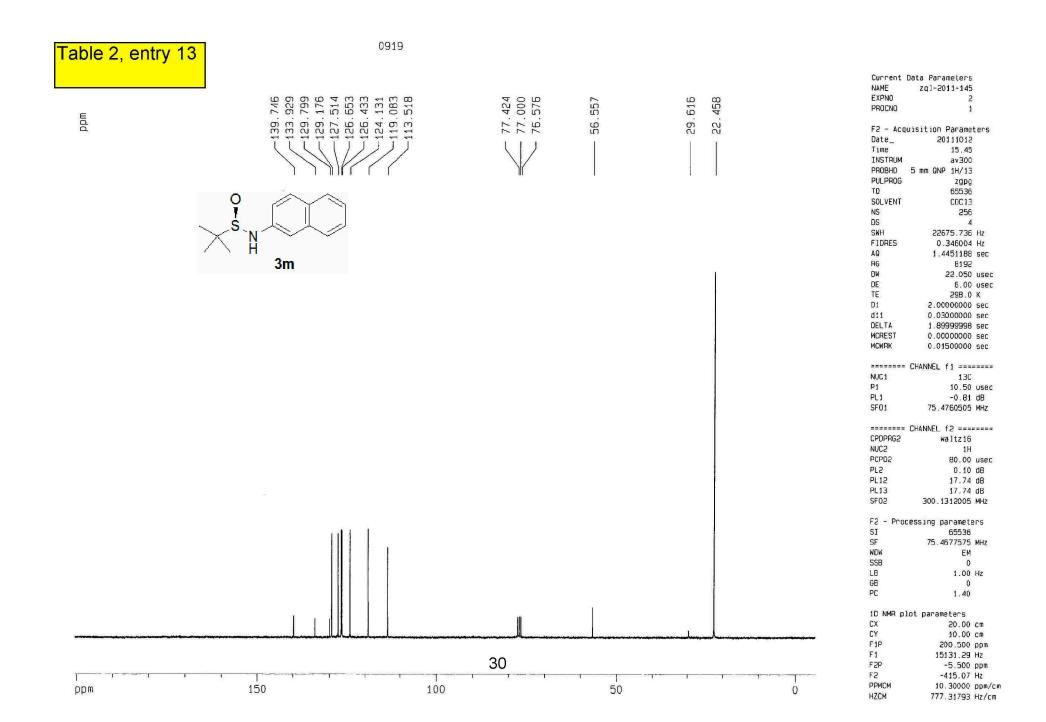
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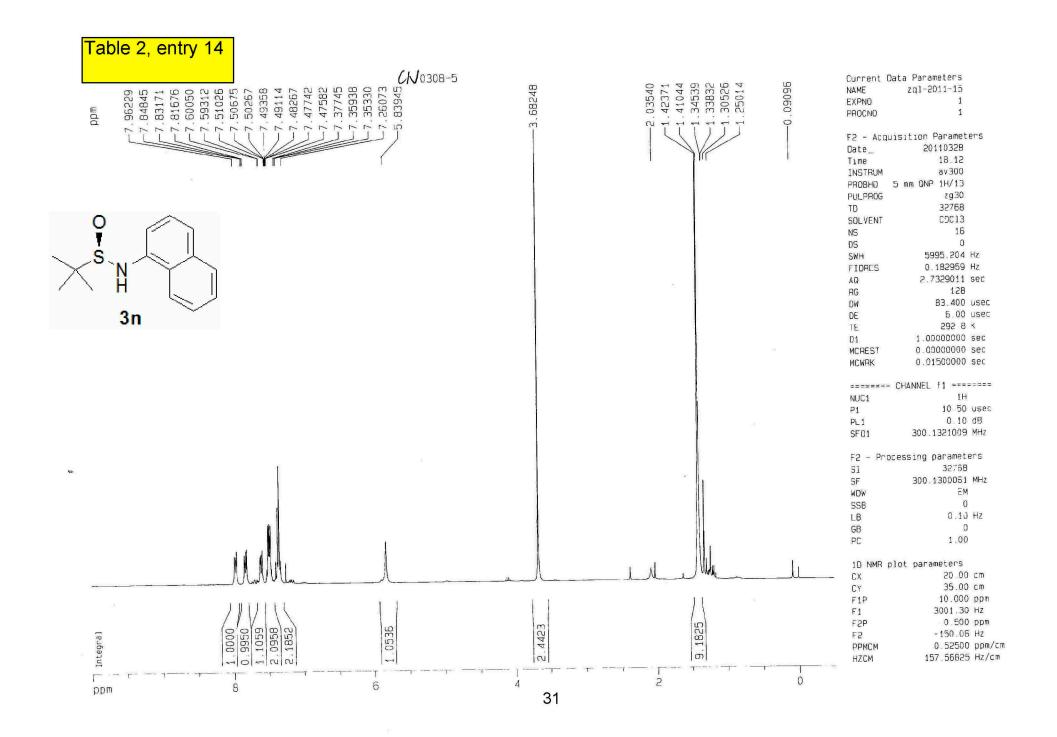


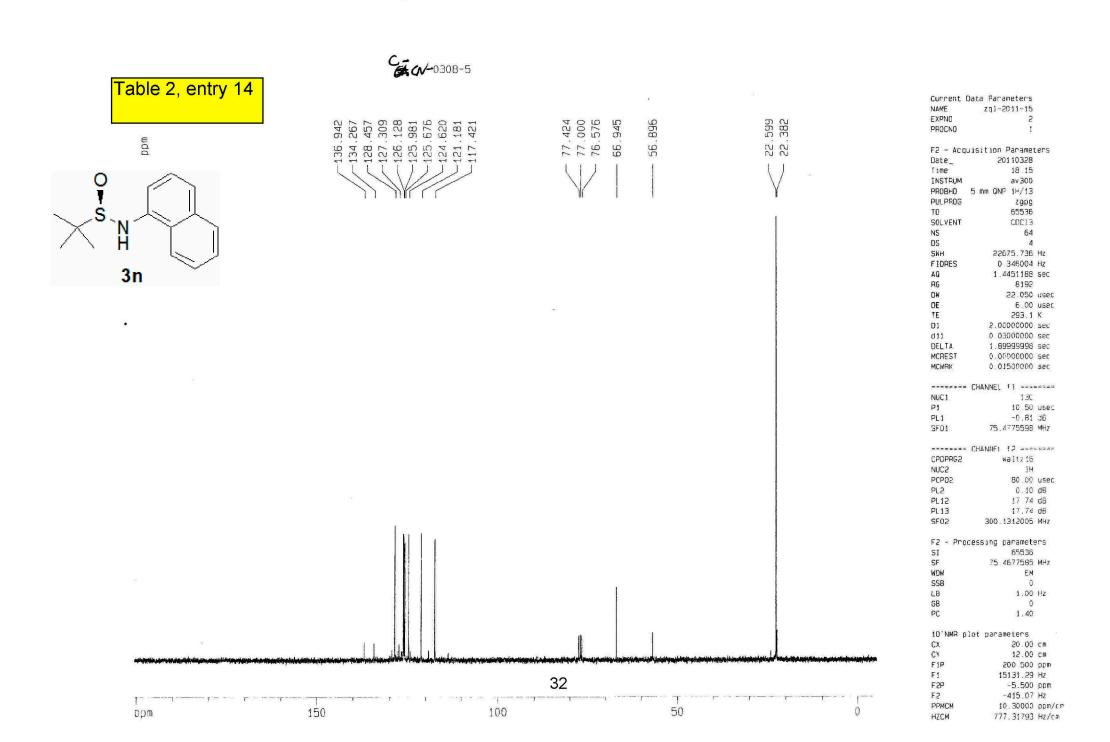


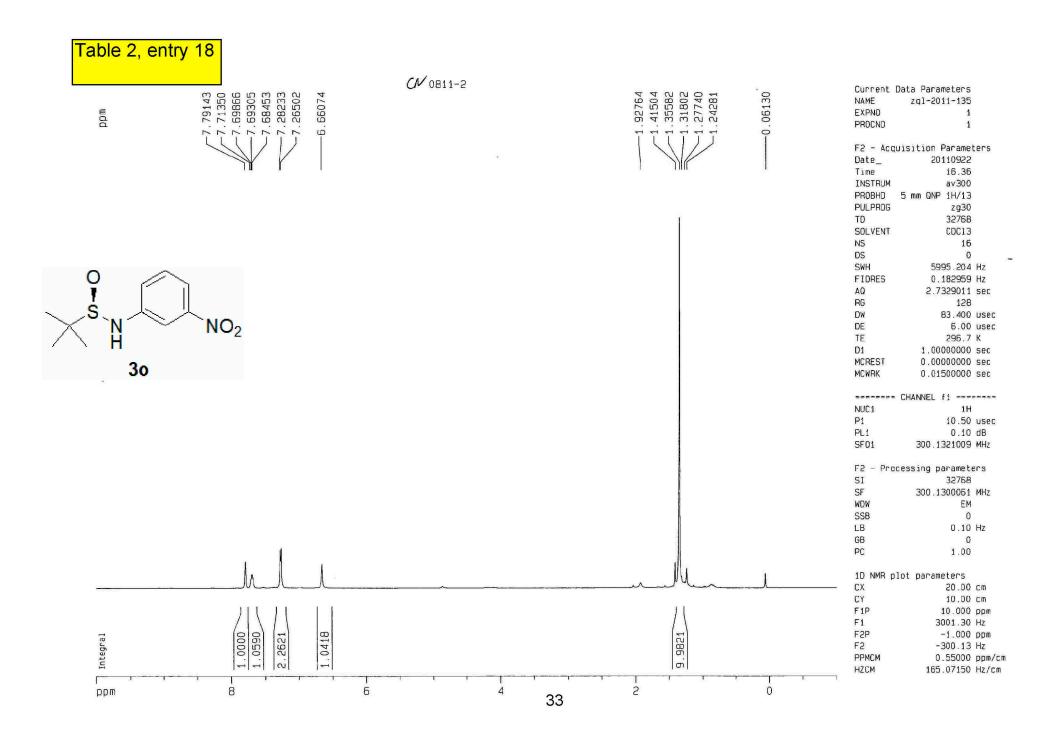


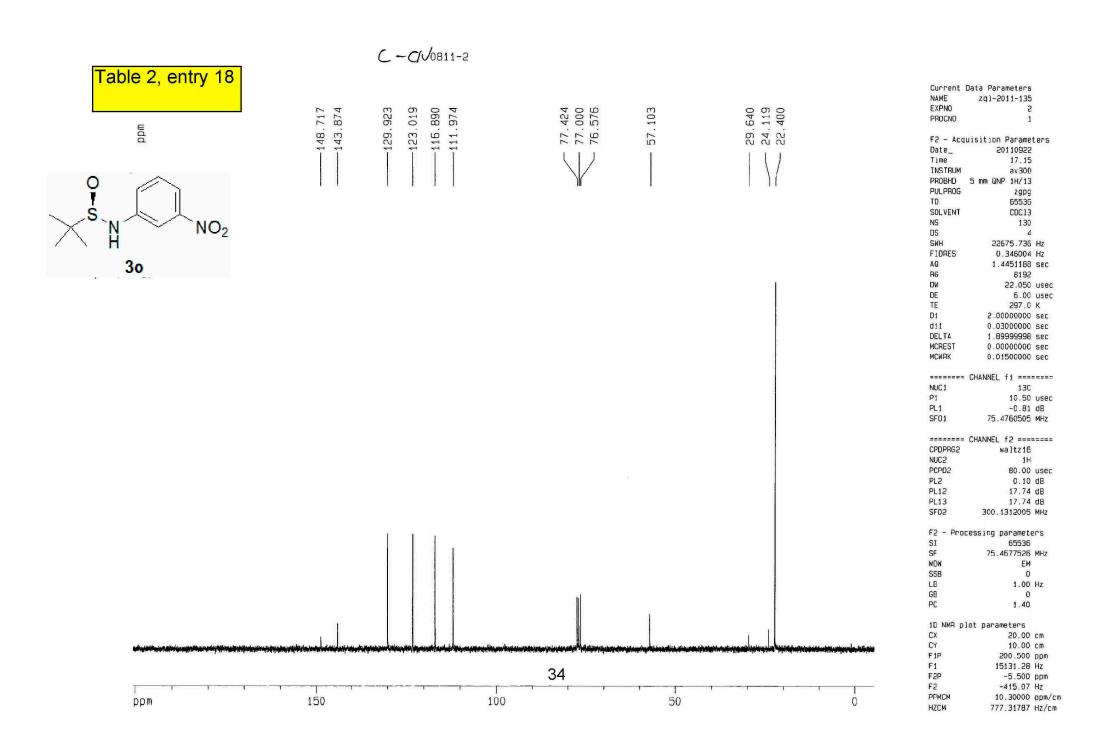


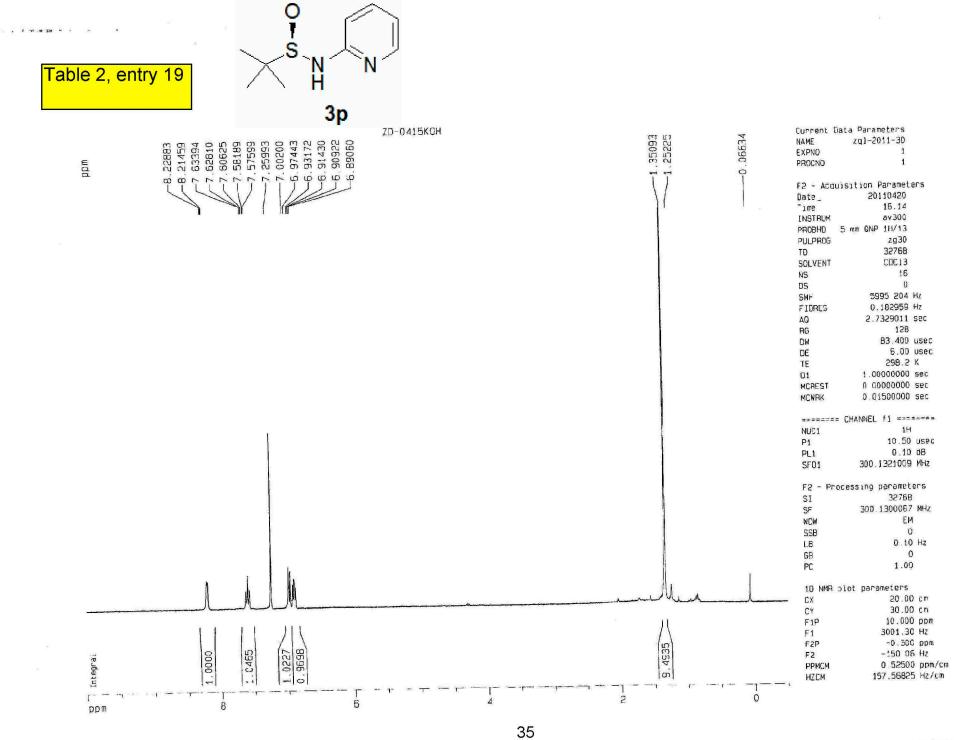






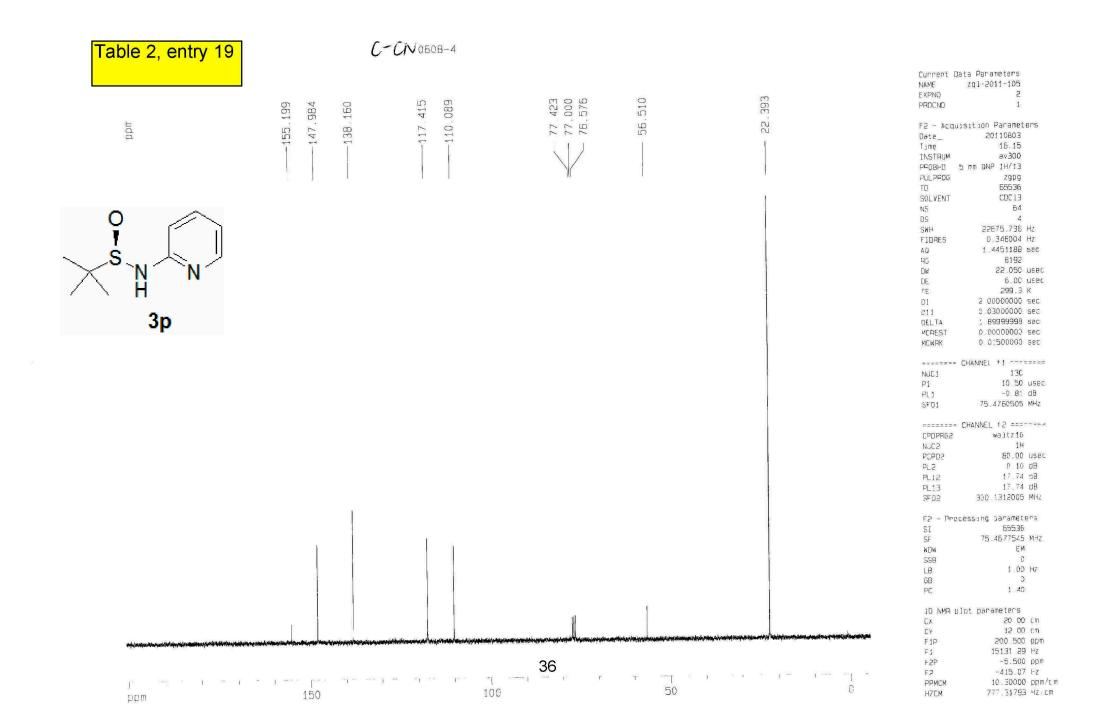


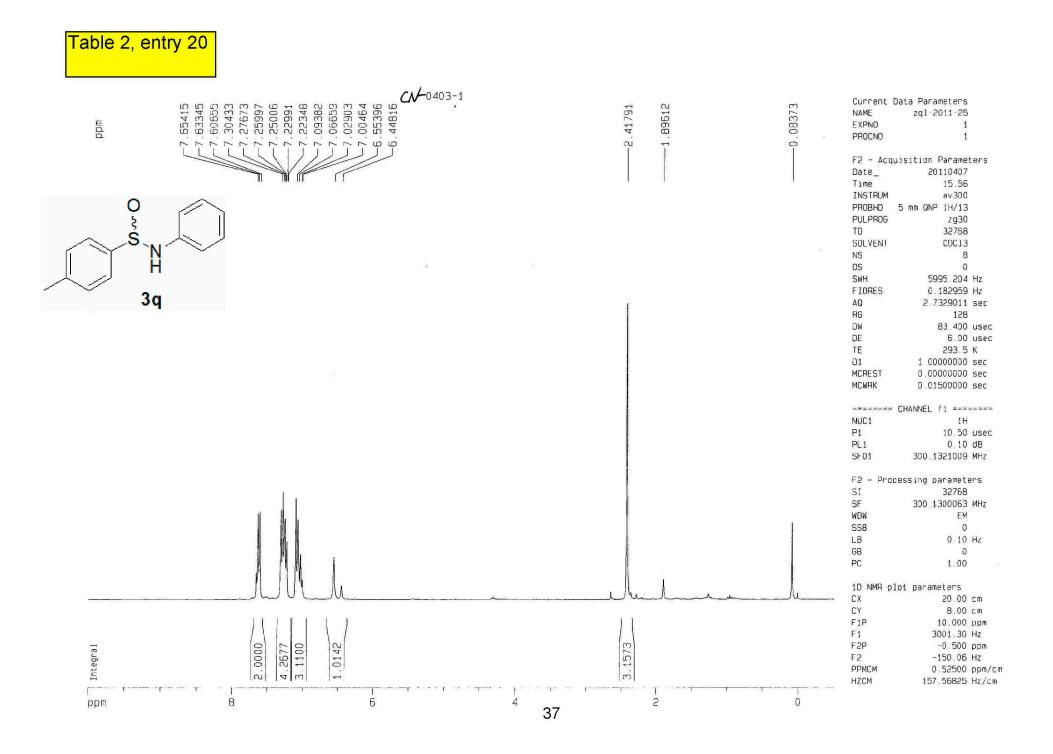


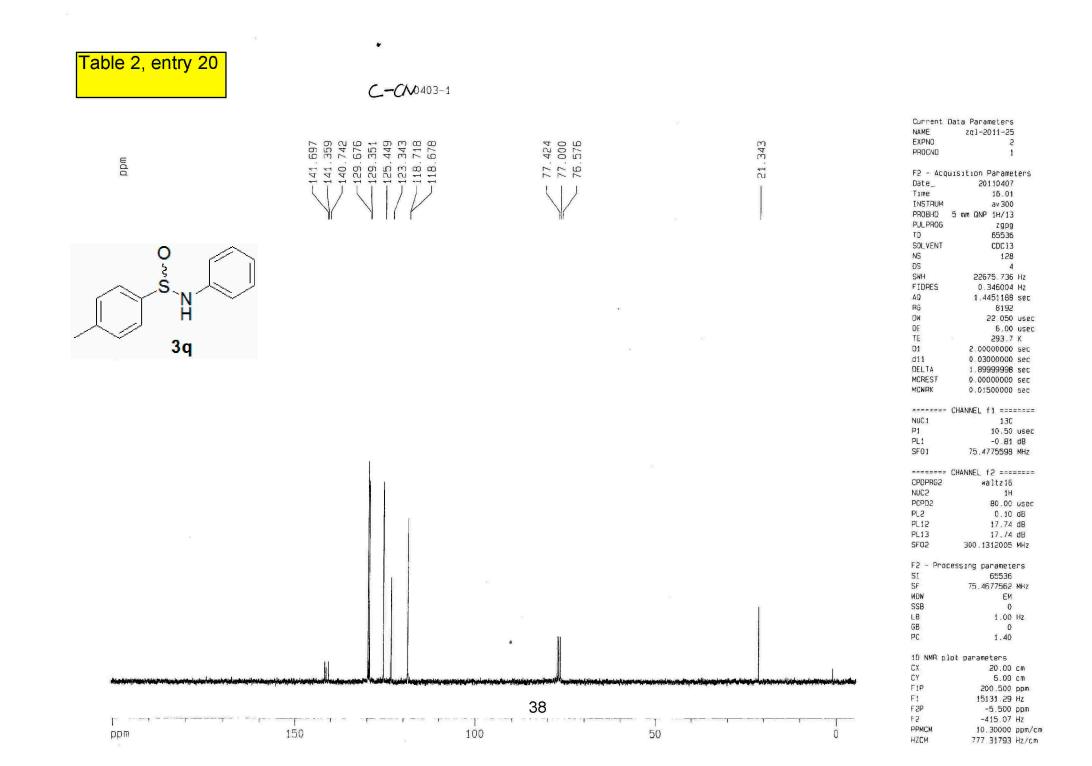


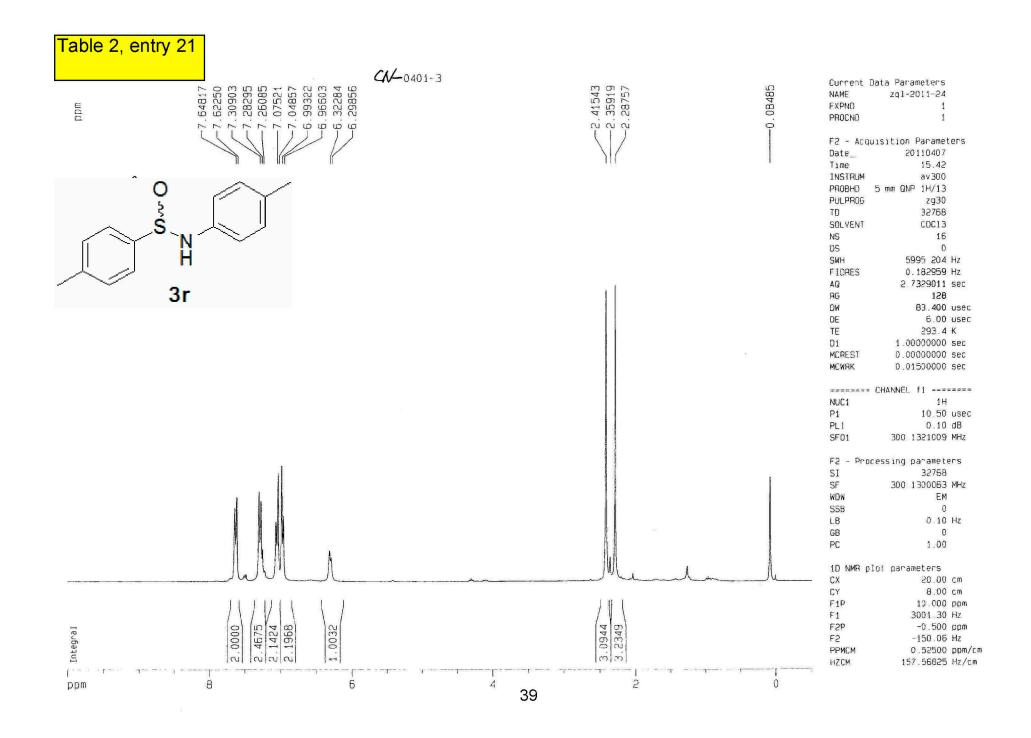
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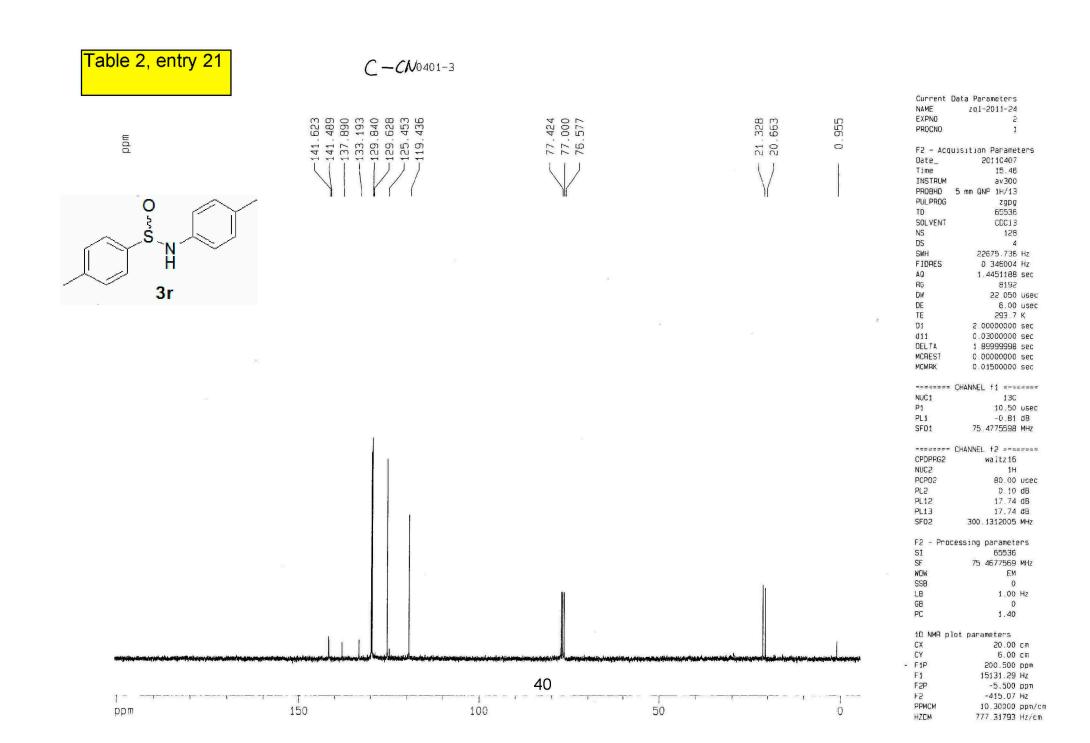
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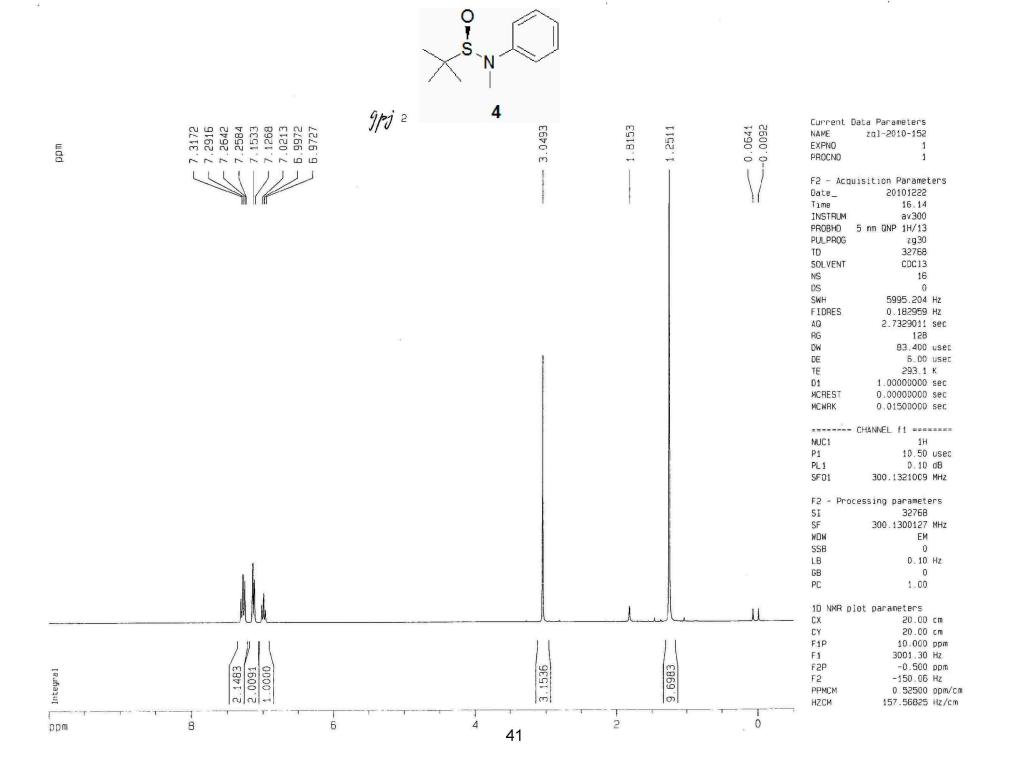


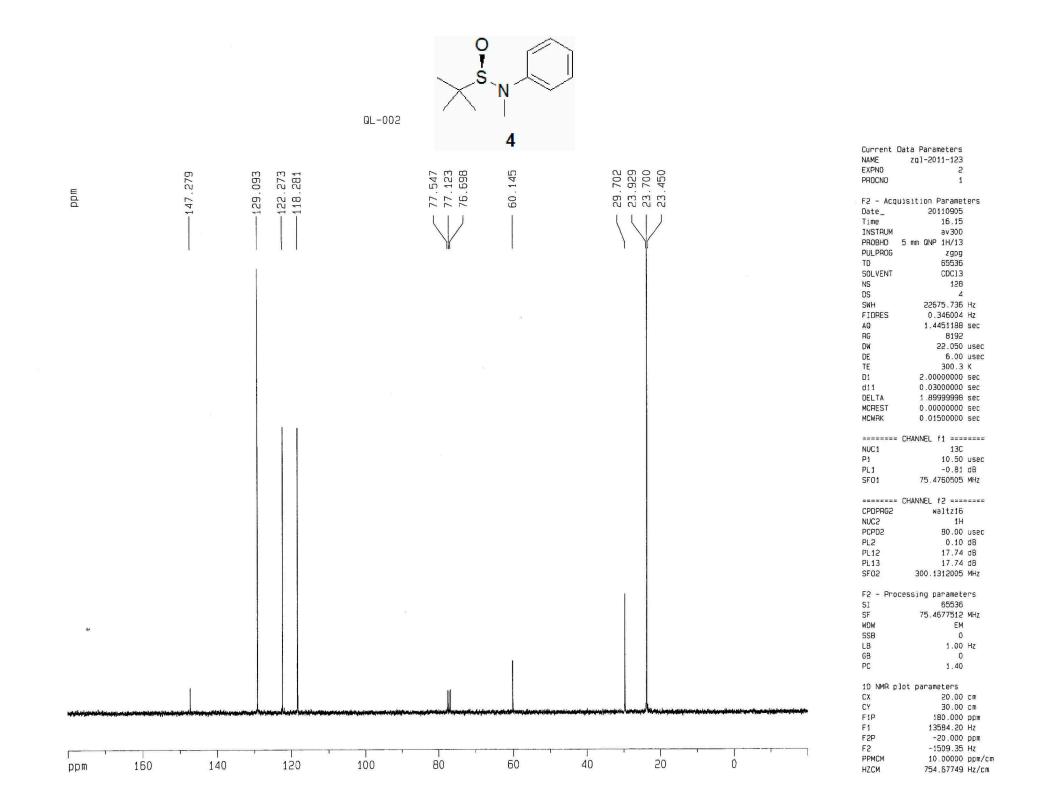


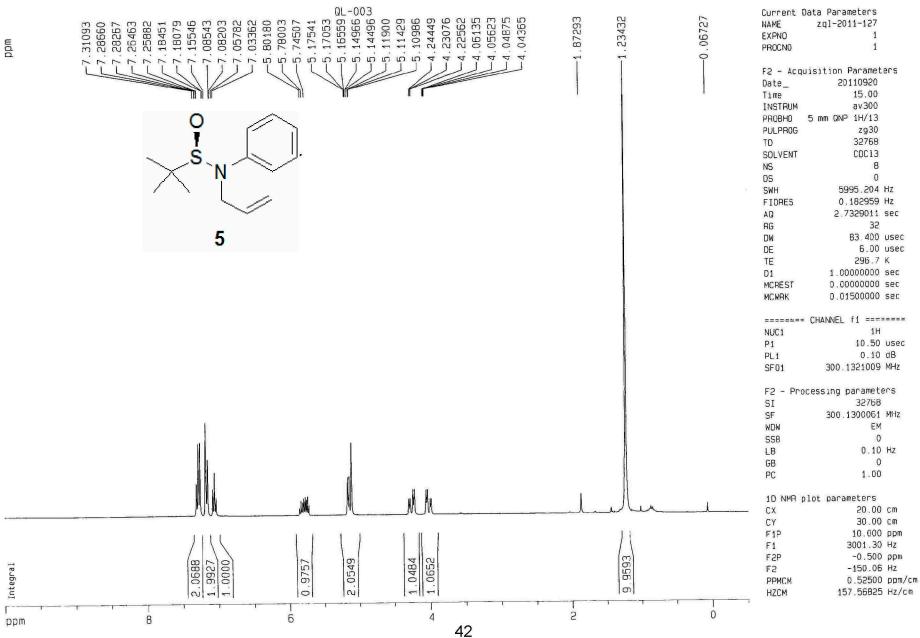


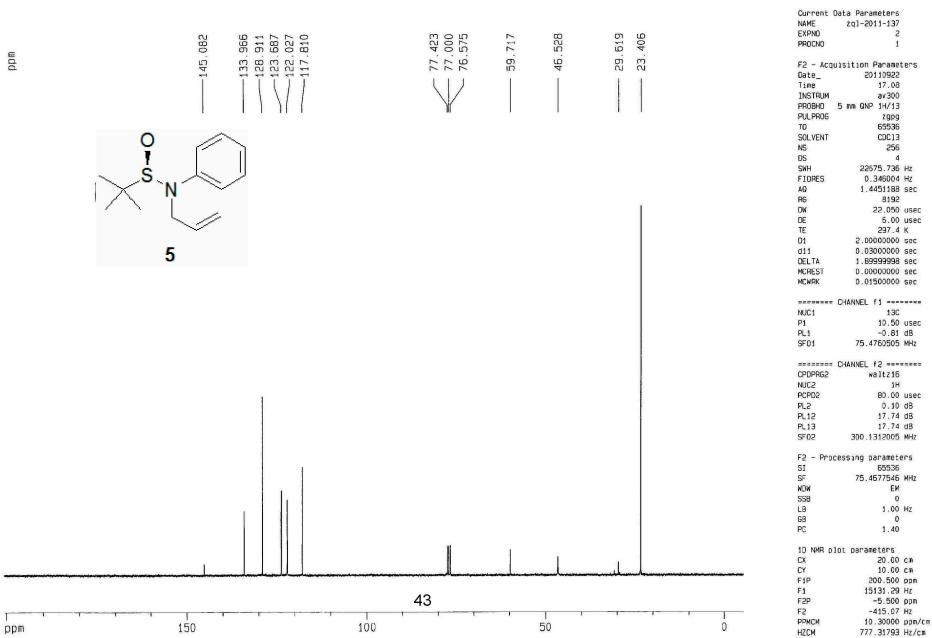












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