

SUPPLEMENTARY INFORMATION

Synthesis, characterisation, X-ray diffraction and biological evaluation of new thiourea derivatives against *Mycobacterium tuberculosis* and cervical cancer

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Table S1: Bond lengths [Å] and angles [°] for compound **TU1**.

Bond lengths [Å]		Bond angles [°]	
S(1)-C(2)	1.680(2)	C(2)-N(1)-C(3)	122.16(17)
O(1)-C(1)	1.216(2)	C(2)-N(1)-C(4)	120.54(16)
N(1)-C(2)	1.331(2)	C(3)-N(1)-C(4)	116.84(16)
N(1)-C(3)	1.465(3)	C(11)-C(10)-C(15)	119.88(18)
N(1)-C(4)	1.481(2)	C(11)-C(10)-C(1)	119.14(17)
C(10)-C(11)	1.393(3)	C(15)-C(10)-C(1)	120.96(17)
C(10)-C(15)	1.392(3)	N(1)-C(4)-C(5)	111.24(16)
C(10)-C(1)	1.494(3)	N(1)-C(4)-C(9)	110.87(16)
C(4)-C(5)	1.521(3)	C(5)-C(4)-C(9)	111.84(17)
C(4)-C(9)	1.522(3)	C(14)-C(15)-C(10)	119.74(19)
C(15)-C(14)	1.388(3)	C(13)-C(12)-C(11)	120.3(2)
C(12)-C(13)	1.383(3)	C(13)-C(14)-C(15)	120.2(2)
C(12)-C(11)	1.386(3)	C(12)-C(13)-C(14)	120.01(19)
C(14)-C(13)	1.385(3)	C(4)-C(5)-C(6)	110.26(18)
C(5)-C(6)	1.528(3)	C(12)-C(11)-C(10)	119.81(19)
C(9)-C(8)	1.534(3)	C(4)-C(9)-C(8)	110.17(17)
C(8)-C(7)	1.518(4)	C(7)-C(8)-C(9)	111.52(19)
C(6)-C(7)	1.519(4)	C(7)-C(6)-C(5)	111.2(2)
C(1)-N(2)	1.388(2)	C(8)-C(7)-C(6)	111.02(19)
C(2)-N(2)	1.401(2)	O(1)-C(1)-N(2)	123.78(18)

Table S2: Atomic coordinates ($\times 10^4$) and equivalent isotropic displacement parameters ($\text{Å}^2 \times 10^3$) for compound **TU1**. $U(\text{eq})$ is defined as one third of the trace of the orthogonalized U_{ij} tensor.

	x	y	z	$U(\text{eq})$
S(1)	1862(1)	3623(1)	4873(1)	28(1)
O(1)	1699(3)	4555(1)	6973(1)	31(1)
N(1)	4607(3)	3381(1)	6055(1)	26(1)
C(10)	1631(4)	6241(1)	6616(1)	23(1)
C(4)	4894(4)	2317(1)	5902(1)	25(1)
C(15)	3124(4)	6928(2)	6328(1)	28(1)
C(12)	-574(4)	7569(2)	7097(1)	33(1)
C(14)	2771(4)	7932(2)	6433(1)	33(1)
C(13)	915(4)	8252(2)	6813(1)	33(1)
C(5)	7238(4)	2124(2)	5569(1)	35(1)
C(11)	-205(4)	6564(2)	7009(1)	29(1)
C(9)	4724(4)	1692(2)	6543(1)	32(1)
C(8)	4991(5)	592(2)	6373(1)	39(1)
C(6)	7467(5)	1025(2)	5402(1)	43(1)
C(7)	7321(5)	392(2)	6038(1)	44(1)
C(1)	1999(3)	5156(1)	6524(1)	24(1)
C(2)	3135(3)	3947(1)	5649(1)	23(1)
C(3)	6232(4)	3794(2)	6613(1)	36(1)
N(2)	2704(3)	4906(1)	5882(1)	26(1)

Table S3: Selected bond lengths [Å] and angles [°] for compound **TU2**.

Bond lengths [Å]		Bond angles [°]	
S(1)-C(2)	1.6661(13)	C(1)-N(2)-C(2)	124.22(12)
O(1)-C(1)	1.214(18)	C(2)-N(1)-C(9)	120.30(11)
N(2)-C(1)	1.389(17)	C(4)-C(3)-C(8)	120.81(13)
N(2)-C(2)	1.395(16)	C(4)-C(3)-N(1)	120.31(13)
O(3)-N(3)	1.221(18)	C(8)-C(3)-N(1)	118.81(13)
N(1)-C(2)	1.349(17)	C(9)-C(14)-C(13)	119.18(15)
N(1)-C(9)	1.447(16)	N(1)-C(2)-N(2)	116.00(11)
N(1)-C(3)	1.448(17)	N(1)-C(2)-S(1)	123.60(10)
O(2)-N(3)	1.226(16)	N(2)-C(2)-S(1)	129.35(10)
N(3)-C(18)	1.474(17)	N(2)-C(1)-C(15)	114.92(12)
C(15)-C(20)	1.389(2)	C(17)-C(18)-C(19)	122.94(13)
C(15)-C(16)	1.397(18)	C(17)-C(18)-N(3)	118.78(12)
C(15)-C(1)	1.496(18)	C(19)-C(18)-N(3)	118.28(13)
C(20)-C(19)	1.389(19)	C(18)-C(19)-C(20)	118.43(14)
C(9)-C(10)	1.380(2)	C(18)-C(17)-C(16)	118.06(12)
C(9)-C(14)	1.387(2)	C(17)-C(16)-C(15)	120.32(14)
C(3)-C(4)	1.382(2)	C(9)-C(10)-C(11)	118.89(14)
C(3)-C(8)	1.386(2)	C(3)-C(4)-C(5)	119.29(15)
C(14)-C(13)	1.388(2)	C(12)-C(13)-C(14)	120.07(14)
C(18)-C(17)	1.380(2)	C(3)-C(8)-C(7)	119.17(16)
C(17)-C(16)	1.389(19)	C(13)-C(12)-C(11)	120.30(14)
C(10)-C(11)	1.392(2)	C(6)-C(7)-C(8)	120.44(15)
C(4)-C(5)	1.389(2)	C(6)-C(5)-C(4)	120.16(16)
C(13)-C(12)	1.379(3)	C(12)-C(11)-C(10)	120.28(16)
C(8)-C(7)	1.388(2)	C(7)-C(6)-C(5)	120.06(15)

Table S4: Atomic coordinates ($\times 10^4$) and equivalent isotropic displacement parameters ($\text{Å}^2 \times 10^3$) for compound **TU2**. U(eq) is defined as one third of the trace of the orthogonalized Uij tensor.

	x	y	z	U(eq)
S(1)	2408(1)	3652(1)	5240(1)	24(1)
O(1)	4979(2)	5989(1)	2647(1)	24(1)
N(2)	2106(2)	5091(1)	3587(1)	20(1)
O(3)	-3087(2)	9595(1)	1054(1)	37(1)
N(1)	4208(2)	3222(1)	3291(1)	20(1)
O(2)	-633(2)	10965(1)	923(1)	39(1)
N(3)	-1408(2)	9883(1)	1162(1)	26(1)
C(15)	1895(2)	7009(1)	2480(1)	18(1)
C(20)	-10(2)	6715(1)	2335(1)	20(1)
C(9)	5468(2)	2211(1)	3630(1)	20(1)
C(1)	3164(2)	6006(1)	2901(1)	19(1)
C(3)	4258(2)	3265(1)	2197(1)	21(1)
C(14)	5050(2)	880(1)	3509(1)	24(1)
C(2)	2976(2)	3991(1)	3985(1)	19(1)
C(18)	-229(2)	8874(1)	1603(1)	21(1)
C(19)	-1092(2)	7653(1)	1889(1)	22(1)
C(17)	1673(2)	9190(1)	1722(1)	23(1)
C(16)	2743(2)	8241(1)	2165(1)	22(1)
C(10)	7116(2)	2579(2)	4011(1)	28(1)
C(4)	2505(2)	3170(1)	1834(1)	26(1)
C(13)	6315(3)	-97(2)	3777(1)	29(1)
C(8)	6097(2)	3339(2)	1522(1)	28(1)

C(12)	7989(3)	260(2)	4138(1)	32(1)
C(7)	6161(3)	3349(2)	471(1)	33(1)
N				
C(11)	8387(3)	1588(2)	4262(1)	34(1)
C(6)	4418(3)	3308(2)	103(1)	33(1)

Table S5: Selected bond lengths [Å] and angles [°] for compound **TU6**.

Bond lengths [Å]		Bond angles [°]	
S(1)-C(2)	1.677(11)	C(2)-N(1)-C(3)	122.65(10)
O(1)-C(1)	1.218(14)	C(2)-N(1)-C(4)	120.49(9)
N(1)-C(2)	1.328(14)	C(3)-N(1)-C(4)	116.71(9)
N(1)-C(3)	1.468(14)	C(1)-N(2)-C(2)	124.91(10)
N(1)-C(4)	1.480(14)	O(1)-C(1)-N(2)	121.75(11)
N(2)-C(1)	1.385(14)	O(1)-C(1)-C(10)	123.51(10)
N(2)-C(2)	1.405(14)	N(2)-C(1)-C(10)	144.72(9)
C(1)-C(10)	1.494(16)	C(11)-C(10)-C(15)	119.96(11)
C(10)-C(11)	1.401(16)	C(11)-C(10)-C(1)	119.63(10)
C(10)-C(15)	1.410(15)	C(15)-C(10)-C(1)	120.29(10)
C(4)-C(9)	1.522(16)	N(1)-C(4)-C(1)	110.69(9)
C(4)-C(5)	1.527(15)	N(1)-C(4)-C(9)	111.27(9)
C(11)-C(12)	1.385(18)	C(9)-C(4)-C(5)	111.96(10)
C(15)-C(14)	1.393(17)	N(1)-C(2)-N(2)	116.51(10)
C(15)-C(16)	1.508(17)	N(1)-C(2)-S(1)	125.90(9)
C(9)-C(8)	1.528(16)	N(2)-C(2)-S(1)	117.58(8)
C(14)-C(13)	1.386(19)	C(12)-C(11)-C(10)	120.97(11)
C(8)-C(7)	1.522(18)	C(14)-C(15)-C(10)	117.68(11)
C(5)-C(6)	1.529(17)	C(14)-C(15)-C(16)	119.08(11)
C(7)-C(6)	1.523(19)	C(10)-C(15)-C(16)	123.24(11)
C(12)-C(13)	1.382(19)	C(4)-C(9)-C(8)	109.97(9)

Table S6: Atomic coordinates ($\times 10^4$) and equivalent isotropic displacement parameters ($\text{Å}^2 \times 10^3$) for compound **TU6**. $U(\text{eq})$ is defined as one third of the trace of the orthogonalized U_{ij} tensor.

	x	y	z	U(eq)
S(1)	7673(1)	4869(1)	5812(1)	26(1)
O(1)	6419(1)	3147(1)	6102(1)	31(1)
N(1)	6344(1)	4018(1)	3660(1)	22(1)
N(2)	5336(1)	4060(1)	5776(1)	23(1)
C(1)	5431(2)	3504(1)	6340(1)	22(1)
C(10)	4201(2)	3379(1)	7208(1)	21(1)
C(4)	7598(2)	4179(1)	2864(1)	22(1)
C(2)	6428(2)	4285(1)	5000(1)	21(1)
C(11)	2542(2)	3673(1)	6774(1)	26(1)
C(15)	4648(2)	2942(1)	8389(1)	23(1)
C(9)	8462(2)	3649(1)	2469(1)	25(1)
C(14)	3374(2)	2811(1)	9057(2)	30(1)
C(16)	6430(2)	2619(1)	8964(2)	30(1)
C(3)	4948(2)	3587(1)	2833(2)	30(1)
C(8)	9790(2)	3824(1)	1689(2)	28(1)
C(5)	6633(2)	4549(1)	1365(2)	30(1)
C(7)	8844(2)	4188(1)	183(2)	33(1)
C(6)	7966(2)	4715(1)	580(2)	33(1)

C(12)	1297(2)	3530(1)	7450(2)	32(1)
C(13)	1714(2)	3094(1)	8584(2)	33(1)

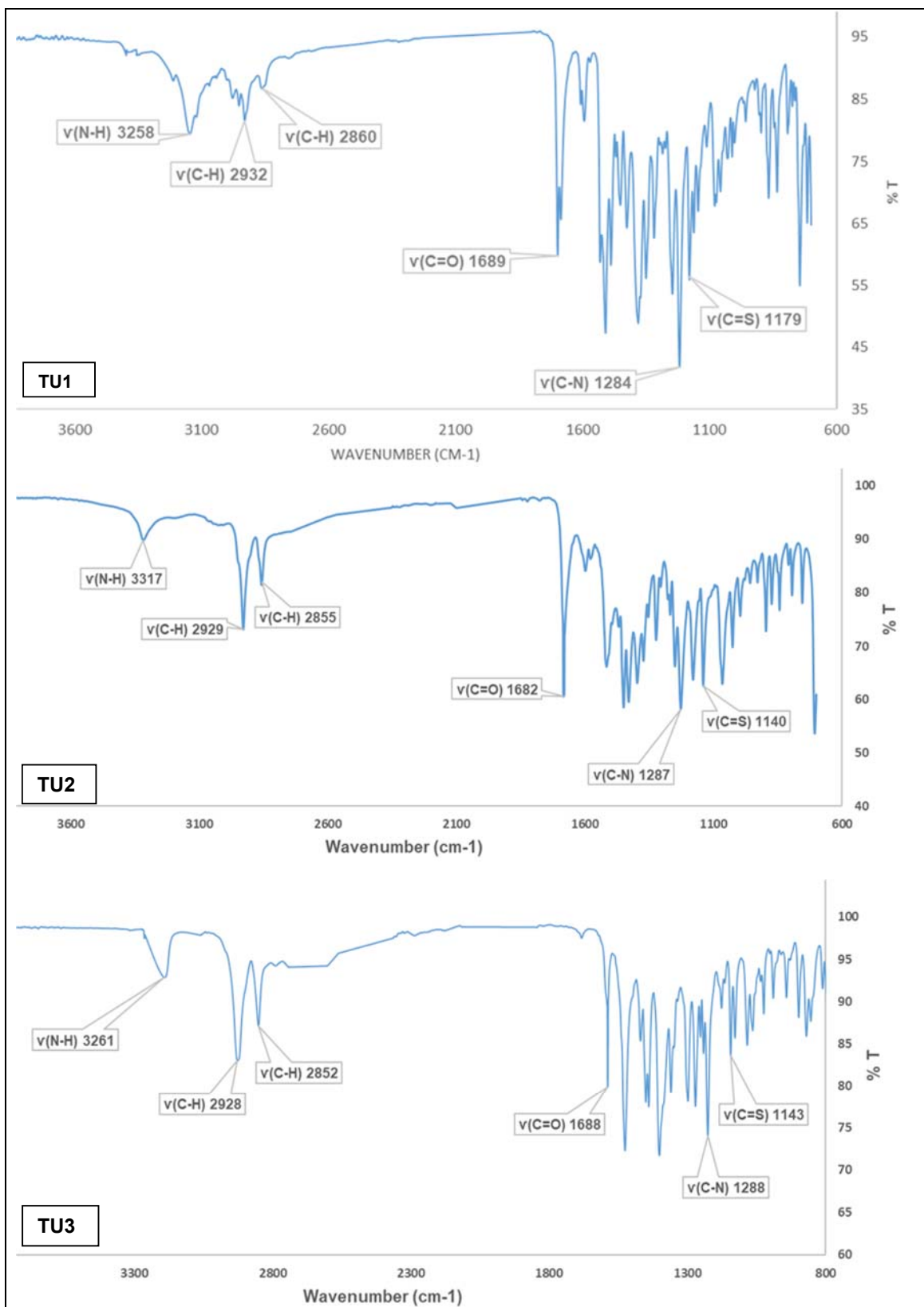


Figure S1: IR spectra for compound TU1 - TU3

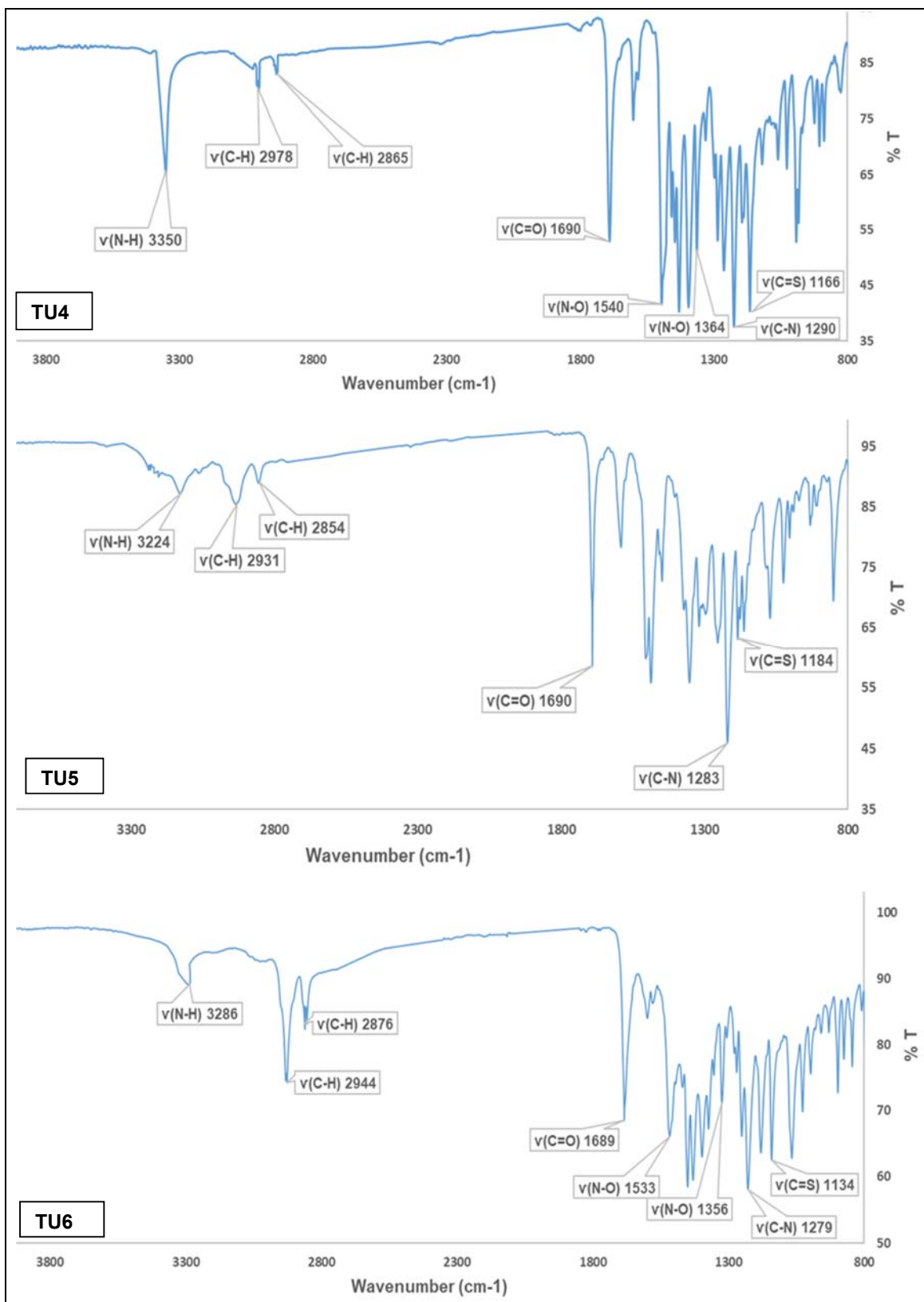


Figure S2: IR spectra for compound TU4 - TU6

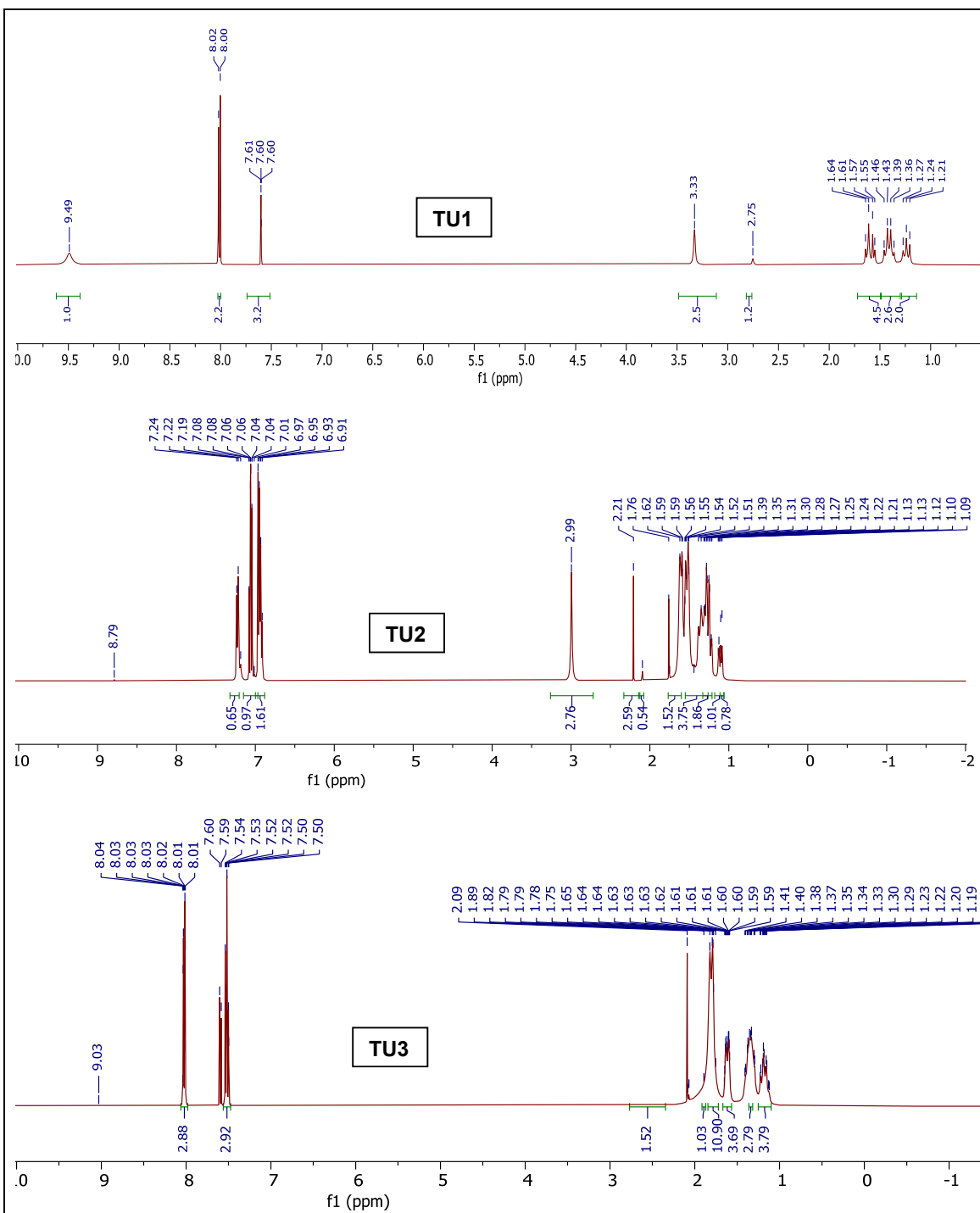


Figure S3: ^1H NMR for compounds TU1 - TU3

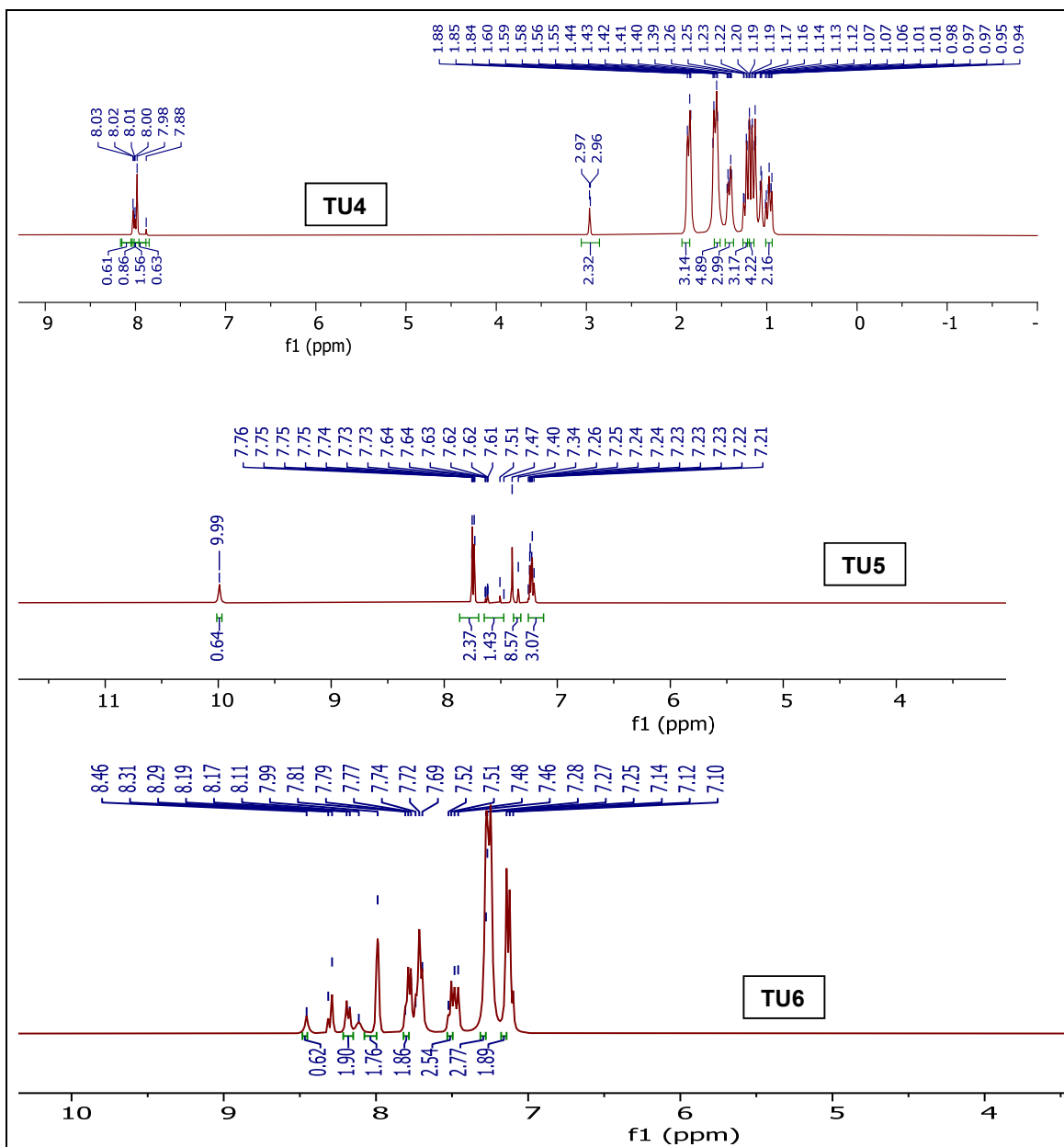


Figure S4: ^1H NMR for compounds TU4 - TU6

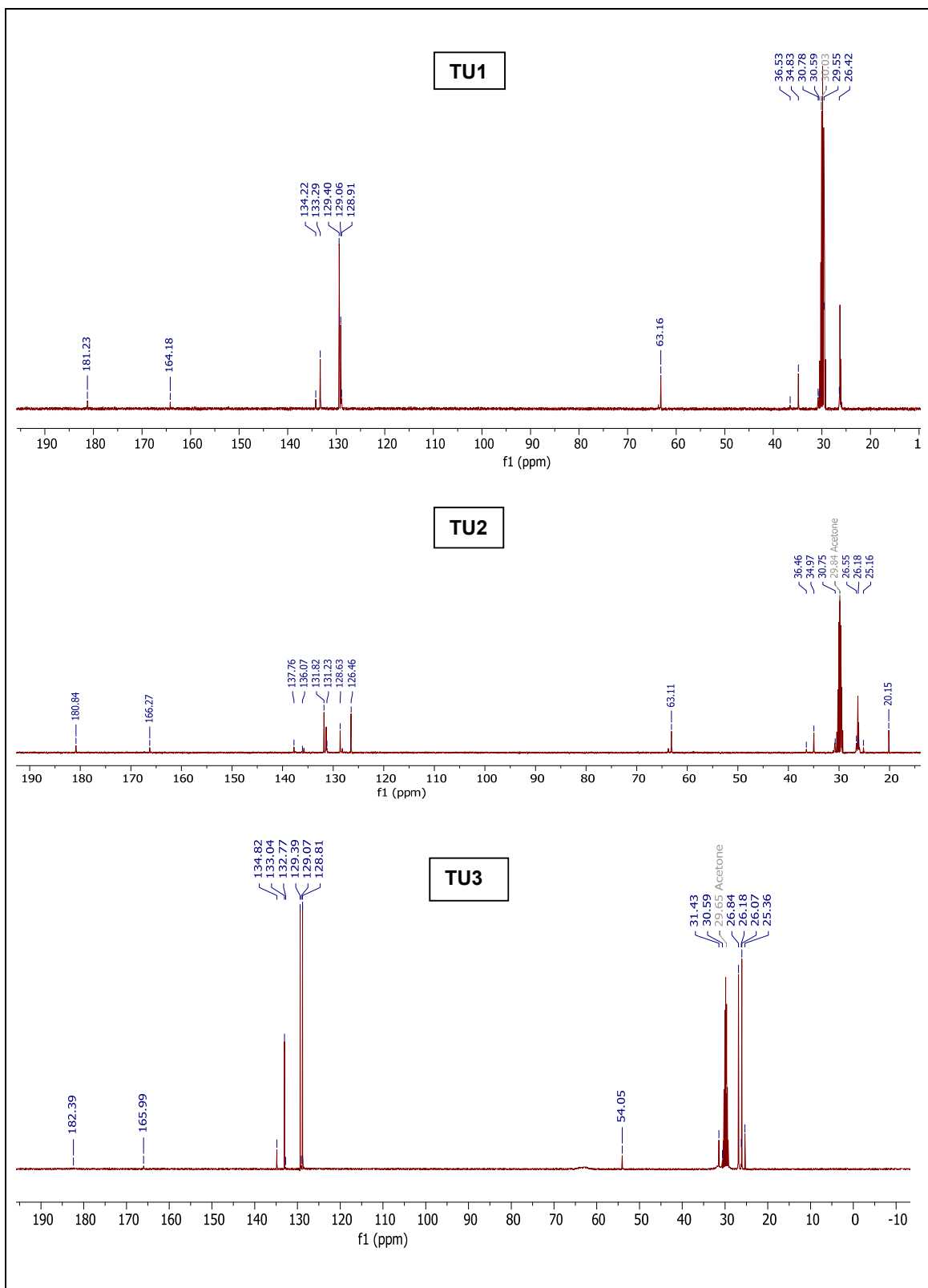


Figure S5: ^{13}C NMR for compound **TU1** - **TU3**

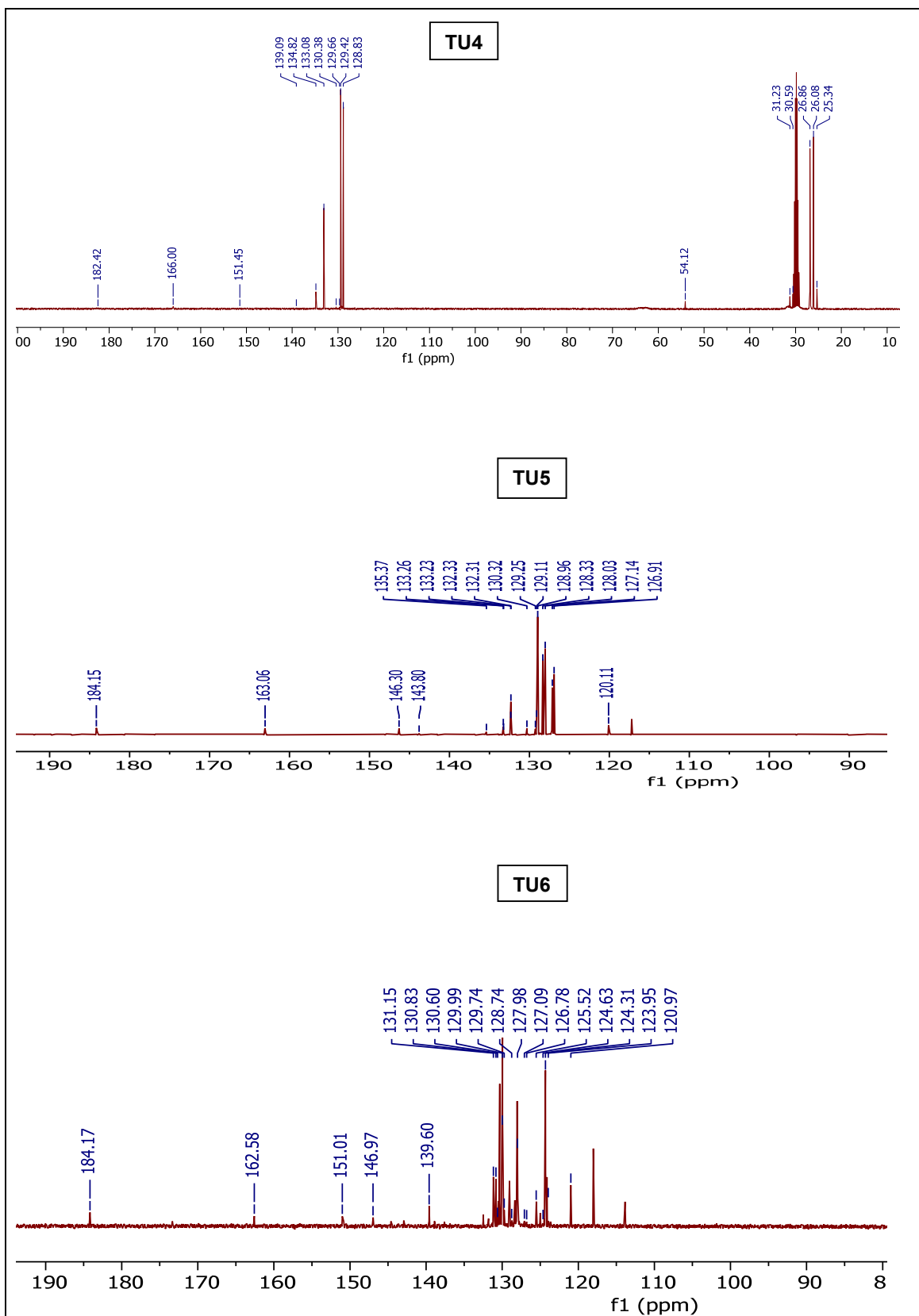


Figure S6: ^{13}C NMR for compound TU4 - TU6

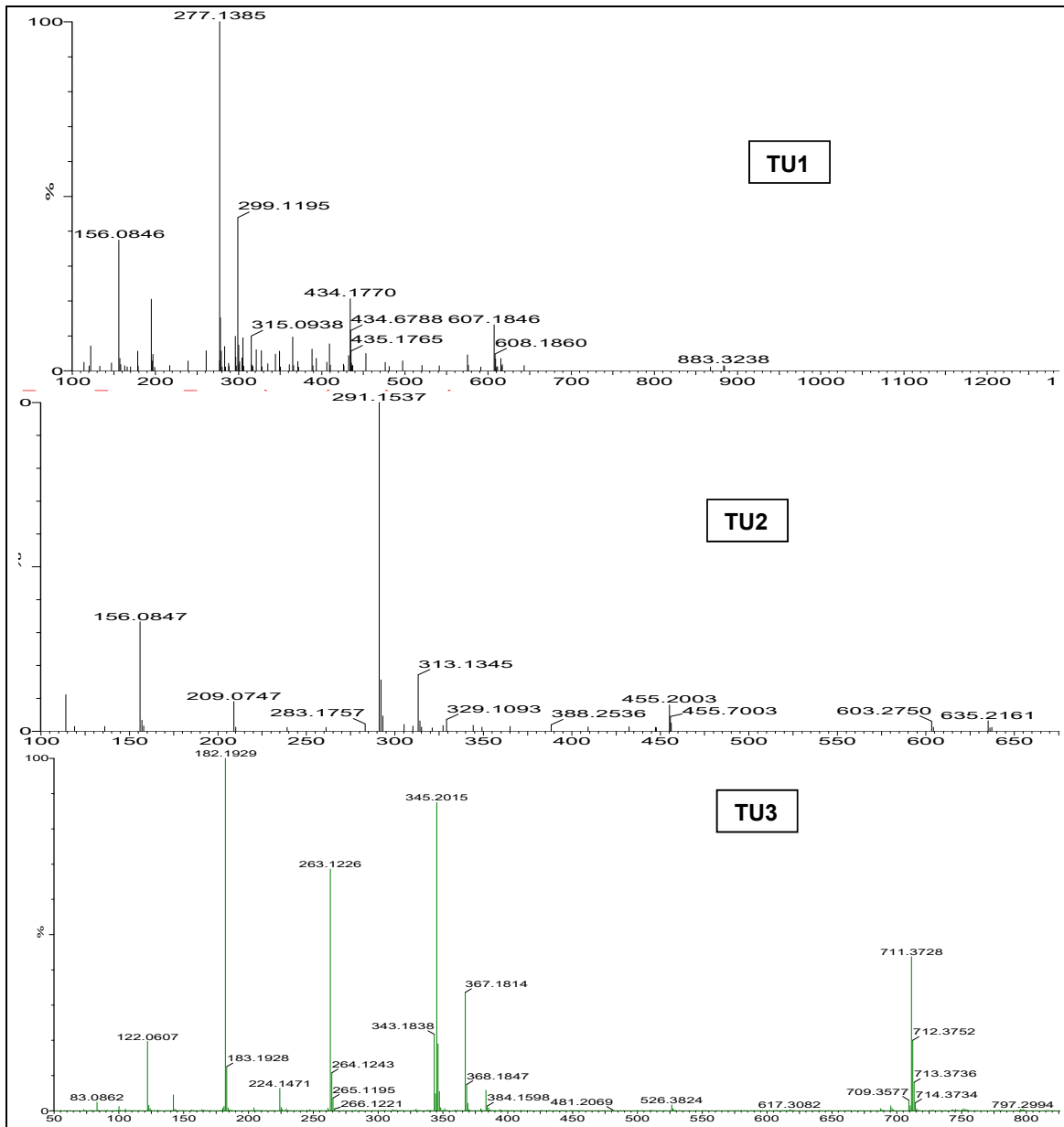


Figure S7: MS for compound TU1 - TU3.

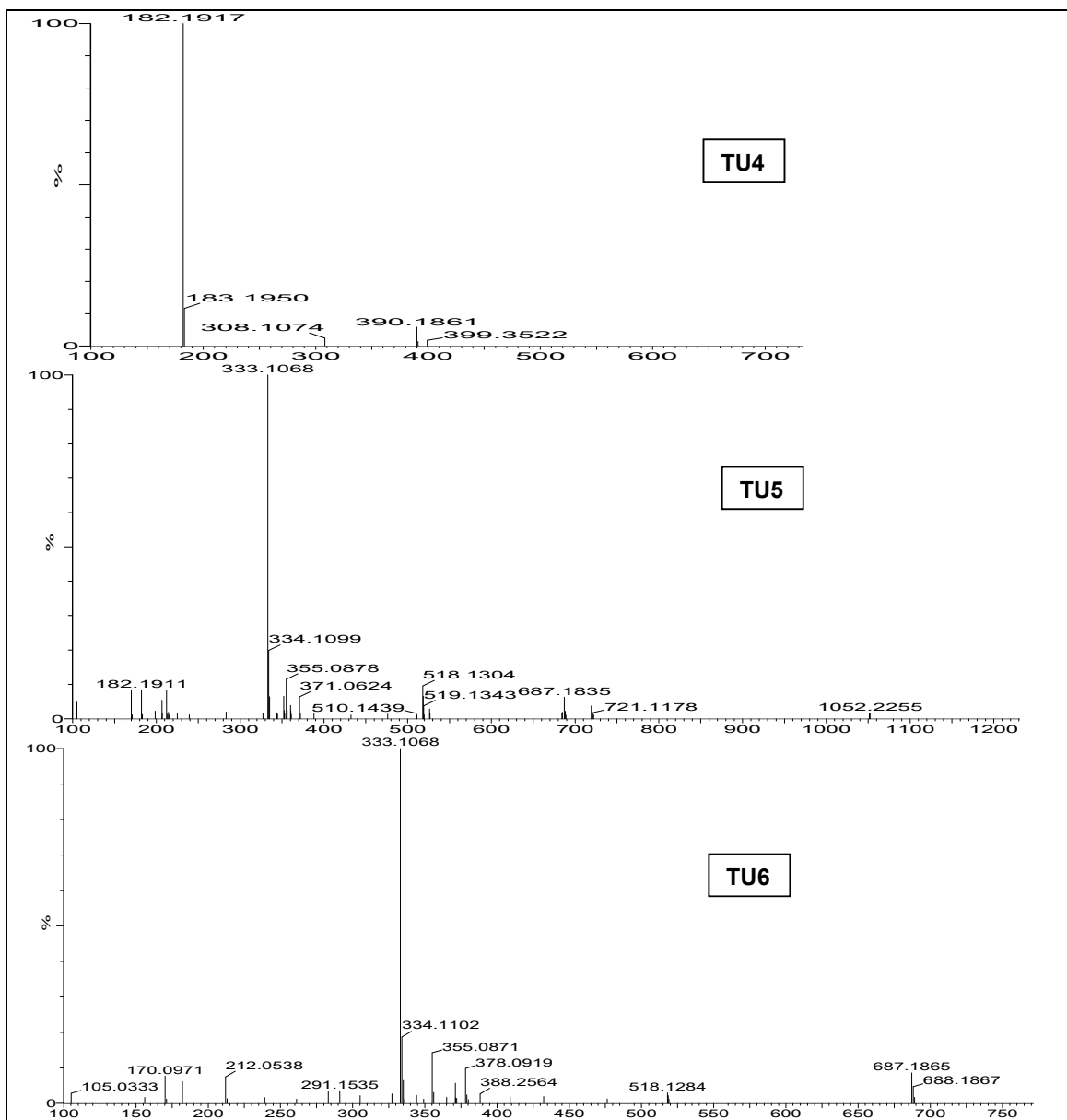


Figure S8: MS for compound TU4 - TU6.

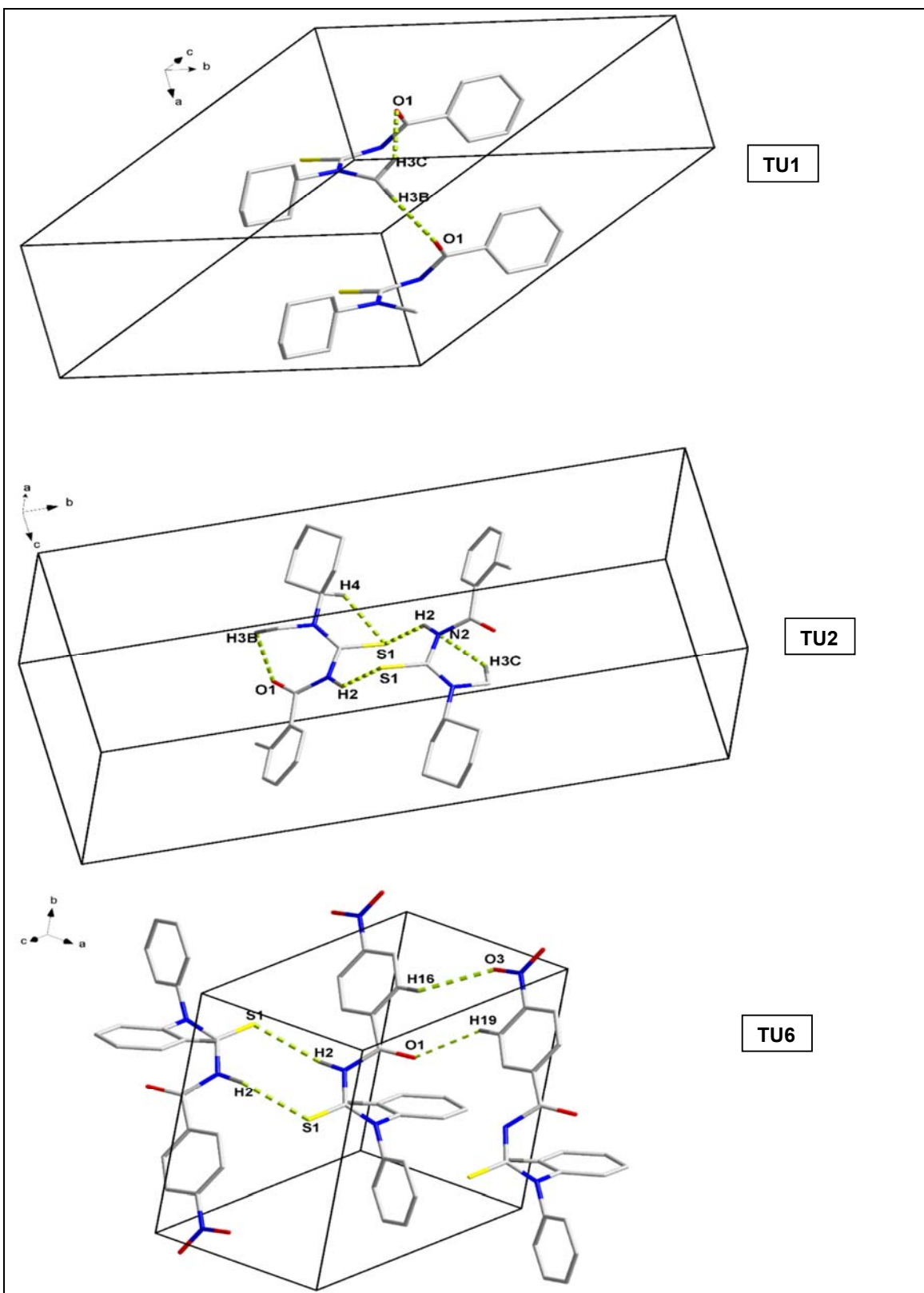


Figure S9: Hydrogen bond interactions (indicated in green dashed lines) for compound **TU1**, **TU2** and **TU6**.

N-(cyclohexyl(methyl)carbomothioyl) benzamide for (TU1)

checkCIF/PLATON report

Structure factors have been supplied for datablock(s) am105_lt_auto

THIS REPORT IS FOR GUIDANCE ONLY. IF USED AS PART OF A REVIEW PROCEDURE FOR PUBLICATION, IT SHOULD NOT REPLACE THE EXPERTISE OF AN EXPERIENCED CRYSTALLOGRAPHIC REFEREE.

No syntax errors found. CIF dictionary Interpreting this report

Datablock: am105_lt_auto

Bond precision: C-C = 0.0022 Å Wavelength=1.54184
Cell: a=5.5312 (1) b=13.5140 (2) c=19.5709 (3)
alpha=90 beta=94.757 (1) gamma=90
Temperature: 150 K

	Calculated	Reported
Volume	1457.86 (4)	1457.86 (4)
Space group	P 21/c	P 1 21/c 1
Hall group	-P 2ybc	-P 2ybc
Moiety formula	C15 H20 N2 O S	C15 H20 N2 O S
Sum formula	C15 H20 N2 O S	C15 H20 N2 O S
Mr	276.39	276.39
Dx, g cm ⁻³	1.259	1.255
Z	4	4
Mu (mm ⁻¹)	1.916	1.915
F000	592.0	588.0
F000'	594.77	
h, k, lmax	7, 17, 24	7, 16, 24
Nref	3136	3043
Tmin, Tmax	0.672, 0.713	0.387, 1.000
Tmin'	0.610	

Correction method= # Reported T Limits: Tmin=0.387 Tmax=1.000
AbsCorr = MULTI-SCAN

Data completeness= 0.970 Theta(max)= 78.810

R(reflections)= 0.0391(2754) wR2(reflections)=
0.0981(3043)
S = 1.059 Npar= 173

The following ALERTS were generated. Each ALERT has the format
test-name_ALERT_alert-type_alert-level.
Click on the hyperlinks for more details of the test.

● **Alert level C**

PLAT068_ALERT_1_C	Reported F000 Differs from Calcd (or Missing)...	Please Check
PLAT230_ALERT_2_C	Hirshfeld Test Diff for S1 --C2 .	6.0 s.u.
PLAT420_ALERT_2_C	D-H Bond Without Acceptor N2 --H2 .	Please Check
PLAT906_ALERT_3_C	Large K Value in the Analysis of Variance	2.152 Check
PLAT911_ALERT_3_C	Missing FCF Refl Between Thmin & STh/L= 0.600	4 Report
PLAT976_ALERT_2_C	Check Calcd Resid. Dens. 0.98Ang From N2 .	-0.52 eA-3
PLAT977_ALERT_2_C	Check Negative Difference Density on H2 .	-0.38 eA-3

● **Alert level G**

PLAT007_ALERT_5_G	Number of Unrefined Donor-H Atoms	1 Report
PLAT912_ALERT_4_G	Missing # of FCF Reflections Above STh/L= 0.600	76 Note
PLAT933_ALERT_2_G	Number of HKL-OMIT Records in Embedded .res File	5 Note
PLAT953_ALERT_1_G	Reported (CIF) and Actual (FCF) Hmax Differ by .	1 Units
PLAT978_ALERT_2_G	Number C-C Bonds with Positive Residual Density.	10 Info

- 0 **ALERT level A** = Most likely a serious problem - resolve or explain
0 **ALERT level B** = A potentially serious problem, consider carefully
7 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight
5 **ALERT level G** = General information/check it is not something unexpected
- 2 ALERT type 1 CIF construction/syntax error, inconsistent or missing data
6 ALERT type 2 Indicator that the structure model may be wrong or deficient
2 ALERT type 3 Indicator that the structure quality may be low
1 ALERT type 4 Improvement, methodology, query or suggestion
1 ALERT type 5 Informative message, check
-

It is advisable to attempt to resolve as many as possible of the alerts in all categories. Often the minor alerts point to easily fixed oversights, errors and omissions in your CIF or refinement strategy, so attention to these fine details can be worthwhile. In order to resolve some of the more serious problems it may be necessary to carry out additional measurements or structure refinements. However, the purpose of your study may justify the reported deviations and the more serious of these should normally be commented upon in the discussion or experimental section of a paper or in the "special_details" fields of the CIF. checkCIF was carefully designed to identify outliers and unusual parameters, but every test has its limitations and alerts that are not important in a particular case may appear. Conversely, the absence of alerts does not guarantee there are no aspects of the results needing attention. It is up to the individual to critically assess their own results and, if necessary, seek expert advice.

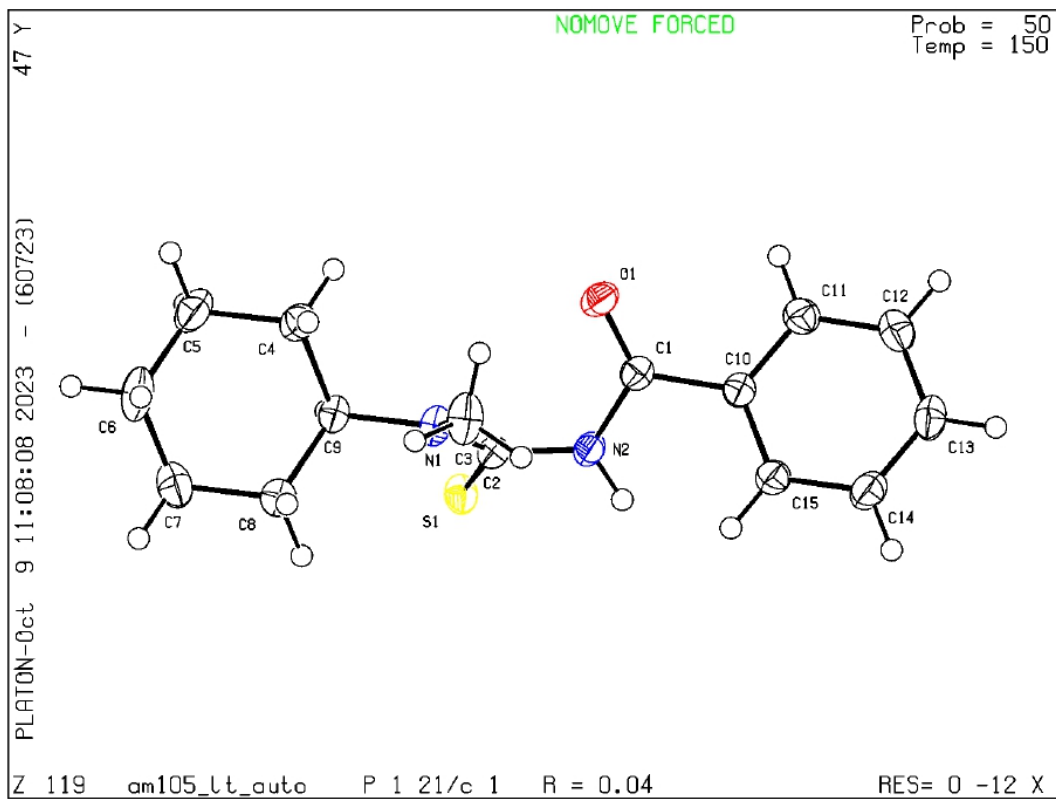
Publication of your CIF in IUCr journals

A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section C* or *E* or *IUCrData*, you should make sure that [full publication checks](#) are run on the final version of your CIF prior to submission.

Publication of your CIF in other journals

Please refer to the *Notes for Authors* of the relevant journal for any special instructions relating to CIF submission.

PLATON version of 06/07/2023; check.def file version of 30/06/2023



The following ALERTS were generated. Each ALERT has the format
test-name_ALERT_alert-type_alert-level.
Click on the hyperlinks for more details of the test.

● **Alert level C**

PLAT230_ALERT_2_C	Hirshfeld Test Diff for	S1	--C2	.	5.2 s.u.
PLAT414_ALERT_2_C	Short Intra D-H..H-X	H2	..H11	.	1.97 Ang.
			x, y, z =	1_555	Check
PLAT976_ALERT_2_C	Check Calcd Resid. Dens.	0.96Ang	From N2	.	-0.41 eA-3

● **Alert level G**

PLAT007_ALERT_5_G	Number of Unrefined Donor-H Atoms	1	Report
PLAT883_ALERT_1_G	No Info/Value for _atom_sites_solution_primary .		Please Do !
PLAT899_ALERT_4_G	SHELXL2018 is Deprecated and Succeeded by SHELXL	2019/3	Note
PLAT910_ALERT_3_G	Missing # of FCF Reflection(s) Below Theta(Min).	1	Note
PLAT912_ALERT_4_G	Missing # of FCF Reflections Above STh/L= 0.600	698	Note
PLAT978_ALERT_2_G	Number C-C Bonds with Positive Residual Density.	13	Info

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1 ALERT type 5 Informative message, check

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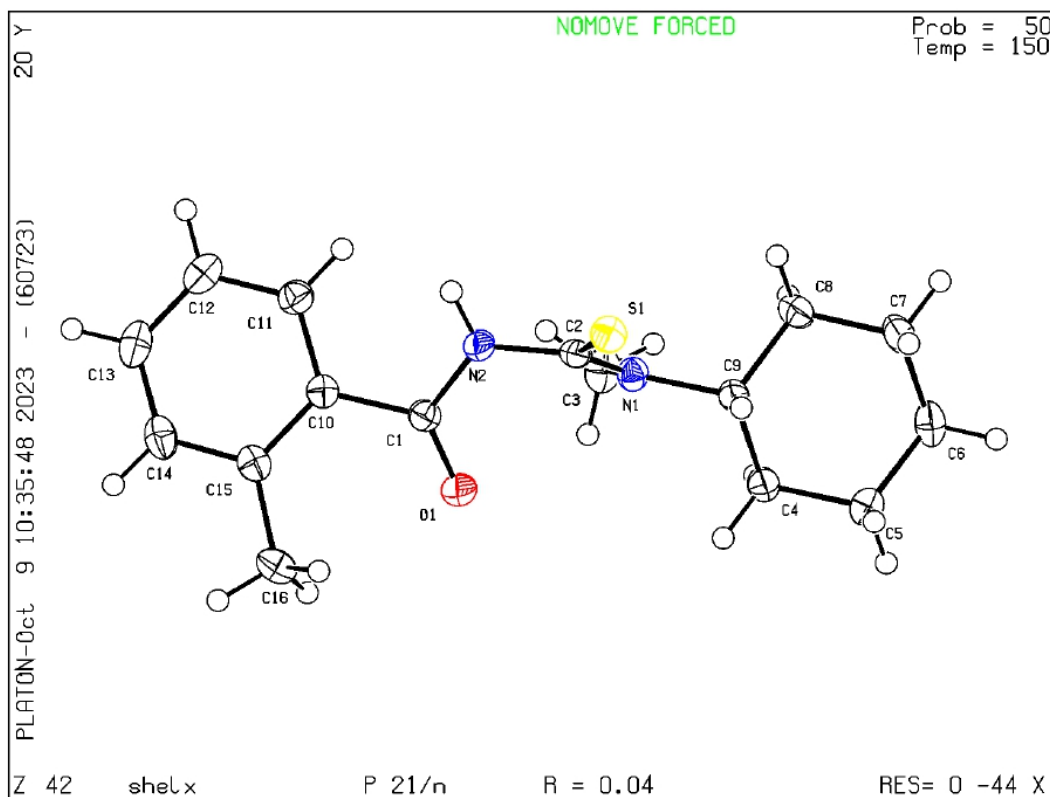
Publication of your CIF in IUCr journals

A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section C* or *E* or *IUCrData*, you should make sure that full publication checks are run on the final version of your CIF prior to submission.

Publication of your CIF in other journals

Please refer to the *Notes for Authors* of the relevant journal for any special instructions relating to CIF submission.

PLATON version of 06/07/2023; check.def file version of 30/06/2023



N-(diphenylcarbamothioyl)-4-nitrobenzamide (TU6)

checkCIF/PLATON report

Structure factors have been supplied for datablock(s) shelx

THIS REPORT IS FOR GUIDANCE ONLY. IF USED AS PART OF A REVIEW PROCEDURE FOR PUBLICATION, IT SHOULD NOT REPLACE THE EXPERTISE OF AN EXPERIENCED CRYSTALLOGRAPHIC REFEREE.

No syntax errors found. CIF dictionary Interpreting this report

Datablock: shelx

Bond precision: C-C = 0.0021 A Wavelength=0.71073
Cell: a=6.8186 (2) b=10.0318 (2) c=13.2816 (3)
alpha=89.529 (2) beta=78.134 (2) gamma=89.102 (2)
Temperature: 150 K

	Calculated	Reported
Volume	888.97 (4)	888.96 (4)
Space group	P -1	P -1
Hall group	-P 1	-P 1
Moiety formula	C20 H15 N3 O3 S	C20 H15 N3 O3 S
Sum formula	C20 H15 N3 O3 S	C20 H15 N3 O3 S
Mr	377.41	377.41
Dx, g cm ⁻³	1.410	1.406
Z	2	2
Mu (mm ⁻¹)	0.209	0.209
F000	392.0	390.0
F000'	392.41	
h, k, lmax	9, 14, 19	8, 13, 17
Nref	5649	4318
Tmin, Tmax	0.951, 0.962	0.664, 1.000
Tmin'	0.951	

Correction method= # Reported T Limits: Tmin=0.664 Tmax=1.000
AbsCorr = MULTI-SCAN

Data completeness= 0.764 Theta(max)= 30.941

R(reflections)= 0.0389 (3503) wR2 (reflections)=
0.1002 (4318)
S = 1.072 Npar= 244

The following ALERTS were generated. Each ALERT has the format
test-name_ALERT_alert-type_alert-level.
Click on the hyperlinks for more details of the test.

Alert level C

PLAT068_ALERT_1_C Reported F000 Differs from Calcd (or Missing)...	Please Check
PLAT911_ALERT_3_C Missing FCF Refl Between Thmin & STh/L= 0.600	40 Report

Alert level G

PLAT007_ALERT_5_G Number of Unrefined Donor-H Atoms	1 Report
PLAT154_ALERT_1_G The s.u.'s on the Cell Angles are Equal ..(Note)	0.002 Degree
PLAT883_ALERT_1_G No Info/Value for _atom_sites_solution_primary .	Please Do !
PLAT899_ALERT_4_G SHELXL2018 is Deprecated and Succeeded by SHELXL	2019/3 Note
PLAT910_ALERT_3_G Missing # of FCF Reflection(s) Below Theta(Min).	2 Note
PLAT912_ALERT_4_G Missing # of FCF Reflections Above STh/L= 0.600	1037 Note
PLAT941_ALERT_3_G Average HKL Measurement Multiplicity	2.5 Low
PLAT952_ALERT_5_G Calculated (ThMax) and CIF-Reported Lmax Differ.	2 Units
PLAT958_ALERT_1_G Calculated (ThMax) and Actual (FCF) Lmax Differ.	2 Units
PLAT978_ALERT_2_G Number C-C Bonds with Positive Residual Density.	11 Info

0 **ALERT level A** = Most likely a serious problem - resolve or explain
0 **ALERT level B** = A potentially serious problem, consider carefully
2 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight
10 **ALERT level G** = General information/check it is not something unexpected

4 ALERT type 1 CIF construction/syntax error, inconsistent or missing data
1 ALERT type 2 Indicator that the structure model may be wrong or deficient
3 ALERT type 3 Indicator that the structure quality may be low
2 ALERT type 4 Improvement, methodology, query or suggestion
2 ALERT type 5 Informative message, check

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