

Supplementary material for publication on line

Influence of complexation of thiosemicarbazone derivatives with Cu (II) ions on their antitumor activity against melanoma cells

Monika Pitucha^{1,*}, Agnieszka Korga-Plewko², Agnieszka Czylkowska³, Bartłomiej Rogalewicz³, Monika Drozd¹ Magdalena Iwan⁴, Joanna Kubik², Ewelina Humeniuk², Grzegorz Adamczuk², Zbigniew Karczmarzyk⁵, Emilia Fornal⁶, Waldemar Wysocki⁵, Paulina Bartnik⁵

¹ Independent Radiopharmacy Unit, Faculty of Pharmacy, Medical University of Lublin, PL-20093 Lublin , Poland

² Independent Medical Biology Unit, Faculty of Pharmacy, Medical University of Lublin, PL-20093 Lublin, Poland; agnieszkakorga@umlub.pl

³ Institute of General and Ecological Chemistry, Faculty of Chemistry, Lodz University of Technology, Zeromskiego 116, 90-924 Lodz, Poland; 211150@edu.p.lodz.pl

⁴ Chair and Department of Toxicology, Faculty of Pharmacy, Medical University of Lublin, PL-20093 Lublin, Poland; magda.iwan@umlub.pl

⁵ Faculty of Science, Siedlce University of Natural Sciences and Humanities, 3 Maja 54, 08-110 Siedlce, Poland

⁶ Chair and Department of Pathophysiology, Faculty of Medicine, Medical University of Lublin, PL-20090 Lublin

* Correspondence: monika.pitucha@umlub.pl; Tel.: 48-81-448-72-40 (MP)

X-ray structure determination

CIF for T2

data_shelx

```
_audit_creation_method      'SHELXL-2014/7'  
_shelx_SHELXL_version_number  '2014/7'  
_chemical_name_systematic    ?  
_chemical_name_common        ?  
_chemical_melting_point      ?  
_chemical_formula_moiety     'C9 H10 Br N3 S'  
_chemical_formula_sum        'C9 H10 Br N3 S'  
_chemical_formula_weight     272.17
```

```
loop_  
_atom_type_symbol  
_atom_type_description  
_atom_type_scat_dispersion_real  
_atom_type_scat_dispersion_imag  
_atom_type_scat_source  
'C' 'C' 0.0033 0.0016  
'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'  
'H' 'H' 0.0000 0.0000
```

'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'

'N' 'N' 0.0061 0.0033

'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'

'S' 'S' 0.1246 0.1234

'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'

'Br' 'Br' -0.2901 2.4595

'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'

_space_group_crystal_system orthorhombic
_space_group_IT_number 61
_space_group_name_H-M_alt 'P b c a'
_space_group_name_Hall '-P 2ac 2ab'

_shelx_space_group_comment

;

The symmetry employed for this shelxl refinement is uniquely defined by the following loop, which should always be used as a source of symmetry information in preference to the above space-group names.

They are only intended as comments.

;

loop_

_space_group_symop_operation_xyz
'x, y, z'
'-x+1/2, -y, z+1/2'
'-x, y+1/2, -z+1/2'
'x+1/2, -y+1/2, -z'
'-x, -y, -z'
'x-1/2, y, -z-1/2'
'x, -y-1/2, z-1/2'
'-x-1/2, y-1/2, z'

_cell_length_a 13.5800(15)
_cell_length_b 8.5494(8)
_cell_length_c 19.0926(18)
_cell_angle_alpha 90
_cell_angle_beta 90
_cell_angle_gamma 90
_cell_volume 2216.7(4)
_cell_formula_units_Z 8
_cell_measurement_temperature 296(2)
_cell_measurement_reflns_used 2646
_cell_measurement_theta_min 2.6100
_cell_measurement_theta_max 26.1330

_exptl_crystal_description plate
_exptl_crystal_colour colourless
_exptl_crystal_density_meas ?
_exptl_crystal_density_method ?
_exptl_crystal_density_diffrn 1.631
_exptl_crystal_F_000 1088
_exptl_transmission_factor_min ?
_exptl_transmission_factor_max ?
_exptl_crystal_size_max 0.5
_exptl_crystal_size_mid 0.5
_exptl_crystal_size_min 0.1
_exptl_absorpt_coefficient_mu 3.862

```

_shelx_estimated_absorpt_T_min  ?
_shelx_estimated_absorpt_T_max  ?
_exptl_absorpt_correction_T_min 0.00545
_exptl_absorpt_correction_T_max 1.00000
_exptl_absorpt_correction_type multi-scan
_exptl_absorpt_process_details
;
CrysAlisPro 1.171.39.46 (Rigaku Oxford Diffraction, 2018)
Empirical absorption correction using spherical harmonics,
implemented in SCALE3 ABSPACK scaling algorithm.
;
_exptl_absorpt_special_details ?
_diffrn_ambient_temperature 296(2)
_diffrn_radiation_wavelength 0.71073
_diffrn_radiation_type MoK\alpha
_diffrn_source ?
_diffrn_measurement_device_type 'KM4 CCD four-circle diffractometer'
_diffrn_measurement_method ?
_diffrn_detector_area_resol_mean ?
_diffrn_reflns_number 12588
_diffrn_reflns_av_unetI/netI 0.0550
_diffrn_reflns_av_R_equivalents 0.0649
_diffrn_reflns_limit_h_min -17
_diffrn_reflns_limit_h_max 17
_diffrn_reflns_limit_k_min -10
_diffrn_reflns_limit_k_max 10
_diffrn_reflns_limit_l_min -19
_diffrn_reflns_limit_l_max 23
_diffrn_reflns_theta_min 2.133
_diffrn_reflns_theta_max 28.098
_diffrn_reflns_theta_full 25.242
_diffrn_measured_fraction_theta_max 0.900
_diffrn_measured_fraction_theta_full 1.000
_diffrn_reflns_Laue_measured_fraction_max 0.900
_diffrn_reflns_Laue_measured_fraction_full 1.000
_diffrn_reflns_point_group_measured_fraction_max 0.900
_diffrn_reflns_point_group_measured_fraction_full 1.000
_reflns_number_total 2431
_reflns_number_gt 1713
_reflns_threshold_expression 'I > 2\s(I)'
_reflns_Friedel_coverage 0.000
_reflns_Friedel_fraction_max .
_reflns_Friedel_fraction_full .

_reflns_special_details
;

```

Reflections were merged by SHELXL according to the crystal class for the calculation of statistics and refinement.

_reflns_Friedel_fraction is defined as the number of unique Friedel pairs measured divided by the number that would be possible theoretically, ignoring centric projections and systematic absences.

;

```

_computing_data_collection
;
```

```

CrysAlisPro, Agilent Technologies,
Version 1.171.37.35h (release 09-02-2015 CrysAlis171 .NET)
(compiled Feb 9 2015,16:26:32)
;
_computing_cell_refinement
;
CrysAlisPro, Agilent Technologies,
Version 1.171.37.35h (release 09-02-2015 CrysAlis171 .NET)
(compiled Feb 9 2015,16:26:32)
;
_computing_data_reduction
;
CrysAlisPro, Agilent Technologies,
Version 1.171.37.35h (release 09-02-2015 CrysAlis171 .NET)
(compiled Feb 9 2015,16:26:32)
;
_computing_structure_solution  'SHELXS-2013/1 (Sheldrick, 2013)'
_computing_structure_refinement 'SHELXL-2014/7 (Sheldrick, 2014)'
_computing_molecular_graphics  ?
_computing_publication_material  ?
_refine_special_details      ?
_refine_ls_structure_factor_coef Fsqd
_refine_ls_matrix_type        full
_refine_ls_weighting_scheme   calc
_refine_ls_weighting_details
'w=1/[s^2^(Fo^2^)+(0.0805P)^2^] where P=(Fo^2^+2Fc^2^)/3'
_atom_sites_solution_primary  refxyz
_atom_sites_solution_secondary ?
_atom_sites_solution_hydrogens mixed
_refine_ls_hydrogen_treatment mixed
_refine_ls_extinction_method  'SHELXL-2014/7 (Sheldrick 2014'
_refine_ls_extinction_coeff   0.0020(6)
_refine_ls_extinction_expression
'Fc^*^=kFc[1+0.001xFc^2^|l^3^/sin(2\q)]^-1/4^'
_refine_ls_number_reflns     2431
_refine_ls_number_parameters 135
_refine_ls_number_restraints  0
_refine_ls_R_factor_all       0.0759
_refine_ls_R_factor_gt        0.0489
_refine_ls_wR_factor_ref      0.1477
_refine_ls_wR_factor_gt        0.1286
_refine_ls_goodness_of_fit_ref 1.009
_refine_ls_restrained_S_all   1.009
_refine_ls_shift/su_max        0.001
_refine_ls_shift/su_mean       0.000

loop_
_atom_site_label
_atom_site_type_symbol
_atom_site_fract_x
_atom_site_fract_y
_atom_site_fract_z
_atom_site_U_iso_or_equiv
_atom_site_adp_type
_atom_site_occupancy
_atom_site_site_symmetry_order
_atom_site_calc_flag

```

```

_atom_site_refinement_flags_posn
_atom_site_refinement_flags_adp
_atom_site_refinement_flags_occupancy
_atom_site_disorder_assembly
_atom_site_disorder_group
C2 C 0.6286(2) 0.1361(4) 0.5668(2) 0.0355(8) Uani 1 1 d . . . .
C6 C 0.5734(3) 0.3942(4) 0.4322(2) 0.0445(9) Uani 1 1 d . . . .
H6 H 0.5320 0.3270 0.4077 0.067 Uiso 1 1 calc R U . . .
C7 C 0.7393(4) 0.1713(5) 0.6660(2) 0.0650(13) Uani 1 1 d . . . .
H7A H 0.6908 0.1313 0.6979 0.097 Uiso 1 1 calc R U . . .
H7B H 0.7727 0.2582 0.6872 0.097 Uiso 1 1 calc R U . . .
H7C H 0.7861 0.0906 0.6553 0.097 Uiso 1 1 calc R U . . .
C21 C 0.5917(3) 0.5502(4) 0.4042(2) 0.0380(9) Uani 1 1 d . . . .
C22 C 0.6431(2) 0.6627(4) 0.4409(2) 0.0420(9) Uani 1 1 d . . . .
H22 H 0.6670 0.6396 0.4853 0.063 Uiso 1 1 calc R U . . .
C23 C 0.6597(3) 0.8085(4) 0.4128(2) 0.0433(9) Uani 1 1 d . . . .
H23 H 0.6946 0.8835 0.4379 0.065 Uiso 1 1 calc R U . . .
C24 C 0.6240(2) 0.8416(4) 0.3471(2) 0.0404(9) Uani 1 1 d . . . .
C25 C 0.5725(3) 0.7340(5) 0.3093(2) 0.0535(11) Uani 1 1 d . . . .
H25 H 0.5490 0.7583 0.2649 0.080 Uiso 1 1 calc R U . . .
C26 C 0.5559(3) 0.5885(5) 0.3382(2) 0.0514(10) Uani 1 1 d . . . .
H26 H 0.5202 0.5147 0.3130 0.077 Uiso 1 1 calc R U . . .
N1 N 0.6915(3) 0.2222(4) 0.60225(18) 0.0472(8) Uani 1 1 d . . . .
H1 H 0.703(4) 0.304(6) 0.586(3) 0.071 Uiso 1 1 d . U . . .
N4 N 0.5902(2) 0.1965(3) 0.50833(18) 0.0435(8) Uani 1 1 d . . . .
H4 H 0.546(3) 0.150(5) 0.490(2) 0.065 Uiso 1 1 d . U . . .
N5 N 0.6121(2) 0.3472(3) 0.48877(17) 0.0386(7) Uani 1 1 d . . . .
S3 S 0.59704(7) -0.04697(10) 0.59064(6) 0.0465(3) Uani 1 1 d . . . .
Br27 Br 0.64839(3) 1.04319(5) 0.30901(2) 0.0548(2) Uani 1 1 d . . . .

```

```

loop_
_atom_site_aniso_label
_atom_site_aniso_U_11
_atom_site_aniso_U_22
_atom_site_aniso_U_33
_atom_site_aniso_U_23
_atom_site_aniso_U_13
_atom_site_aniso_U_12
C2 0.0300(16) 0.0337(19) 0.043(2) -0.0003(15) -0.0009(16) 0.0040(14)
C6 0.0403(19) 0.036(2) 0.058(3) 0.0004(18) -0.0064(18) -0.0028(15)
C7 0.084(3) 0.058(3) 0.053(3) 0.009(2) -0.021(3) -0.021(2)
C21 0.0338(18) 0.036(2) 0.044(2) 0.0017(15) -0.0036(17) 0.0000(14)
C22 0.044(2) 0.041(2) 0.041(2) 0.0001(17) -0.0047(17) 0.0002(15)
C23 0.045(2) 0.0322(19) 0.052(3) -0.0025(17) -0.0034(17) -0.0040(15)
C24 0.0344(17) 0.0375(19) 0.049(2) 0.0065(17) 0.0033(18) -0.0012(14)
C25 0.061(3) 0.051(2) 0.048(3) 0.0087(19) -0.015(2) -0.005(2)
C26 0.057(2) 0.049(2) 0.048(2) 0.001(2) -0.019(2) -0.0117(18)
N1 0.055(2) 0.0360(17) 0.050(2) 0.0054(15) -0.0085(17) -0.0097(15)
N4 0.0383(16) 0.0337(16) 0.059(2) 0.0055(15) -0.0096(15) -0.0071(12)
N5 0.0383(15) 0.0313(15) 0.0464(19) 0.0046(13) -0.0020(15) -0.0025(12)
S3 0.0461(6) 0.0321(5) 0.0613(7) 0.0075(4) -0.0076(5) -0.0041(4)
Br27 0.0630(4) 0.0426(3) 0.0589(4) 0.01154(18) 0.0052(2) -0.00537(17)

```

```

_geom_special_details
;
All esds (except the esd in the dihedral angle between two l.s. planes)
are estimated using the full covariance matrix. The cell esds are taken

```

into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

;

```
loop_
  _geom_bond_atom_site_label_1
  _geom_bond_atom_site_label_2
  _geom_bond_distance
  _geom_bond_site_symmetry_2
  _geom_bond_publ_flag
C2 N1 1.316(5). ?
C2 N4 1.335(5). ?
C2 S3 1.686(4). ?
C6 N5 1.266(5). ?
C6 C21 1.458(5). ?
C6 H6 0.9300 . ?
C7 N1 1.446(5). ?
C7 H7A 0.9600 . ?
C7 H7B 0.9600 . ?
C7 H7C 0.9600 . ?
C21 C22 1.379(5). ?
C21 C26 1.391(5). ?
C22 C23 1.375(6). ?
C22 H22 0.9300 . ?
C23 C24 1.374(5). ?
C23 H23 0.9300 . ?
C24 C25 1.362(5). ?
C24 Br27 1.900(3). ?
C25 C26 1.379(6). ?
C25 H25 0.9300 . ?
C26 H26 0.9300 . ?
N1 H1 0.78(5). ?
N4 N5 1.373(4). ?
N4 H4 0.80(4). ?
```

```
loop_
  _geom_angle_atom_site_label_1
  _geom_angle_atom_site_label_2
  _geom_angle_atom_site_label_3
  _geom_angle
  _geom_angle_site_symmetry_1
  _geom_angle_site_symmetry_3
  _geom_angle_publ_flag
N1 C2 N4 117.9(3) . . ?
N1 C2 S3 123.0(3) . . ?
N4 C2 S3 119.1(3) . . ?
N5 C6 C21 122.2(3) . . ?
N5 C6 H6 118.9 . . ?
C21 C6 H6 118.9 . . ?
N1 C7 H7A 109.5 . . ?
N1 C7 H7B 109.5 . . ?
H7A C7 H7B 109.5 . . ?
N1 C7 H7C 109.5 . . ?
H7A C7 H7C 109.5 . . ?
H7B C7 H7C 109.5 . . ?
```

C22 C21 C26 118.2(3) . . ?
C22 C21 C6 122.6(4) . . ?
C26 C21 C6 119.3(3) . . ?
C23 C22 C21 121.2(4) . . ?
C23 C22 H22 119.4 . . ?
C21 C22 H22 119.4 . . ?
C24 C23 C22 118.9(3) . . ?
C24 C23 H23 120.5 . . ?
C22 C23 H23 120.5 . . ?
C25 C24 C23 121.8(3) . . ?
C25 C24 Br27 119.9(3) . . ?
C23 C24 Br27 118.3(3) . . ?
C24 C25 C26 118.7(4) . . ?
C24 C25 H25 120.6 . . ?
C26 C25 H25 120.6 . . ?
C25 C26 C21 121.2(4) . . ?
C25 C26 H26 119.4 . . ?
C21 C26 H26 119.4 . . ?
C2 N1 C7 123.8(3) . . ?
C2 N1 H1 116(4) . . ?
C7 N1 H1 120(4) . . ?
C2 N4 N5 120.4(3) . . ?
C2 N4 H4 118(3) . . ?
N5 N4 H4 120(3) . . ?
C6 N5 N4 116.1(3) . . ?

loop_
_geom_torsion_atom_site_label_1
_geom_torsion_atom_site_label_2
_geom_torsion_atom_site_label_3
_geom_torsion_atom_site_label_4
_geom_torsion
_geom_torsion_site_symmetry_1
_geom_torsion_site_symmetry_2
_geom_torsion_site_symmetry_3
_geom_torsion_site_symmetry_4
_geom_torsion_publ_flag
N5 C6 C21 C22 -7.3(6) ?
N5 C6 C21 C26 172.9(4) ?
C26 C21 C22 C23 -0.6(6) ?
C6 C21 C22 C23 179.6(3) ?
C21 C22 C23 C24 0.1(5) ?
C22 C23 C24 C25 0.2(6) ?
C22 C23 C24 Br27 -179.6(3) ?
C23 C24 C25 C26 0.2(6) ?
Br27 C24 C25 C26 179.9(3) ?
C24 C25 C26 C21 -0.7(7) ?
C22 C21 C26 C25 1.0(6) ?
C6 C21 C26 C25 -179.2(4) ?
N4 C2 N1 C7 -179.5(4) ?
S3 C2 N1 C7 -1.1(6) ?
N1 C2 N4 N5 -3.7(5) ?
S3 C2 N4 N5 177.9(3) ?
C21 C6 N5 N4 -178.7(3) ?
C2 N4 N5 C6 179.2(3) ?

_refine_diff_density_max 0.675

```

_refine_diff_density_min -0.457
_refine_diff_density_rms 0.107

_shelx_res_file
;

shelx.res created by SHELXL-2014/7

TITL T2_20191007_03 in Pbca
CELL 0.71073 13.5800 8.5494 19.0926 90.000 90.000 90.000
ZERR 8.00 0.0015 0.0008 0.0018 0.000 0.000 0.000
LATM 1
SYMM 1/2 - X, - Y, 1/2 + Z
SYMM - X, 1/2 + Y, 1/2 - Z
SYMM 1/2 + X, 1/2 - Y, - Z
SFAC C H N S BR
UNIT 72 80 24 8 8
MERG 2
FMAP 2
PLAN -30
ACTA
BOND $H
CONF
L.S. 30
WGHT 0.080500
EXTI 0.001960
FVAR 2.47847
MOLE 1
C2 1 0.628599 0.136142 0.566764 11.00000 0.03004 0.03371 =
0.04288 -0.00034 -0.00089 0.00396
C6 1 0.573400 0.394177 0.432235 11.00000 0.04033 0.03561 =
0.05770 0.00042 -0.00637 -0.00281
AFIX 43
H6 2 0.531971 0.327005 0.407748 11.00000 -1.50000
AFIX 0
C7 1 0.739275 0.171311 0.665992 11.00000 0.08381 0.05812 =
0.05304 0.00926 -0.02125 -0.02098
AFIX 137
H7A 2 0.690776 0.131273 0.697879 11.00000 -1.50000
H7B 2 0.772686 0.258247 0.687160 11.00000 -1.50000
H7C 2 0.786069 0.090630 0.655257 11.00000 -1.50000
AFIX 0
C21 1 0.591673 0.550154 0.404239 11.00000 0.03380 0.03596 =
0.04431 0.00171 -0.00362 0.00003
C22 1 0.643125 0.662699 0.440858 11.00000 0.04368 0.04113 =
0.04129 0.00013 -0.00471 0.00020
AFIX 43
H22 2 0.667025 0.639558 0.485313 11.00000 -1.50000
AFIX 0
C23 1 0.659718 0.808515 0.412813 11.00000 0.04542 0.03219 =
0.05233 -0.00254 -0.00341 -0.00401
AFIX 43
H23 2 0.694576 0.883537 0.437901 11.00000 -1.50000
AFIX 0
C24 1 0.623969 0.841576 0.347133 11.00000 0.03444 0.03752 =
0.04933 0.00652 0.00332 -0.00125

```

C25 1 0.572496 0.734044 0.309309 11.00000 0.06066 0.05145 =
 0.04846 0.00874 -0.01545 -0.00540
 AFIX 43
 H25 2 0.548972 0.758294 0.264871 11.00000 -1.50000
 AFIX 0
 C26 1 0.555921 0.588490 0.338169 11.00000 0.05670 0.04922 =
 0.04818 0.00135 -0.01900 -0.01165
 AFIX 43
 H26 2 0.520195 0.514657 0.312992 11.00000 -1.50000
 AFIX 0
 N1 3 0.691525 0.222195 0.602253 11.00000 0.05524 0.03596 =
 0.05031 0.00542 -0.00846 -0.00966
 H1 2 0.703428 0.304255 0.586235 11.00000 -1.50000
 N4 3 0.590234 0.196545 0.508332 11.00000 0.03826 0.03370 =
 0.05856 0.00552 -0.00962 -0.00714
 H4 2 0.545851 0.150487 0.489752 11.00000 -1.50000
 N5 3 0.612054 0.347160 0.488766 11.00000 0.03826 0.03127 =
 0.04640 0.00456 -0.00202 -0.00253
 S3 4 0.597036 -0.046970 0.590640 11.00000 0.04613 0.03205 =
 0.06130 0.00747 -0.00762 -0.00415
 BR27 5 0.648392 1.043191 0.309010 11.00000 0.06302 0.04261 =
 0.05891 0.01154 0.00523 -0.00537
 HKLF 4

REM T2_20191007_03 in Pbca
 REM R1 = 0.0489 for 1713 Fo > 4sig(Fo) and 0.0759 for all 2431 data
 REM 135 parameters refined using 0 restraints

END

WGHT 0.0806 0.0000

REM Highest difference peak 0.675, deepest hole -0.457, 1-sigma level 0.107
 Q1 1 0.7393 1.0172 0.3084 11.00000 0.05 0.67
 Q2 1 0.6866 1.0520 0.3563 11.00000 0.05 0.51
 Q3 1 0.5556 1.0251 0.3087 11.00000 0.05 0.51
 Q4 1 0.6481 0.8915 0.3087 11.00000 0.05 0.47
 Q5 1 0.7321 0.9108 0.3058 11.00000 0.05 0.41
 Q6 1 0.7632 1.1983 0.3082 11.00000 0.05 0.39
 Q7 1 0.6503 1.1956 0.3104 11.00000 0.05 0.38
 Q8 1 0.6022 1.0703 0.3587 11.00000 0.05 0.38
 Q9 1 0.5229 -0.0122 0.5885 11.00000 0.05 0.38
 Q10 1 0.6944 1.0479 0.2581 11.00000 0.05 0.37
 Q11 1 0.6588 0.7615 0.3090 11.00000 0.05 0.36
 Q12 1 0.6988 -0.0535 0.5932 11.00000 0.05 0.35
 Q13 1 0.6437 -0.0471 0.6403 11.00000 0.05 0.35
 Q14 1 0.5615 0.9315 0.3067 11.00000 0.05 0.35
 Q15 1 0.6023 1.0274 0.2580 11.00000 0.05 0.31
 Q16 1 0.4765 0.5989 0.4016 11.00000 0.05 0.30
 Q17 1 0.7105 0.8007 0.3010 11.00000 0.05 0.30
 Q18 1 0.4563 0.5559 0.3193 11.00000 0.05 0.29
 Q19 1 0.5565 0.7469 0.1882 11.00000 0.05 0.28
 Q20 1 0.7359 0.0590 0.6829 11.00000 0.05 0.28
 Q21 1 0.5843 1.2311 0.3045 11.00000 0.05 0.27
 Q22 1 0.6666 1.3203 0.3094 11.00000 0.05 0.27
 Q23 1 0.7618 0.3288 0.5913 11.00000 0.05 0.27
 Q24 1 0.6815 1.2592 0.3004 11.00000 0.05 0.26

```

Q25 1 0.4334 0.7941 0.3078 11.00000 0.05 0.25
Q26 1 0.4826 0.0672 0.5842 11.00000 0.05 0.25
Q27 1 0.7406 0.2973 0.5732 11.00000 0.05 0.25
Q28 1 0.4815 0.4325 0.3137 11.00000 0.05 0.25
Q29 1 0.6620 0.9586 0.2489 11.00000 0.05 0.24
Q30 1 0.5563 -0.0359 0.6415 11.00000 0.05 0.24
;

```

CIF for T3

data_shelx

```

_audit_creation_method      'SHELXL-2014/7'
_shelx_SHELXL_version_number  '2014/7'
_chemical_name_systematic    ?
_chemical_name_common       ?
_chemical_melting_point     ?
_chemical_formula_moiety    'C11 H15 N3 O2 S'
_chemical_formula_sum        'C11 H15 N3 O2 S'
_chemical_formula_weight     253.32

```

```

loop_
_atom_type_symbol
_atom_type_description
_atom_type_scat_dispersion_real
_atom_type_scat_dispersion_imag
_atom_type_scat_source
'C' 'C' 0.0033 0.0016
'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
'H' 'H' 0.0000 0.0000
'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
'N' 'N' 0.0061 0.0033
'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
'O' 'O' 0.0106 0.0060
'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
'S' 'S' 0.1246 0.1234
'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'

```

```

_space_group_crystal_system    monoclinic
_space_group_IT_number         14
_space_group_name_H-M_alt      'P 21/c'
_space_group_name_Hall          '-P 2ybc'

```

```

_shelx_space_group_comment
;
```

The symmetry employed for this shelxl refinement is uniquely defined by the following loop, which should always be used as a source of symmetry information in preference to the above space-group names.

They are only intended as comments.

```
;

```

```

loop_
_space_group_symop_operation_xyz
'x, y, z'
'-x, y+1/2, -z+1/2'
'-x, -y, -z'

```

'x, -y-1/2, z-1/2'

_cell_length_a 8.5409(7)
_cell_length_b 8.1329(5)
_cell_length_c 18.8922(12)
_cell_angle_alpha 90
_cell_angle_beta 102.099(7)
_cell_angle_gamma 90
_cell_volume 1283.15(16)
_cell_formula_units_Z 4
_cell_measurement_temperature 296(2)
_cell_measurement_reflns_used 4130
_cell_measurement_theta_min 2.2510
_cell_measurement_theta_max 27.8070

_exptl_crystal_description block
_exptl_crystal_colour colourless
_exptl_crystal_density_meas ?
_exptl_crystal_density_method ?
_exptl_crystal_density_diffrrn 1.311
_exptl_crystal_F_000 536
_exptl_transmission_factor_min ?
_exptl_transmission_factor_max ?
_exptl_crystal_size_max 0.5
_exptl_crystal_size_mid 0.5
_exptl_crystal_size_min 0.3
_exptl_absorpt_coefficient_mu 0.247
_shelx_estimated_absorpt_T_min ?
_shelx_estimated_absorpt_T_max ?
_exptl_absorpt_correction_T_min 0.75249
_exptl_absorpt_correction_T_max 1.00000
_exptl_absorpt_correction_type multi-scan
_exptl_absorpt_process_details
;

CrysAlisPro, Agilent Technologies,
Version 1.171.37.35h (release 09-02-2015 CrysAlis171 .NET)
(compiled Feb 9 2015,16:26:32)

Empirical absorption correction using spherical harmonics,
implemented in SCALE3 ABSPACK scaling algorithm.

;
_exptl_absorpt_special_details ?
_diffrrn_ambient_temperature 296(2)
_diffrrn_radiation_wavelength 0.71073
_diffrrn_radiation_type MoK\alpha
_diffrrn_source ?
_diffrrn_measurement_device_type 'KM4 CCD four-circle diffractometer'
_diffrrn_measurement_method ?
_diffrrn_detector_area_resol_mean ?
_diffrrn_reflns_number 8182
_diffrrn_reflns_av_unetI/netI 0.0156
_diffrrn_reflns_av_R_equivalents 0.0177
_diffrrn_reflns_limit_h_min -10
_diffrrn_reflns_limit_h_max 10
_diffrrn_reflns_limit_k_min -9
_diffrrn_reflns_limit_k_max 6
_diffrrn_reflns_limit_l_min -22
_diffrrn_reflns_limit_l_max 22

```

_diffrn_reflns_theta_min      2.205
_diffrn_reflns_theta_max      27.778
_diffrn_reflns_theta_full     25.242
_diffrn_measured_fraction_theta_max 0.879
_diffrn_measured_fraction_theta_full 1.000
_diffrn_reflns_Laue_measured_fraction_max 0.879
_diffrn_reflns_Laue_measured_fraction_full 1.000
_diffrn_reflns_point_group_measured_fraction_max 0.879
_diffrn_reflns_point_group_measured_fraction_full 1.000
_reflns_number_total         2666
_reflns_number_gt             2366
_reflns_threshold_expression  'I > 2\s(I)'
_reflns_Friedel_coverage     0.000
_reflns_Friedel_fraction_max .
_reflns_Friedel_fraction_full .

_reflns_special_details
;
Reflections were merged by SHELXL according to the crystal
class for the calculation of statistics and refinement.

_reflns_Friedel_fraction is defined as the number of unique
Friedel pairs measured divided by the number that would be
possible theoretically, ignoring centric projections and
systematic absences.
;

_computing_data_collection
;
CrysAlisPro, Agilent Technologies,
Version 1.171.37.35h (release 09-02-2015 CrysAlis171 .NET)
(compiled Feb 9 2015,16:26:32)
;
_computing_cell_refinement
;
CrysAlisPro, Agilent Technologies,
Version 1.171.37.35h (release 09-02-2015 CrysAlis171 .NET)
(compiled Feb 9 2015,16:26:32)
;
_computing_data_reduction
;
CrysAlisPro, Agilent Technologies,
Version 1.171.37.35h (release 09-02-2015 CrysAlis171 .NET)
(compiled Feb 9 2015,16:26:32)
;
_computing_structure_solution  'SHELXS-2013/1 (Sheldrick, 2013)'
_computing_structure_refinement 'SHELXL-2014/7 (Sheldrick, 2014)'
_computing_molecular_graphics   ?
_computing_publication_material ?
_refine_special_details        ?
_refine_ls_structure_factor_coef Fsqd
_refine_ls_matrix_type         full
_refine_ls_weighting_scheme    calc
_refine_ls_weighting_details   ?
'w=1/[s^2^(Fo^2^)+(0.0426P)^2^+0.3621P] where P=(Fo^2^+2Fc^2^)/3'
_atom_sites_solution_primary   refxyz
_atom_sites_solution_secondary ?

```

```

_atom_sites_solution_hydrogens mixed
_refine_ls_hydrogen_treatment mixed
_refine_ls_extinction_method 'SHELXL-2014/7 (Sheldrick 2014'
_refine_ls_extinction_coef 0.0142(17)
_refine_ls_extinction_expression
'Fc^*^=kFc[1+0.001xFc^2\|A^3/sin(2\q)]^-1/4^'
_refine_ls_number_reflns 2666
_refine_ls_number_parameters 164
_refine_ls_number_restraints 0
_refine_ls_R_factor_all 0.0362
_refine_ls_R_factor_gt 0.0313
_refine_ls_wR_factor_ref 0.0881
_refine_ls_wR_factor_gt 0.0846
_refine_ls_goodness_of_fit_ref 1.045
_refine_ls_restrained_S_all 1.045
_refine_ls_shift/su_max 0.000
_refine_ls_shift/su_mean 0.000

```

loop_

```

_atom_site_label
_atom_site_type_symbol
_atom_site_fract_x
_atom_site_fract_y
_atom_site_fract_z
_atom_site_U_iso_or_equiv
_atom_site_adp_type
_atom_site_occupancy
_atom_site_site_symmetry_order
_atom_site_calc_flag
_atom_site_refinement_flags_posn
_atom_site_refinement_flags_adp
_atom_site_refinement_flags_occupancy
_atom_site_disorder_assembly
_atom_site_disorder_group
C2 C 0.99910(15) -0.12039(15) 0.11931(7) 0.0311(3) Uani 1 1 d .....
C6 C 0.64184(15) 0.10073(18) 0.07411(7) 0.0376(3) Uani 1 1 d .....
H6 H 0.6523 0.1226 0.0270 0.056 Uiso 1 1 calc R U ...
C7 C 1.1177(2) -0.2616(2) 0.23209(8) 0.0546(4) Uani 1 1 d .....
H7A H 1.1537 -0.3494 0.2056 0.082 Uiso 1 1 calc R U ...
H7B H 1.0787 -0.3061 0.2722 0.082 Uiso 1 1 calc R U ...
H7C H 1.2051 -0.1881 0.2498 0.082 Uiso 1 1 calc R U ...
C21 C 0.50276(16) 0.16718(16) 0.09795(7) 0.0352(3) Uani 1 1 d .....
C22 C 0.47854(15) 0.14654(16) 0.16873(7) 0.0332(3) Uani 1 1 d .....
H22 H 0.5534 0.0895 0.2028 0.050 Uiso 1 1 calc R U ...
C23 C 0.34378(15) 0.21090(16) 0.18766(7) 0.0336(3) Uani 1 1 d .....
C24 C 0.22811(16) 0.29395(17) 0.13577(7) 0.0361(3) Uani 1 1 d .....
C25 C 0.25324(18) 0.3157(2) 0.06673(8) 0.0461(4) Uani 1 1 d .....
H25 H 0.1784 0.3723 0.0325 0.069 Uiso 1 1 calc R U ...
C26 C 0.39027(18) 0.2532(2) 0.04822(8) 0.0459(4) Uani 1 1 d .....
H26 H 0.4069 0.2693 0.0016 0.069 Uiso 1 1 calc R U ...
C28 C 0.4277(2) 0.1368(3) 0.31217(8) 0.0589(5) Uani 1 1 d .....
H28A H 0.4518 0.0253 0.3016 0.088 Uiso 1 1 calc R U ...
H28B H 0.3893 0.1400 0.3564 0.088 Uiso 1 1 calc R U ...
H28C H 0.5228 0.2025 0.3173 0.088 Uiso 1 1 calc R U ...
C29 C -0.03031(18) 0.4177(2) 0.10612(9) 0.0511(4) Uani 1 1 d .....
H29A H 0.0074 0.5164 0.0873 0.077 Uiso 1 1 calc R U ...
H29B H -0.1190 0.4435 0.1281 0.077 Uiso 1 1 calc R U ...

```

H29C H -0.0642 0.3408 0.0674 0.077 Uiso 1 1 calc R U . . .
 N1 N 0.99009(15) -0.17262(15) 0.18497(6) 0.0397(3) Uani 1 1 d
 H1 H 0.910(2) -0.151(2) 0.1999(11) 0.060 Uiso 1 1 d . U
 N4 N 0.87451(13) -0.03239(15) 0.08246(6) 0.0384(3) Uani 1 1 d
 H4 H 0.875(2) 0.003(2) 0.0399(10) 0.058 Uiso 1 1 d . U
 N5 N 0.74994(12) 0.01429(14) 0.11432(6) 0.0343(3) Uani 1 1 d
 O27 O 0.30829(12) 0.19967(15) 0.25478(5) 0.0484(3) Uani 1 1 d
 O28 O 0.09617(12) 0.34676(13) 0.15915(6) 0.0470(3) Uani 1 1 d
 S3 S 1.15525(4) -0.16089(5) 0.07998(2) 0.04145(14) Uani 1 1 d

loop_
 _atom_site_aniso_label
 _atom_site_aniso_U_11
 _atom_site_aniso_U_22
 _atom_site_aniso_U_33
 _atom_site_aniso_U_23
 _atom_site_aniso_U_13
 _atom_site_aniso_U_12
 C2 0.0310(6) 0.0328(6) 0.0308(6) -0.0020(5) 0.0095(5) -0.0009(5)
 C6 0.0331(7) 0.0505(8) 0.0315(7) -0.0005(6) 0.0118(5) 0.0045(6)
 C7 0.0624(10) 0.0661(10) 0.0351(8) 0.0077(7) 0.0096(7) 0.0237(8)
 C21 0.0305(6) 0.0428(7) 0.0335(7) -0.0023(5) 0.0096(5) 0.0028(5)
 C22 0.0282(6) 0.0394(7) 0.0320(7) -0.0015(5) 0.0061(5) 0.0022(5)
 C23 0.0320(6) 0.0388(7) 0.0314(6) -0.0038(5) 0.0099(5) 0.0003(5)
 C24 0.0309(6) 0.0381(7) 0.0408(7) -0.0033(6) 0.0109(6) 0.0054(5)
 C25 0.0408(8) 0.0585(9) 0.0392(8) 0.0082(6) 0.0090(6) 0.0170(7)
 C26 0.0433(8) 0.0626(10) 0.0343(7) 0.0063(6) 0.0134(6) 0.0122(7)
 C28 0.0537(10) 0.0899(13) 0.0341(8) 0.0095(8) 0.0118(7) 0.0140(9)
 C29 0.0365(8) 0.0547(9) 0.0623(10) 0.0063(8) 0.0109(7) 0.0139(7)
 N1 0.0397(7) 0.0502(7) 0.0322(6) 0.0050(5) 0.0145(5) 0.0101(5)
 N4 0.0339(6) 0.0546(7) 0.0302(6) 0.0059(5) 0.0146(5) 0.0108(5)
 N5 0.0299(5) 0.0434(6) 0.0321(6) -0.0029(5) 0.0124(4) 0.0034(5)
 O27 0.0401(6) 0.0754(7) 0.0326(5) 0.0034(5) 0.0142(4) 0.0150(5)
 O28 0.0379(5) 0.0587(6) 0.0478(6) 0.0048(5) 0.0164(5) 0.0182(5)
 S3 0.0340(2) 0.0526(2) 0.0424(2) 0.00832(15) 0.01854(15) 0.00904(14)

_geom_special_details
 ;
 All esds (except the esd in the dihedral angle between two l.s. planes)
 are estimated using the full covariance matrix. The cell esds are taken
 into account individually in the estimation of esds in distances, angles
 and torsion angles; correlations between esds in cell parameters are only
 used when they are defined by crystal symmetry. An approximate (isotropic)
 treatment of cell esds is used for estimating esds involving l.s. planes.
 ;

loop_
 _geom_bond_atom_site_label_1
 _geom_bond_atom_site_label_2
 _geom_bond_distance
 _geom_bond_site_symmetry_2
 _geom_bond_publ_flag
 C2 N1 1.3285(17) . ?
 C2 N4 1.3487(17) . ?
 C2 S3 1.6896(13) . ?
 C6 N5 1.2766(17) . ?
 C6 C21 1.4595(18) . ?

C6 H6 0.9300 . ?
 C7 N1 1.4479(18) . ?
 C7 H7A 0.9600 . ?
 C7 H7B 0.9600 . ?
 C7 H7C 0.9600 . ?
 C21 C26 1.3846(19) . ?
 C21 C22 1.4053(18) . ?
 C22 C23 1.3781(18) . ?
 C22 H22 0.9300 . ?
 C23 O27 1.3671(16) . ?
 C23 C24 1.4098(19) . ?
 C24 O28 1.3631(16) . ?
 C24 C25 1.377(2) . ?
 C25 C26 1.386(2) . ?
 C25 H25 0.9300 . ?
 C26 H26 0.9300 . ?
 C28 O27 1.4192(18) . ?
 C28 H28A 0.9600 . ?
 C28 H28B 0.9600 . ?
 C28 H28C 0.9600 . ?
 C29 O28 1.4317(18) . ?
 C29 H29A 0.9600 . ?
 C29 H29B 0.9600 . ?
 C29 H29C 0.9600 . ?
 N1 H1 0.813(19) . ?
 N4 N5 1.3809(14) . ?
 N4 H4 0.854(19) . ?

loop_
 _geom_angle_atom_site_label_1
 _geom_angle_atom_site_label_2
 _geom_angle_atom_site_label_3
 _geom_angle
 _geom_angle_site_symmetry_1
 _geom_angle_site_symmetry_3
 _geom_angle_publ_flag
 N1 C2 N4 117.25(11) . . ?
 N1 C2 S3 123.63(10) . . ?
 N4 C2 S3 119.11(9) . . ?
 N5 C6 C21 123.66(12) . . ?
 N5 C6 H6 118.2 . . ?
 C21 C6 H6 118.2 . . ?
 N1 C7 H7A 109.5 . . ?
 N1 C7 H7B 109.5 . . ?
 H7A C7 H7B 109.5 . . ?
 N1 C7 H7C 109.5 . . ?
 H7A C7 H7C 109.5 . . ?
 H7B C7 H7C 109.5 . . ?
 C26 C21 C22 119.10(12) . . ?
 C26 C21 C6 118.14(12) . . ?
 C22 C21 C6 122.75(12) . . ?
 C23 C22 C21 119.94(12) . . ?
 C23 C22 H22 120.0 . . ?
 C21 C22 H22 120.0 . . ?
 O27 C23 C22 125.10(12) . . ?
 O27 C23 C24 114.69(11) . . ?
 C22 C23 C24 120.21(12) . . ?

O28 C24 C25 124.75(12) . . ?
 O28 C24 C23 115.60(12) . . ?
 C25 C24 C23 119.65(12) . . ?
 C24 C25 C26 119.94(13) . . ?
 C24 C25 H25 120.0 . . ?
 C26 C25 H25 120.0 . . ?
 C21 C26 C25 121.11(13) . . ?
 C21 C26 H26 119.4 . . ?
 C25 C26 H26 119.4 . . ?
 O27 C28 H28A 109.5 . . ?
 O27 C28 H28B 109.5 . . ?
 H28A C28 H28B 109.5 . . ?
 O27 C28 H28C 109.5 . . ?
 H28A C28 H28C 109.5 . . ?
 H28B C28 H28C 109.5 . . ?
 O28 C29 H29A 109.5 . . ?
 O28 C29 H29B 109.5 . . ?
 H29A C29 H29B 109.5 . . ?
 O28 C29 H29C 109.5 . . ?
 H29A C29 H29C 109.5 . . ?
 H29B C29 H29C 109.5 . . ?
 C2 N1 C7 123.39(12) . . ?
 C2 N1 H1 118.2(14) . . ?
 C7 N1 H1 118.3(14) . . ?
 C2 N4 N5 121.16(11) . . ?
 C2 N4 H4 120.0(12) . . ?
 N5 N4 H4 118.6(12) . . ?
 C6 N5 N4 114.79(11) . . ?
 C23 O27 C28 117.97(11) . . ?
 C24 O28 C29 117.00(12) . . ?

loop_
 _geom_torsion_atom_site_label_1
 _geom_torsion_atom_site_label_2
 _geom_torsion_atom_site_label_3
 _geom_torsion_atom_site_label_4
 _geom_torsion
 _geom_torsion_site_symmetry_1
 _geom_torsion_site_symmetry_2
 _geom_torsion_site_symmetry_3
 _geom_torsion_site_symmetry_4
 _geom_torsion_publ_flag
 N5 C6 C21 C26 -177.27(14) ?
 N5 C6 C21 C22 2.4(2) ?
 C26 C21 C22 C23 0.3(2) ?
 C6 C21 C22 C23 -179.32(13) ?
 C21 C22 C23 O27 -179.46(13) ?
 C21 C22 C23 C24 1.6(2) ?
 O27 C23 C24 O28 -1.78(18) ?
 C22 C23 C24 O28 177.31(12) ?
 O27 C23 C24 C25 178.55(13) ?
 C22 C23 C24 C25 -2.4(2) ?
 O28 C24 C25 C26 -178.37(14) ?
 C23 C24 C25 C26 1.3(2) ?
 C22 C21 C26 C25 -1.4(2) ?
 C6 C21 C26 C25 178.24(14) ?
 C24 C25 C26 C21 0.6(2) ?

N4 C2 N1 C7 -177.01(14) . . . ?
 S3 C2 N1 C7 4.0(2) . . . ?
 N1 C2 N4 N5 5.86(19) . . . ?
 S3 C2 N4 N5 -175.08(10) . . . ?
 C21 C6 N5 N4 -177.95(12) . . . ?
 C2 N4 N5 C6 178.25(13) . . . ?
 C22 C23 O27 C28 8.9(2) . . . ?
 C24 C23 O27 C28 -172.08(14) . . . ?
 C25 C24 O28 C29 5.4(2) . . . ?
 C23 C24 O28 C29 -174.28(12) . . . ?

_refine_diff_density_max 0.219
 _refine_diff_density_min -0.159
 _refine_diff_density_rms 0.034

_shelx_res_file
 ;

shelx.res created by SHELXL-2014/7

TITL T3_20191008 in P2(1)/c
 CELL 0.71073 8.5409 8.1329 18.8922 90.000 102.099 90.000
 ZERR 4.00 0.0007 0.0005 0.0012 0.000 0.007 0.000
 LATT 1
 SYMM -X, 1/2 +Y, 1/2 -Z
 SFAC C H N O S
 UNIT 44 60 12 8 4
 MERG 2
 FMAP 2
 PLAN 20
 ACTA
 BOND \$H
 CONF
 L.S. 40
 WGHT 0.042600 0.362100
 EXTI 0.014162
 FVAR 6.08435
 MOLE 1
 C2 1 0.999099 -0.120394 0.119315 11.00000 0.03104 0.03278 =
 0.03078 -0.00204 0.00947 -0.00088
 C6 1 0.641836 0.100731 0.074110 11.00000 0.03307 0.05051 =
 0.03152 -0.00047 0.01182 0.00452
 AFIX 43
 H6 2 0.652287 0.122583 0.027002 11.00000 -1.50000
 AFIX 0
 C7 1 1.117663 -0.261567 0.232090 11.00000 0.06237 0.06610 =
 0.03506 0.00772 0.00955 0.02366
 AFIX 137
 H7A 2 1.153710 -0.349450 0.205557 11.00000 -1.50000
 H7B 2 1.078739 -0.306097 0.272203 11.00000 -1.50000
 H7C 2 1.205127 -0.188129 0.249751 11.00000 -1.50000
 AFIX 0
 C21 1 0.502762 0.167179 0.097949 11.00000 0.03046 0.04284 =
 0.03352 -0.00231 0.00959 0.00281
 C22 1 0.478541 0.146542 0.168727 11.00000 0.02816 0.03938 =
 0.03201 -0.00153 0.00609 0.00216

AFIX 43
 H22 2 0.553388 0.089506 0.202759 11.00000 -1.50000
 AFIX 0
 C23 1 0.343779 0.210897 0.187661 11.00000 0.03196 0.03879 =
 0.03138 -0.00380 0.00986 0.00028
 C24 1 0.228111 0.293953 0.135767 11.00000 0.03090 0.03811 =
 0.04075 -0.00332 0.01085 0.00539
 C25 1 0.253236 0.315732 0.066726 11.00000 0.04082 0.05854 =
 0.03919 0.00820 0.00895 0.01700
 AFIX 43
 H25 2 0.178350 0.372329 0.032541 11.00000 -1.50000
 AFIX 0
 C26 1 0.390267 0.253214 0.048219 11.00000 0.04326 0.06255 =
 0.03431 0.00635 0.01338 0.01216
 AFIX 43
 H26 2 0.406905 0.269337 0.001619 11.00000 -1.50000
 AFIX 0
 C28 1 0.427714 0.136765 0.312169 11.00000 0.05374 0.08994 =
 0.03410 0.00954 0.01182 0.01398
 AFIX 137
 H28A 2 0.451836 0.025259 0.301640 11.00000 -1.50000
 H28B 2 0.389313 0.139969 0.356406 11.00000 -1.50000
 H28C 2 0.522775 0.202516 0.317283 11.00000 -1.50000
 AFIX 0
 C29 1 -0.030307 0.417699 0.106122 11.00000 0.03646 0.05474 =
 0.06234 0.00635 0.01086 0.01389
 AFIX 137
 H29A 2 0.007395 0.516352 0.087317 11.00000 -1.50000
 H29B 2 -0.118956 0.443527 0.128101 11.00000 -1.50000
 H29C 2 -0.064168 0.340787 0.067392 11.00000 -1.50000
 AFIX 0
 N1 3 0.990085 -0.172617 0.184969 11.00000 0.03968 0.05023 =
 0.03222 0.00497 0.01451 0.01014
 H1 2 0.909968 -0.150916 0.199939 11.00000 -1.50000
 N4 3 0.874511 -0.032386 0.082455 11.00000 0.03394 0.05458 =
 0.03015 0.00595 0.01462 0.01075
 H4 2 0.875478 0.002624 0.039936 11.00000 -1.50000
 N5 3 0.749940 0.014286 0.114316 11.00000 0.02994 0.04337 =
 0.03210 -0.00287 0.01236 0.00341
 O27 4 0.308287 0.199673 0.254777 11.00000 0.04009 0.07540 =
 0.03262 0.00340 0.01422 0.01500
 O28 4 0.096170 0.346761 0.159149 11.00000 0.03788 0.05868 =
 0.04783 0.00484 0.01639 0.01818
 S3 5 1.155254 -0.160892 0.079984 11.00000 0.03397 0.05264 =
 0.04238 0.00832 0.01854 0.00904
 HKLF 4

REM T3_20191008 in P2(1)/c
 REM R1 = 0.0313 for 2366 Fo > 4sig(Fo) and 0.0362 for all 2666 data
 REM 164 parameters refined using 0 restraints

END

WGHT 0.0424 0.3629

REM Highest difference peak 0.219, deepest hole -0.159, 1-sigma level 0.034
 Q1 1 0.4927 0.1595 0.1325 11.00000 0.05 0.22

Q2	1	1.0874	-0.1043	0.1085	11.00000	0.05	0.20
Q3	1	0.5510	0.1229	0.0843	11.00000	0.05	0.19
Q4	1	0.4294	0.1742	0.0695	11.00000	0.05	0.18
Q5	1	0.3886	0.1550	0.1742	11.00000	0.05	0.18
Q6	1	0.2311	0.2859	0.1019	11.00000	0.05	0.17
Q7	1	0.3100	0.2990	0.1685	11.00000	0.05	0.17
Q8	1	0.2589	0.3260	0.1092	11.00000	0.05	0.16
Q9	1	0.4690	0.2316	0.0827	11.00000	0.05	0.16
Q10	1	0.4187	0.2131	0.1838	11.00000	0.05	0.16
Q11	1	0.3236	0.2715	0.0606	11.00000	0.05	0.15
Q12	1	0.3435	0.3116	0.0683	11.00000	0.05	0.14
Q13	1	0.6600	0.1877	0.0228	11.00000	0.05	0.14
Q14	1	0.9991	-0.1430	0.1495	11.00000	0.05	0.14
Q15	1	1.2655	-0.1819	0.1230	11.00000	0.05	0.14
Q16	1	1.2364	-0.2546	0.1108	11.00000	0.05	0.14
Q17	1	0.3341	0.1989	0.2100	11.00000	0.05	0.13
Q18	1	0.3390	0.2816	0.2475	11.00000	0.05	0.13
Q19	1	1.0569	-0.1689	0.0389	11.00000	0.05	0.13
Q20	1	0.4457	0.3191	0.0009	11.00000	0.05	0.13
	;						

CIF for T5

data_shelx

```
_audit_creation_method      'SHELXL-2014/7'
_shelx_SHELXL_version_number   '2014/7'
_chemical_name_systematic    ?
_chemical_name_common       ?
_chemical_melting_point     ?
_chemical_formula_moiety    'C9 H9 Cl2 N3 S'
_chemical_formula_sum        'C9 H9 Cl2 N3 S'
_chemical_formula_weight     262.15
```

```
loop_
_atom_type_symbol
_atom_type_description
_atom_type_scat_dispersion_real
_atom_type_scat_dispersion_imag
_atom_type_scat_source
'C' 'C' 0.0033 0.0016
'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
'H' 'H' 0.0000 0.0000
'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
'N' 'N' 0.0061 0.0033
'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
'S' 'S' 0.1246 0.1234
'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
'Cl' 'Cl' 0.1484 0.1585
'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
```

```
_space_group_crystal_system      monoclinic
_space_group_IT_number          15
_space_group_name_H-M_alt       'C 2/c'
_space_group_name_Hall          '-C 2yc'
```

```

_shelx_space_group_comment
;
The symmetry employed for this shelxl refinement is uniquely defined
by the following loop, which should always be used as a source of
symmetry information in preference to the above space-group names.
They are only intended as comments.
;

loop_
  _space_group_symop_operation_xyz
  'x, y, z'
  '-x, y, -z+1/2'
  'x+1/2, y+1/2, z'
  '-x+1/2, y+1/2, -z+1/2'
  '-x, -y, -z'
  'x, -y, z-1/2'
  '-x+1/2, -y+1/2, -z'
  'x+1/2, -y+1/2, z-1/2'

  _cell_length_a      12.8992(10)
  _cell_length_b      10.0640(8)
  _cell_length_c      18.4826(13)
  _cell_angle_alpha    90
  _cell_angle_beta     96.754(7)
  _cell_angle_gamma    90
  _cell_volume        2382.7(3)
  _cell_formula_units_Z   8
  _cell_measurement_temperature 296(2)
  _cell_measurement_reflns_used 754
  _cell_measurement_theta_min   3.1440
  _cell_measurement_theta_max   27.3160

  _exptl_crystal_description    block
  _exptl_crystal_colour        colourless
  _exptl_crystal_density_meas   ?
  _exptl_crystal_density_method ?
  _exptl_crystal_density_diffrn 1.462
  _exptl_crystal_F_000         1072
  _exptl_transmission_factor_min ?
  _exptl_transmission_factor_max ?
  _exptl_crystal_size_max       0.5
  _exptl_crystal_size_mid       0.4
  _exptl_crystal_size_min       0.3
  _exptl_absorpt_coefficient_mu 0.690
  _shelx_estimated_absorpt_T_min ?
  _shelx_estimated_absorpt_T_max ?
  _exptl_absorpt_correction_T_min 0.24257
  _exptl_absorpt_correction_T_max 1.00000
  _exptl_absorpt_correction_type multi-scan
  _exptl_absorpt_process_details
;
CrysAlisPro 1.171.40.53 (Rigaku Oxford Diffraction, 2019)
Empirical absorption correction using spherical harmonics,
implemented in SCALE3 ABSPACK scaling algorithm.
;
  _exptl_absorpt_special_details ?

```

```

._diffrn_ambient_temperature      296(2)
._diffrn_radiation_wavelength    0.71073
._diffrn_radiation_type         MoK\alpha
._diffrn_source                 ?
._diffrn_measurement_device_type 'KM4 CCD four-circle diffractometer'
._diffrn_measurement_method     ?
._diffrn_detector_area_resol_mean ?
._diffrn_reflns_number          6897
._diffrn_reflns_av_unetI/netI   0.0204
._diffrn_reflns_av_R_equivalents 0.0187
._diffrn_reflns_limit_h_min     -15
._diffrn_reflns_limit_h_max     15
._diffrn_reflns_limit_k_min     -8
._diffrn_reflns_limit_k_max     12
._diffrn_reflns_limit_l_min     -18
._diffrn_reflns_limit_l_max     22
._diffrn_reflns_theta_min       2.219
._diffrn_reflns_theta_max       28.065
._diffrn_reflns_theta_full      25.242
._diffrn_measured_fraction_theta_max 0.860
._diffrn_measured_fraction_theta_full 1.000
._diffrn_reflns_Laue_measured_fraction_max 0.860
._diffrn_reflns_Laue_measured_fraction_full 1.000
._diffrn_reflns_point_group_measured_fraction_max 0.860
._diffrn_reflns_point_group_measured_fraction_full 1.000
._reflns_number_total          2493
._reflns_number_gt              1981
._reflns_threshold_expression   'I > 2\s(I)'
._reflns_Friedel_coverage      0.000
._reflns_Friedel_fraction_max  .
._reflns_Friedel_fraction_full .

```

._reflns_special_details

;

Reflections were merged by SHELXL according to the crystal class for the calculation of statistics and refinement.

._reflns_Friedel_fraction is defined as the number of unique Friedel pairs measured divided by the number that would be possible theoretically, ignoring centric projections and systematic absences.

;

._computing_data_collection

;

CrysAlisPro, Agilent Technologies,

Version 1.171.37.35h (release 09-02-2015 CrysAlis171 .NET)

(compiled Feb 9 2015,16:26:32)

;

._computing_cell_refinement

;

CrysAlisPro, Agilent Technologies,

Version 1.171.37.35h (release 09-02-2015 CrysAlis171 .NET)

(compiled Feb 9 2015,16:26:32)

;

._computing_data_reduction

;

CrysAlisPro, Agilent Technologies,
 Version 1.171.37.35h (release 09-02-2015 CrysAlis171 .NET)
 (compiled Feb 9 2015,16:26:32)

;

_computing_structure_solution 'SHELXS-2013/1 (Sheldrick, 2013)'
 _computing_structure_refinement 'SHELXL-2014/7 (Sheldrick, 2014)'
 _computing_molecular_graphics ?
 _computing_publication_material ?
 _refine_special_details ?
 _refine_ls_structure_factor_coef Fsqd
 _refine_ls_matrix_type full
 _refine_ls_weighting_scheme calc
 _refine_ls_weighting_details
 'w=1/[s^2^(Fo^2^)+(0.0665P)^2^+2.3540P] where P=(Fo^2^+2Fc^2^)/3'
 _atom_sites_solution_primary refxyz
 _atom_sites_solution_secondary ?
 _atom_sites_solution_hydrogens mixed
 _refine_ls_hydrogen_treatment mixed
 _refine_ls_extinction_method none
 _refine_ls_extinction_coeff .
 _refine_ls_number_reflns 2493
 _refine_ls_number_parameters 143
 _refine_ls_number_restraints 0
 _refine_ls_R_factor_all 0.0568
 _refine_ls_R_factor_gt 0.0439
 _refine_ls_wR_factor_ref 0.1379
 _refine_ls_wR_factor_gt 0.1266
 _refine_ls_goodness_of_fit_ref 1.066
 _refine_ls_restrained_S_all 1.066
 _refine_ls_shift/su_max 0.001
 _refine_ls_shift/su_mean 0.000

loop_
 _atom_site_label
 _atom_site_type_symbol
 _atom_site_fract_x
 _atom_site_fract_y
 _atom_site_fract_z
 _atom_site_U_iso_or_equiv
 _atom_site_adp_type
 _atom_site_occupancy
 _atom_site_site_symmetry_order
 _atom_site_calc_flag
 _atom_site_refinement_flags_posn
 _atom_site_refinement_flags_adp
 _atom_site_refinement_flags_occupancy
 _atom_site_disorder_assembly
 _atom_site_disorder_group
 C2 C 0.16570(19) 0.9349(3) 0.74732(13) 0.0479(6) Uani 1 1 d
 C6 C 0.07636(19) 0.8279(3) 0.90895(13) 0.0500(6) Uani 1 1 d
 H6 H 0.0247 0.8922 0.9098 0.075 Uiso 1 1 calc R U ...
 C7 C 0.3129(3) 0.8474(4) 0.68877(18) 0.0766(9) Uani 1 1 d
 H7A H 0.2744 0.8420 0.6411 0.115 Uiso 1 1 calc R U ...
 H7B H 0.3593 0.7729 0.6962 0.115 Uiso 1 1 calc R U ...
 H7C H 0.3526 0.9283 0.6928 0.115 Uiso 1 1 calc R U ...
 C21 C 0.09089(18) 0.7301(2) 0.96791(13) 0.0456(5) Uani 1 1 d
 C22 C 0.16299(19) 0.6274(3) 0.96739(13) 0.0474(6) Uani 1 1 d

H22 H 0.2045 0.6206 0.9296 0.071 Uiso 1 1 calc R U . . .
 C23 C 0.1726(2) 0.5352(3) 1.02344(14) 0.0514(6) Uani 1 1 d
 C24 C 0.1105(2) 0.5437(3) 1.07997(14) 0.0550(7) Uani 1 1 d
 C25 C 0.0411(2) 0.6462(4) 1.08106(15) 0.0660(8) Uani 1 1 d
 H25 H 0.0007 0.6537 1.1194 0.099 Uiso 1 1 calc R U . . .
 C26 C 0.0306(2) 0.7386(3) 1.02553(15) 0.0615(7) Uani 1 1 d
 H26 H -0.0173 0.8074 1.0266 0.092 Uiso 1 1 calc R U . . .
 N1 N 0.24033(19) 0.8462(2) 0.74377(13) 0.0563(6) Uani 1 1 d
 H1 H 0.252(3) 0.793(4) 0.7728(19) 0.085 Uiso 1 1 d . U . . .
 N4 N 0.11103(18) 0.9260(2) 0.80540(13) 0.0534(6) Uani 1 1 d
 H4 H 0.064(3) 0.974(4) 0.809(2) 0.080 Uiso 1 1 d . U . . .
 N5 N 0.13225(15) 0.8277(2) 0.85666(11) 0.0477(5) Uani 1 1 d
 S3 S 0.13423(6) 1.05573(7) 0.68487(4) 0.0590(2) Uani 1 1 d
 Cl23 Cl 0.26246(8) 0.40935(8) 1.02152(5) 0.0844(3) Uani 1 1 d
 Cl24 Cl 0.11927(8) 0.42480(9) 1.14854(4) 0.0825(3) Uani 1 1 d

loop_
 _atom_site_aniso_label
 _atom_site_aniso_U_11
 _atom_site_aniso_U_22
 _atom_site_aniso_U_33
 _atom_site_aniso_U_23
 _atom_site_aniso_U_13
 _atom_site_aniso_U_12
 C2 0.0512(13) 0.0475(14) 0.0443(13) 0.0008(10) 0.0036(10) -0.0135(11)
 C6 0.0466(13) 0.0515(15) 0.0521(14) 0.0039(11) 0.0065(11) 0.0035(11)
 C7 0.087(2) 0.080(2) 0.0699(19) 0.0020(17) 0.0361(17) 0.0058(18)
 C21 0.0420(12) 0.0499(14) 0.0443(12) 0.0019(10) 0.0031(9) -0.0032(10)
 C22 0.0509(13) 0.0480(14) 0.0435(13) -0.0031(11) 0.0062(10) -0.0035(11)
 C23 0.0576(14) 0.0426(13) 0.0513(14) -0.0045(11) -0.0041(11) -0.0020(11)
 C24 0.0588(15) 0.0628(17) 0.0414(14) 0.0043(12) -0.0031(11) -0.0187(13)
 C25 0.0596(16) 0.093(2) 0.0473(15) 0.0061(15) 0.0160(12) -0.0056(16)
 C26 0.0516(15) 0.074(2) 0.0603(17) 0.0052(14) 0.0135(12) 0.0072(13)
 N1 0.0652(14) 0.0538(14) 0.0520(13) 0.0053(10) 0.0149(11) -0.0013(11)
 N4 0.0517(12) 0.0561(14) 0.0529(13) 0.0126(10) 0.0078(10) 0.0051(10)
 N5 0.0469(11) 0.0482(12) 0.0473(11) 0.0070(9) 0.0029(9) -0.0017(9)
 S3 0.0660(4) 0.0533(4) 0.0566(4) 0.0129(3) 0.0023(3) -0.0137(3)
 Cl23 0.1153(7) 0.0582(5) 0.0789(6) 0.0015(4) 0.0077(5) 0.0310(4)
 Cl24 0.1014(6) 0.0856(6) 0.0561(5) 0.0224(4) -0.0099(4) -0.0293(5)

_geom_special_details
 ;
 All esds (except the esd in the dihedral angle between two l.s. planes)
 are estimated using the full covariance matrix. The cell esds are taken
 into account individually in the estimation of esds in distances, angles
 and torsion angles; correlations between esds in cell parameters are only
 used when they are defined by crystal symmetry. An approximate (isotropic)
 treatment of cell esds is used for estimating esds involving l.s. planes.
 ;

loop_
 _geom_bond_atom_site_label_1
 _geom_bond_atom_site_label_2
 _geom_bond_distance
 _geom_bond_site_symmetry_2
 _geom_bond_publ_flag
 C2 N1 1.320(4) . ?

C2 N4 1.355(3) . ?
 C2 S3 1.693(3) . ?
 C6 N5 1.272(3) . ?
 C6 C21 1.464(3) . ?
 C6 H6 0.9300 . ?
 C7 N1 1.461(4) . ?
 C7 H7A 0.9600 . ?
 C7 H7B 0.9600 . ?
 C7 H7C 0.9600 . ?
 C21 C22 1.391(4) . ?
 C21 C26 1.393(4) . ?
 C22 C23 1.385(4) . ?
 C22 H22 0.9300 . ?
 C23 C24 1.392(4) . ?
 C23 Cl23 1.720(3) . ?
 C24 C25 1.368(4) . ?
 C24 Cl24 1.737(3) . ?
 C25 C26 1.379(4) . ?
 C25 H25 0.9300 . ?
 C26 H26 0.9300 . ?
 N1 H1 0.76(4) . ?
 N4 N5 1.375(3) . ?
 N4 H4 0.79(4) . ?

loop_
 _geom_angle_atom_site_label_1
 _geom_angle_atom_site_label_2
 _geom_angle_atom_site_label_3
 _geom_angle
 _geom_angle_site_symmetry_1
 _geom_angle_site_symmetry_3
 _geom_angle_publ_flag
 N1 C2 N4 116.5(2) . . ?
 N1 C2 S3 124.7(2) . . ?
 N4 C2 S3 118.8(2) . . ?
 N5 C6 C21 122.0(2) . . ?
 N5 C6 H6 119.0 . . ?
 C21 C6 H6 119.0 . . ?
 N1 C7 H7A 109.5 . . ?
 N1 C7 H7B 109.5 . . ?
 H7A C7 H7B 109.5 . . ?
 N1 C7 H7C 109.5 . . ?
 H7A C7 H7C 109.5 . . ?
 H7B C7 H7C 109.5 . . ?
 C22 C21 C26 119.0(2) . . ?
 C22 C21 C6 121.4(2) . . ?
 C26 C21 C6 119.6(2) . . ?
 C23 C22 C21 119.6(2) . . ?
 C23 C22 H22 120.2 . . ?
 C21 C22 H22 120.2 . . ?
 C22 C23 C24 120.7(2) . . ?
 C22 C23 Cl23 118.6(2) . . ?
 C24 C23 Cl23 120.6(2) . . ?
 C25 C24 C23 119.6(2) . . ?
 C25 C24 Cl24 119.7(2) . . ?
 C23 C24 Cl24 120.8(2) . . ?
 C24 C25 C26 120.2(3) . . ?

C24 C25 H25 119.9 . . ?

C26 C25 H25 119.9 . . ?

C25 C26 C21 120.9(3) . . ?

C25 C26 H26 119.6 . . ?

C21 C26 H26 119.6 . . ?

C2 N1 C7 124.0(3) . . ?

C2 N1 H1 121(3) . . ?

C7 N1 H1 114(3) . . ?

C2 N4 N5 120.9(2) . . ?

C2 N4 H4 120(3) . . ?

N5 N4 H4 119(3) . . ?

C6 N5 N4 116.0(2) . . ?

loop_

_geom_torsion_atom_site_label_1

_geom_torsion_atom_site_label_2

_geom_torsion_atom_site_label_3

_geom_torsion_atom_site_label_4

_geom_torsion

_geom_torsion_site_symmetry_1

_geom_torsion_site_symmetry_2

_geom_torsion_site_symmetry_3

_geom_torsion_site_symmetry_4

_geom_torsion_publ_flag

N5 C6 C21 C22 -3.6(4) ?

N5 C6 C21 C26 177.1(3) ?

C26 C21 C22 C23 0.7(4) ?

C6 C21 C22 C23 -178.6(2) ?

C21 C22 C23 C24 0.4(4) ?

C21 C22 C23 Cl23 -179.92(19) ?

C22 C23 C24 C25 -1.6(4) ?

Cl23 C23 C24 C25 178.7(2) ?

C22 C23 C24 Cl24 177.68(19) ?

Cl23 C23 C24 Cl24 -2.0(3) ?

C23 C24 C25 C26 1.7(4) ?

Cl24 C24 C25 C26 -177.6(2) ?

C24 C25 C26 C21 -0.6(4) ?

C22 C21 C26 C25 -0.6(4) ?

C6 C21 C26 C25 178.6(3) ?

N4 C2 N1 C7 -174.5(3) ?

S3 C2 N1 C7 5.9(4) ?

N1 C2 N4 N5 -1.1(4) ?

S3 C2 N4 N5 178.55(18) ?

C21 C6 N5 N4 -179.3(2) ?

C2 N4 N5 C6 -179.8(2) ?

_refine_diff_density_max 0.559

_refine_diff_density_min -0.344

_refine_diff_density_rms 0.048

_shelx_res_file

;

shelx.res created by SHELXL-2014/7

TITL T5_20191015 in C2/c

CELL 0.71073 12.8992 10.0640 18.4826 90.000 96.754 90.000
 ZERR 8.00 0.0010 0.0008 0.0013 0.000 0.007 0.000
 LATT 7
 SYMM - X, Y, 1/2 - Z
 SFAC C H N S CL
 UNIT 72 72 24 8 16
 MERG 2
 FMAP 2
 PLAN -10
 ACTA
 BOND \$H
 CONF
 L.S. 30
 WGHT 0.066500 2.354000
 FVAR 2.52979
 MOLE 1
 C2 1 0.165703 0.934857 0.747320 11.00000 0.05125 0.04754 =
 0.04427 0.00079 0.00358 -0.01352
 C6 1 0.076356 0.827928 0.908951 11.00000 0.04660 0.05149 =
 0.05210 0.00386 0.00645 0.00346
 AFIX 43
 H6 2 0.024689 0.892181 0.909764 11.00000 -1.50000
 AFIX 0
 C7 1 0.312882 0.847437 0.688771 11.00000 0.08681 0.07955 =
 0.06986 0.00203 0.03614 0.00584
 AFIX 137
 H7A 2 0.274412 0.841998 0.641137 11.00000 -1.50000
 H7B 2 0.359308 0.772853 0.696217 11.00000 -1.50000
 H7C 2 0.352618 0.928302 0.692811 11.00000 -1.50000
 AFIX 0
 C21 1 0.090886 0.730084 0.967914 11.00000 0.04203 0.04995 =
 0.04426 0.00188 0.00311 -0.00316
 C22 1 0.162991 0.627359 0.967388 11.00000 0.05085 0.04797 =
 0.04348 -0.00307 0.00621 -0.00346
 AFIX 43
 H22 2 0.204452 0.620620 0.929649 11.00000 -1.50000
 AFIX 0
 C23 1 0.172617 0.535202 1.023439 11.00000 0.05765 0.04263 =
 0.05133 -0.00454 -0.00407 -0.00198
 C24 1 0.110514 0.543726 1.079967 11.00000 0.05875 0.06278 =
 0.04145 0.00433 -0.00306 -0.01872
 C25 1 0.041052 0.646234 1.081056 11.00000 0.05964 0.09346 =
 0.04731 0.00607 0.01597 -0.00564
 AFIX 43
 H25 2 0.000706 0.653685 1.119379 11.00000 -1.50000
 AFIX 0
 C26 1 0.030639 0.738559 1.025534 11.00000 0.05156 0.07427 =
 0.06035 0.00520 0.01354 0.00725
 AFIX 43
 H26 2 -0.017255 0.807403 1.026580 11.00000 -1.50000
 AFIX 0
 N1 3 0.240326 0.846156 0.743769 11.00000 0.06521 0.05378 =
 0.05196 0.00532 0.01486 -0.00133
 H1 2 0.251674 0.792910 0.772806 11.00000 -1.50000
 N4 3 0.111026 0.926031 0.805404 11.00000 0.05172 0.05610 =
 0.05291 0.01258 0.00781 0.00511
 H4 2 0.063833 0.974032 0.808952 11.00000 -1.50000

N5 3 0.132246 0.827714 0.856657 11.00000 0.04695 0.04824 =
 0.04733 0.00698 0.00292 -0.00167
 S3 4 0.134225 1.055731 0.684875 11.00000 0.06602 0.05328 =
 0.05664 0.01286 0.00232 -0.01371
 CL23 5 0.262456 0.409351 1.021518 11.00000 0.11529 0.05822 =
 0.07886 0.00149 0.00765 0.03099
 CL24 5 0.119271 0.424798 1.148541 11.00000 0.10141 0.08560 =
 0.05606 0.02240 -0.00990 -0.02930
 HKLF 4

REM T5_20191015 in C2/c
 REM R1 = 0.0439 for 1981 Fo > 4sig(Fo) and 0.0568 for all 2493 data
 REM 143 parameters refined using 0 restraints

END

WGHT 0.0653 2.4399

REM Highest difference peak 0.559, deepest hole -0.344, 1-sigma level 0.048
 Q1 1 0.0130 0.5889 1.1483 11.00000 0.05 0.56
 Q2 1 0.1869 0.3654 1.1163 11.00000 0.05 0.53
 Q3 1 0.0606 0.5799 1.0632 11.00000 0.05 0.21
 Q4 1 0.0958 1.0978 0.7081 11.00000 0.05 0.21
 Q5 1 0.1240 0.6857 0.9687 11.00000 0.05 0.19
 Q6 1 0.0841 0.7772 0.9407 11.00000 0.05 0.19
 Q7 1 0.0612 0.4755 1.1436 11.00000 0.05 0.19
 Q8 1 0.1447 0.5058 1.0579 11.00000 0.05 0.16
 Q9 1 0.1017 0.5188 1.0972 11.00000 0.05 0.15
 Q10 1 0.2041 0.4886 1.0169 11.00000 0.05 0.15
 ;

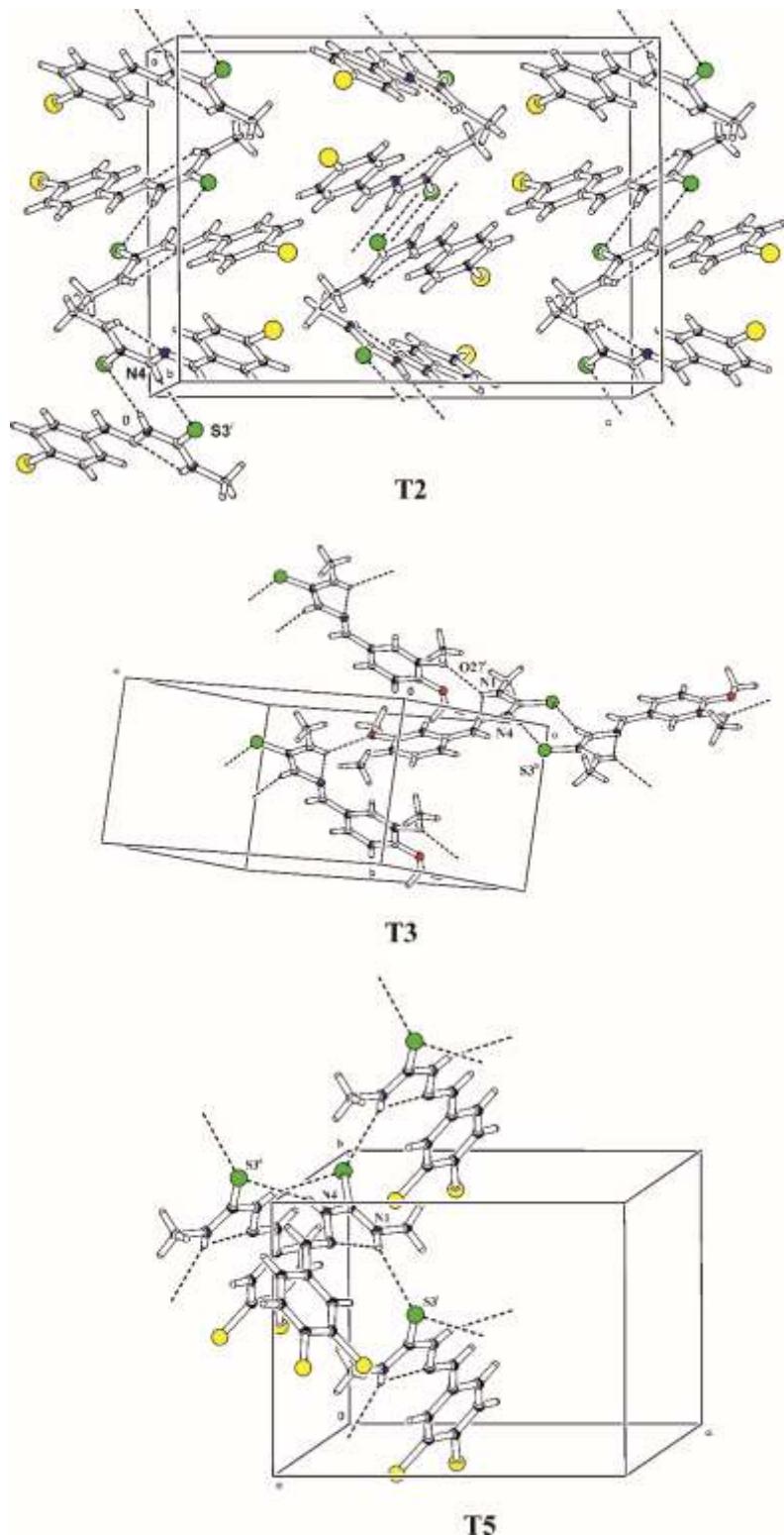


Figure 1S. The molecular packing of **T2**, **T3** and **T5**. Dashed lines indicate intermolecular hydrogen bonds; **T2**: $i = 1-x, -y, 1-z$; **T3**: $i = 1-x, -1/2+y, 1/2-z$; $ii = 2-x, -y, -z$; **T5**: $i = 1/2-x, -1/2+y, 3/2-z$; $ii = -x, y, 2-z$.

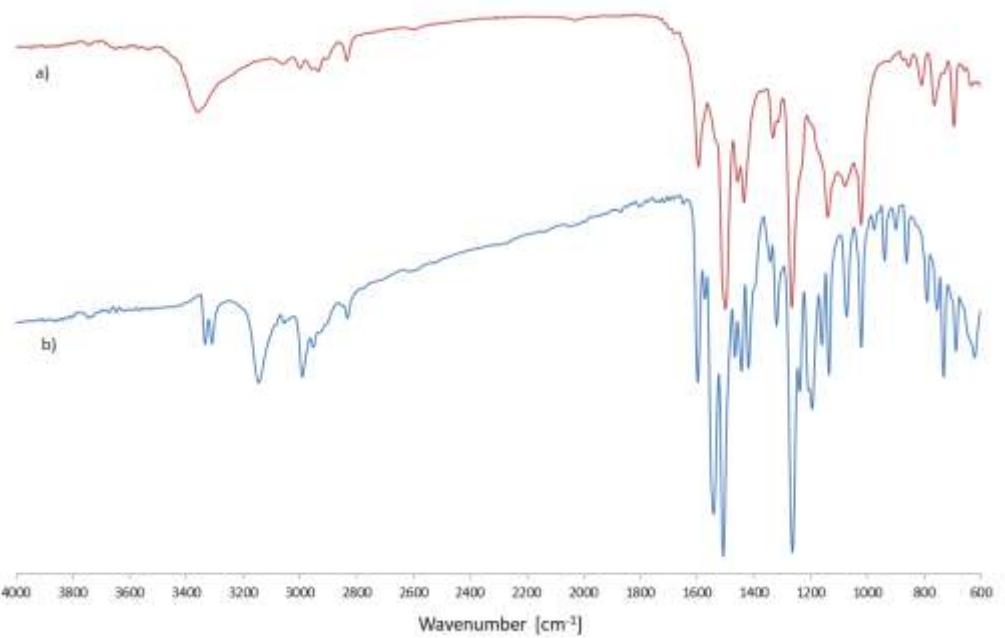


Figure 2S. Infrared spectra of a) Cu(T1)Cl₂ and b) T1

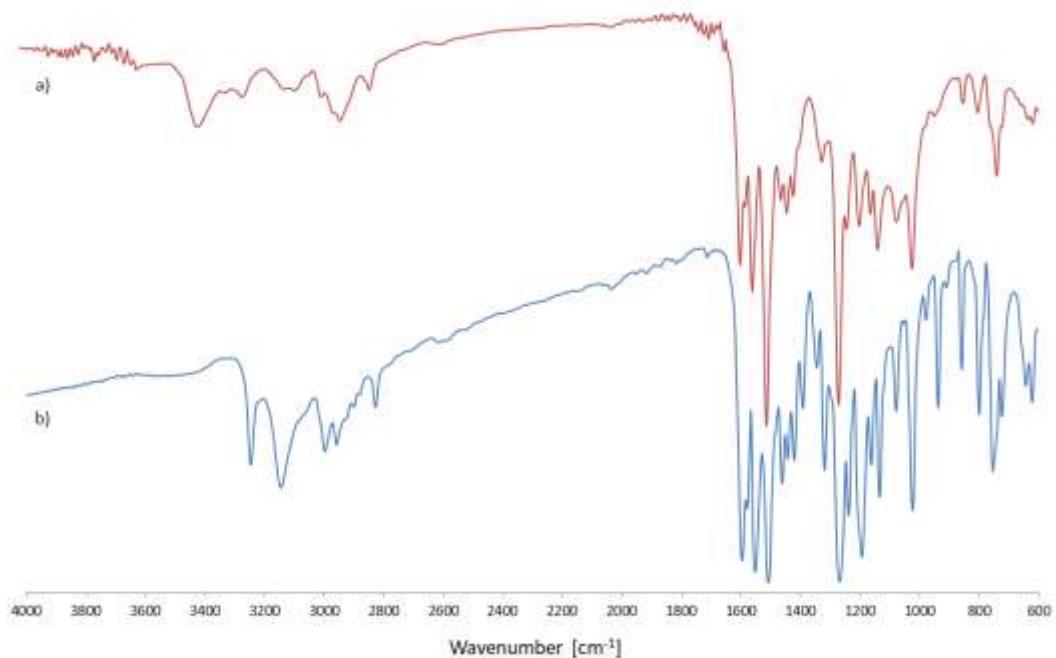


Figure 3S. Infrared spectra of a) Cu(T10)₂Cl₂ and b) T10

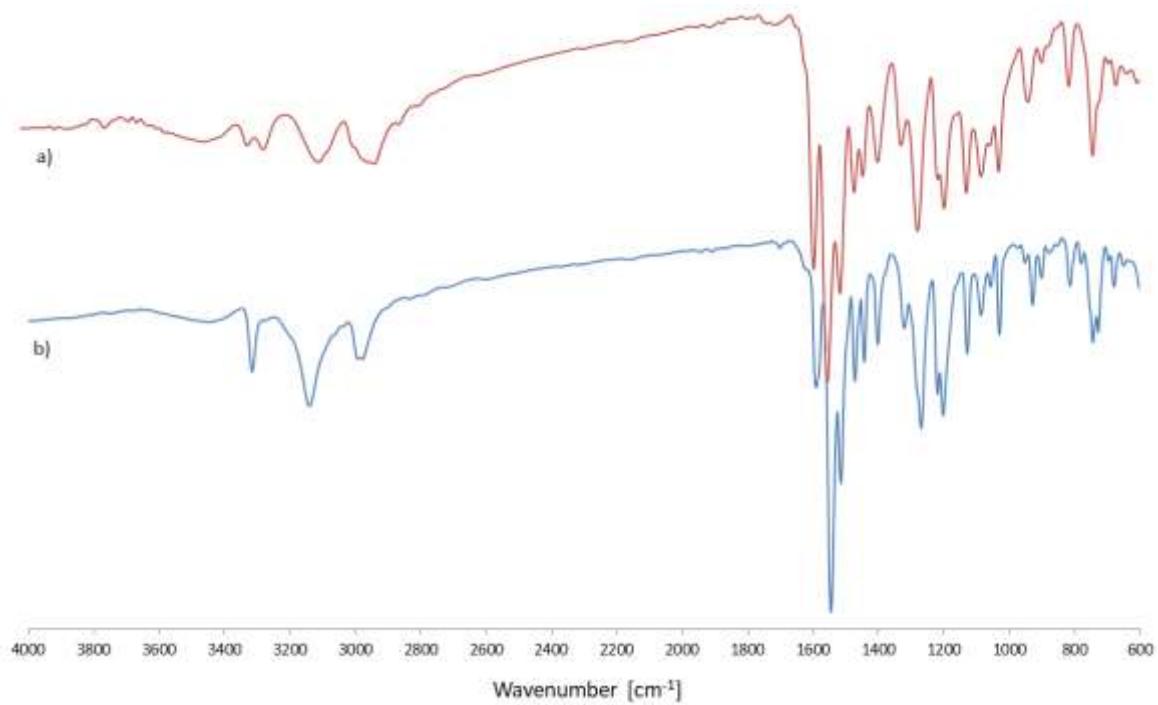


Figure 4S. Infrared spectra of a) Cu(T12)₂Cl₂ and b) T12

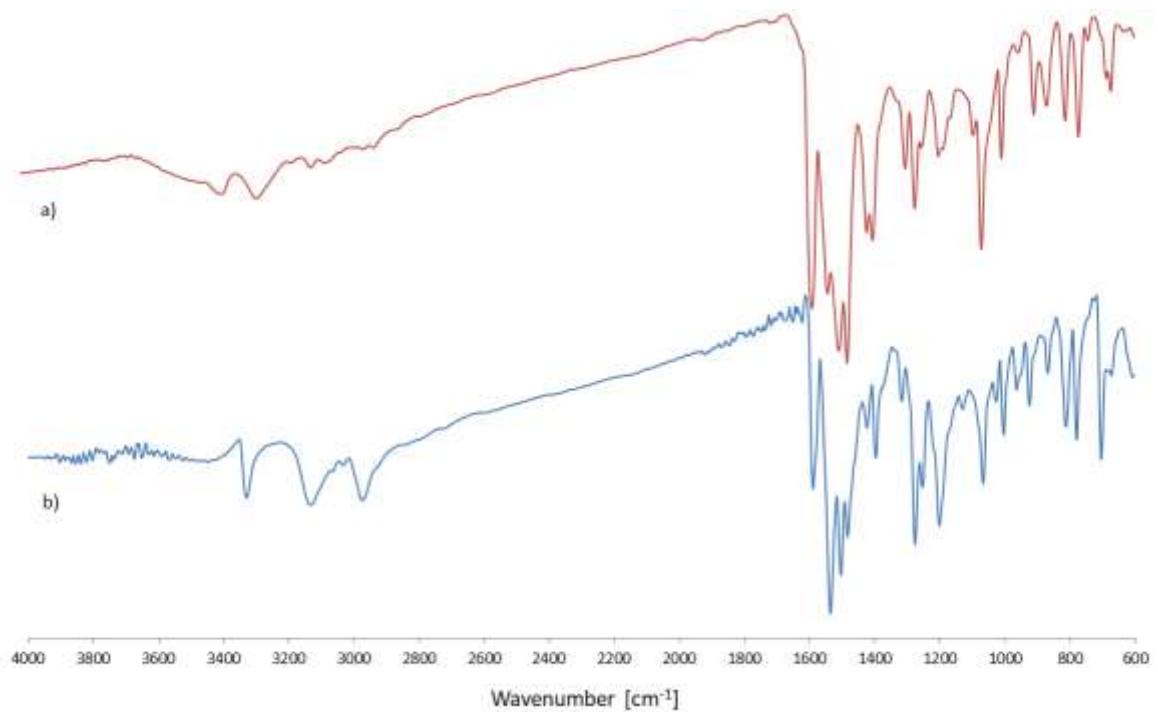


Figure 5S. Infrared spectra of a) Cu(T13)Cl₂ and b) T13

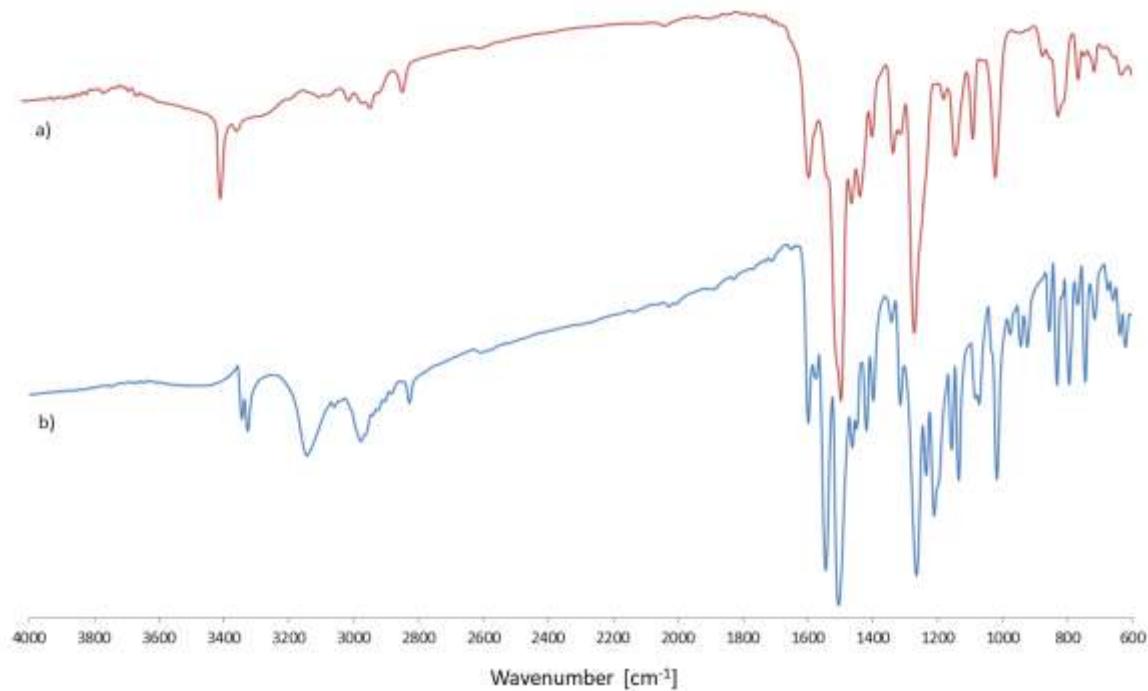


Figure 6S. Infrared spectra of a) Cu(T16)Cl₂ and b) T16

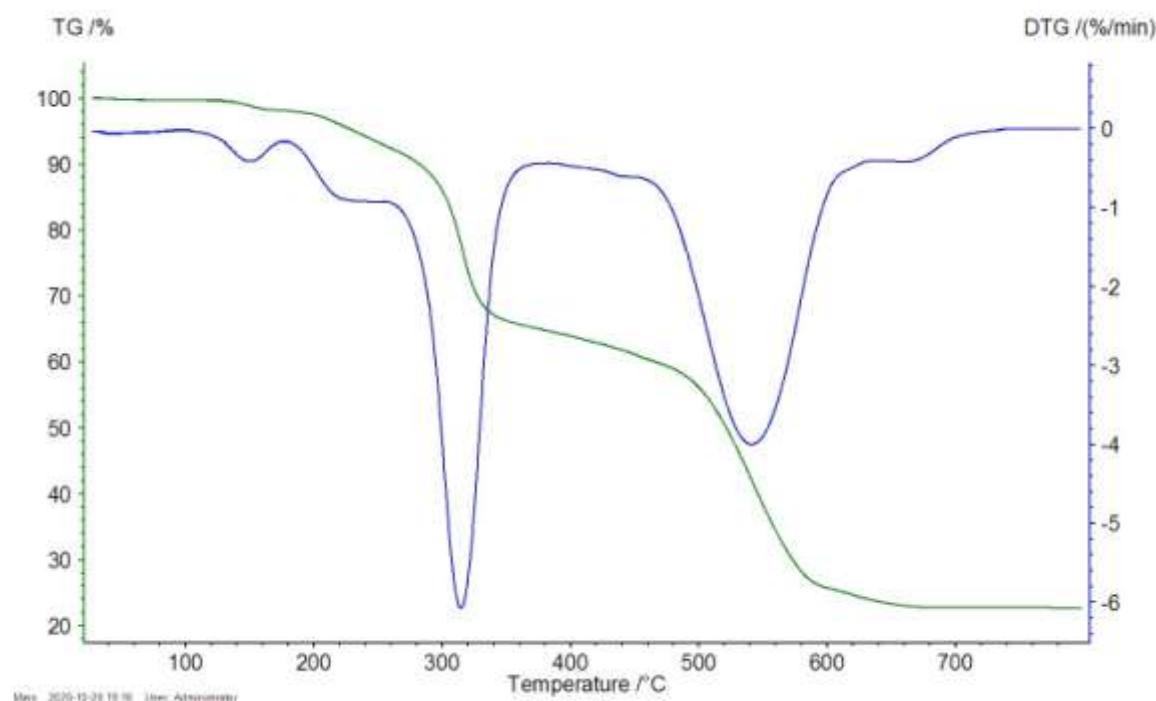


Figure 7S. TG and DTG curves of Cu(T16)Cl₂ in air atmosphere

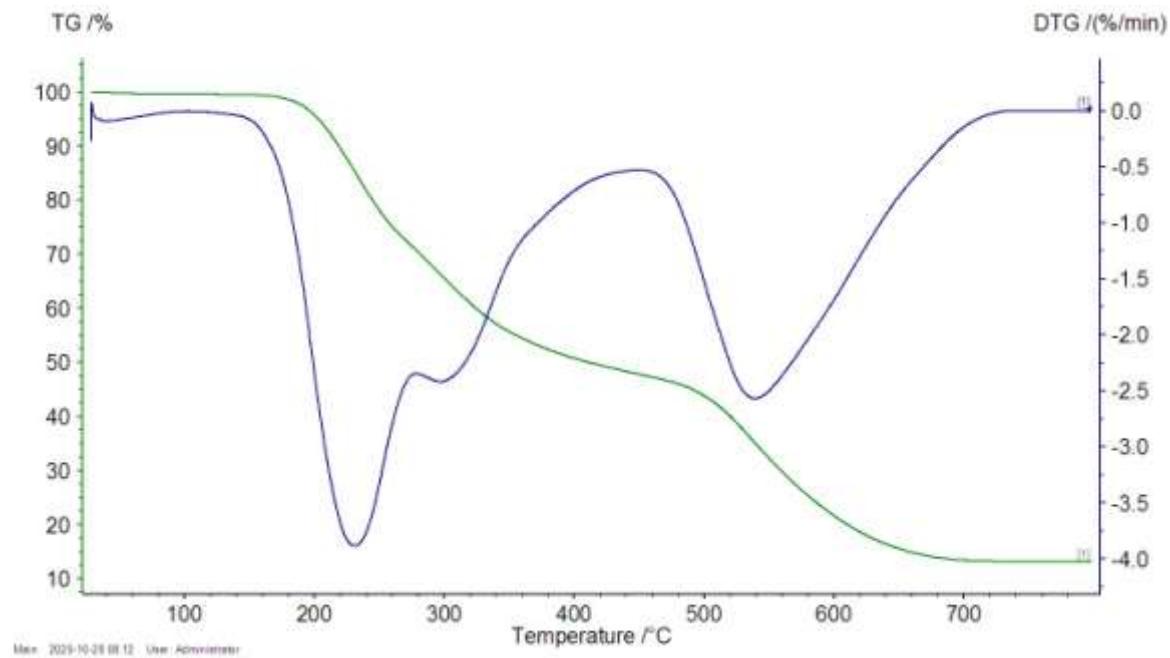


Figure 8S. TG and DTG curves of Cu(T10)₂Cl₂ in air atmosphere

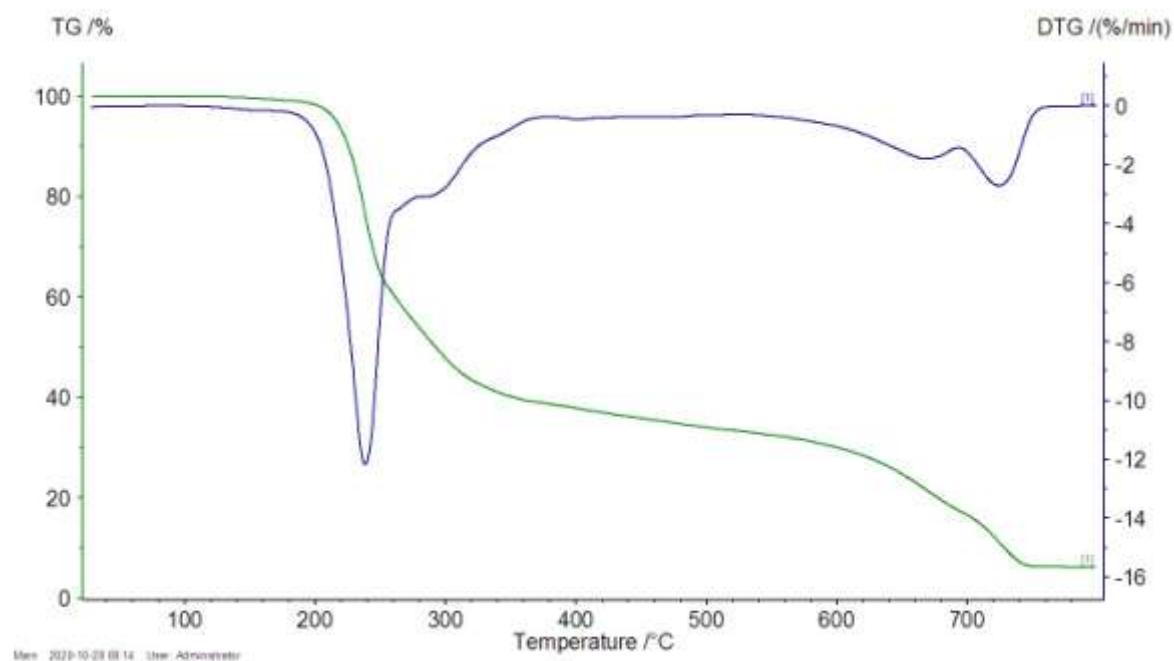


Figure 9S. TG and DTG curves of Cu(T12)₂Cl₂ in air atmosphere

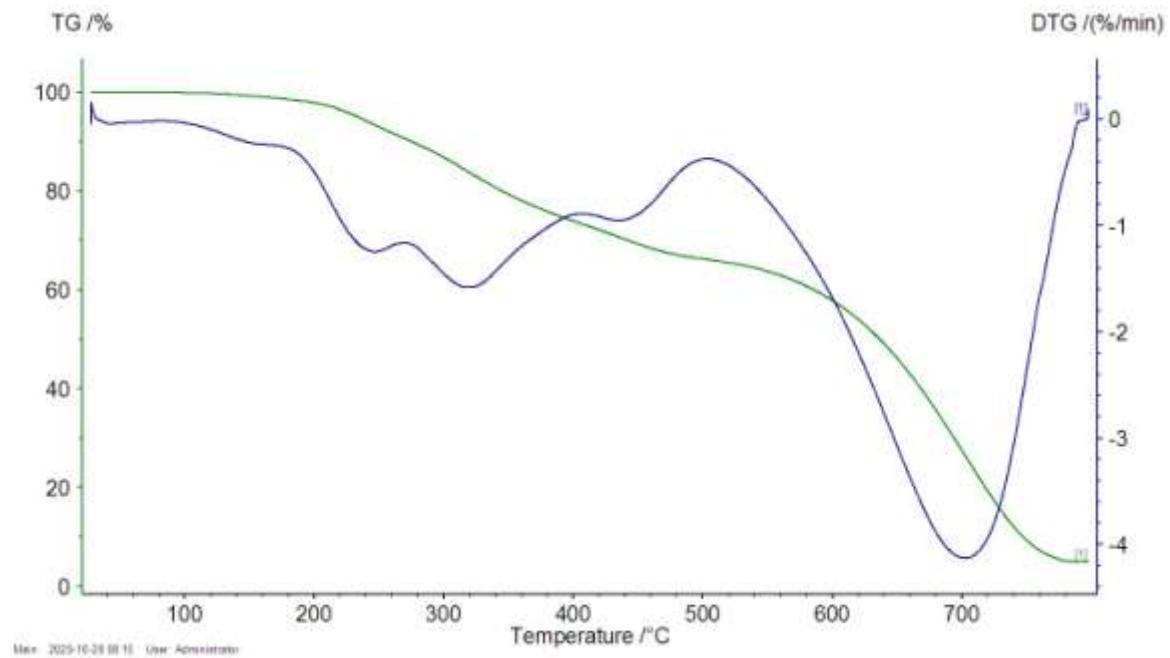


Figure 10S. TG and DTG curves of Cu(T13)Cl₂ in air atmosphere

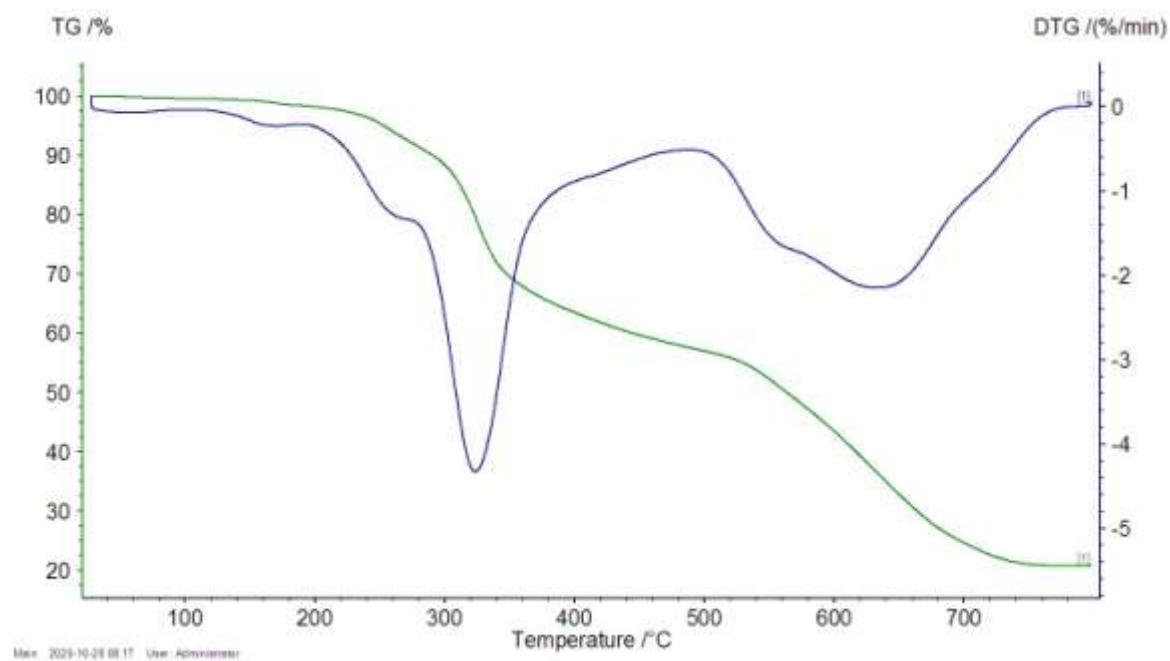
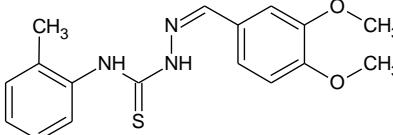
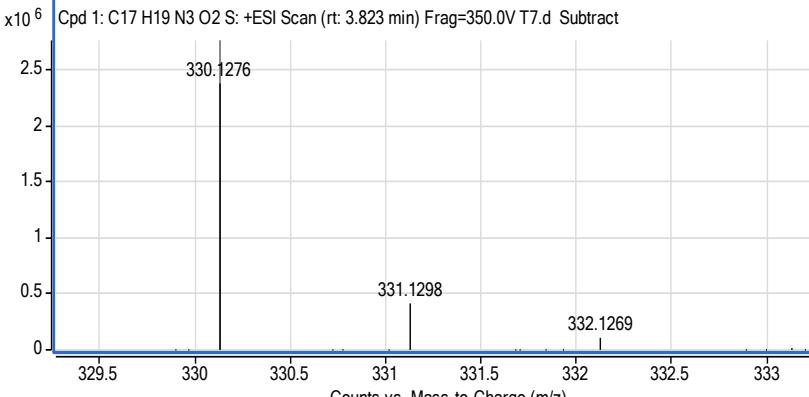
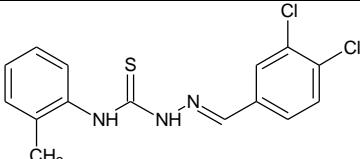
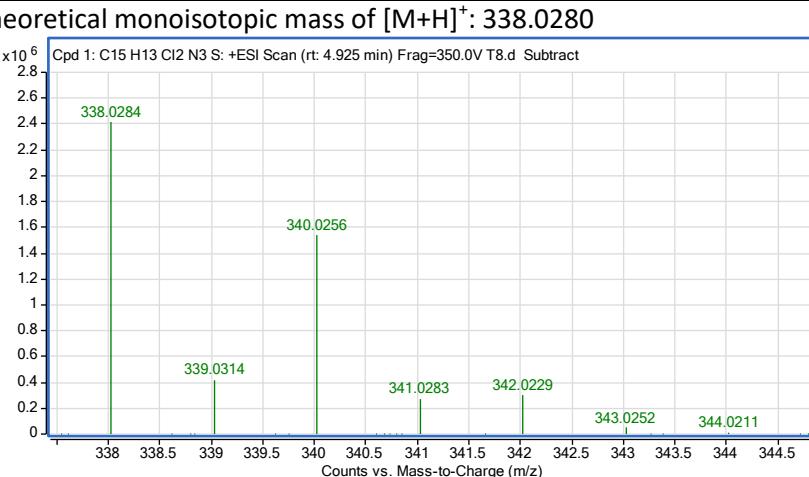
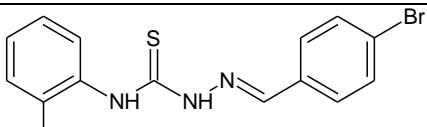


Figure 11S. TG and DTG curves of Cu(T16)Cl₂ in air atmosphere

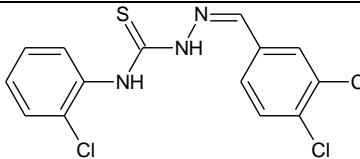
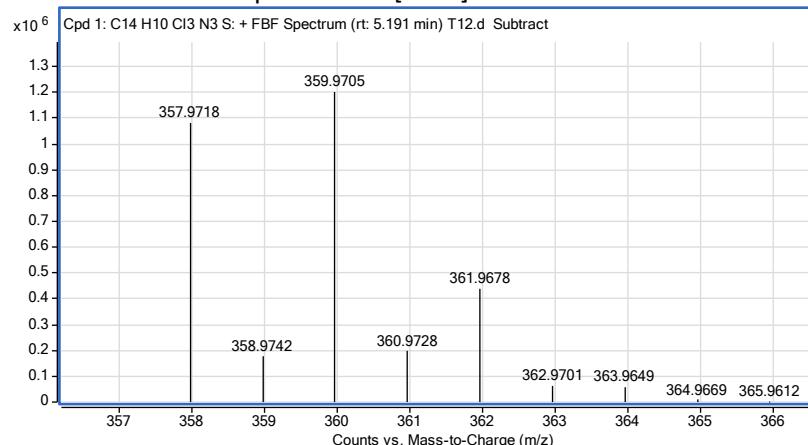
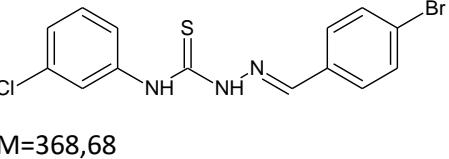
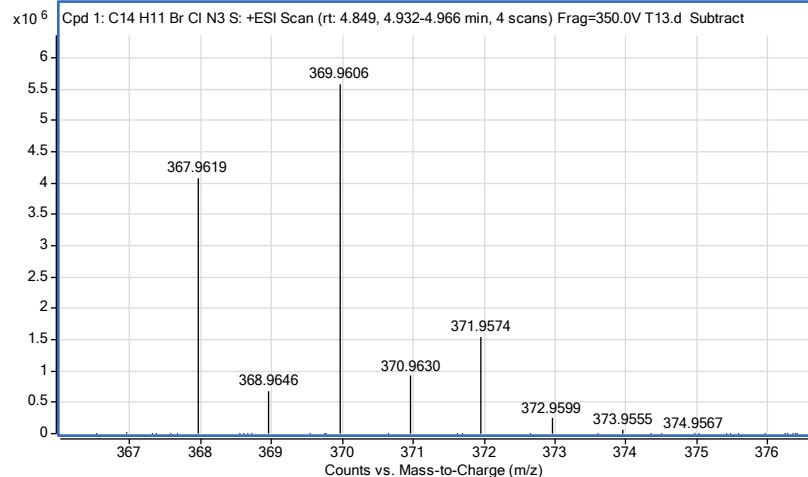
Table 1S. Spectrometry mass (MS) for new compounds.

LP.	Compound structure	Compound formula, theoretical and measured monoisotopic mass, and mass difference in ppm	Theoretical monoisotopic mass of protonated ion and acquired LC/QTOF HRMS spectra
T3	 $M=253.32$	<p>C11 H15 N3 O2S Calculated monoisotopic mass: 253.0885 Measured monoisotopic mass: 253.0889 Mass error: 1.55 ppm</p>	<p>Theoretical monoisotopic mass of $[M+H]^+$: 254.0958</p> <p>x10⁵ Cpd 1: C11 H15 N3 O2 S: +ESI Scan (rt: 2.539-2.556, 2.855-3.371 min, 34 scans) Frag=350.0V T3.d Subtract</p> <p>254.0962</p> <p>255.0988</p> <p>256.0938</p> <p>Counts vs. Mass-to-Charge (m/z)</p>
T4	 $M=223.31$	<p>C10H14N4S Calculated monoisotopic mass: 222.0939 Measured monoisotopic mass: 222.0944 Mass error: 2.14 ppm</p>	<p>Theoretical monoisotopic mass of $[M+H]^+$: 223.1012</p> <p>x10⁶ Cpd 1: C10 H14 N4 S: +ESI Scan (rt: 1.662, 1.762 min, 2 scans) Frag=350.0V T4.d Subtract</p> <p>223.1016</p> <p>224.1046</p> <p>225.0982</p> <p>Counts vs. Mass-to-Charge (m/z)</p>

T5	 M=262,26	<p>C9H9N3SCl2 Calculated monoisotopic mass: 260.9894 Measured monoisotopic mass: 260.9896 Mass error: 0.55 ppm</p>	<p>Theoretical monoisotopic mass of [M+H]⁺: 261.9967</p>
T6	 M=348,26	<p>C15H14N3SBr Calculated monoisotopic mass: 347.0092 Measured monoisotopic mass: 347.0096 Mass error: 1.14 ppm</p>	<p>Theoretical monoisotopic mass of [M+H]⁺: 348.0165</p>

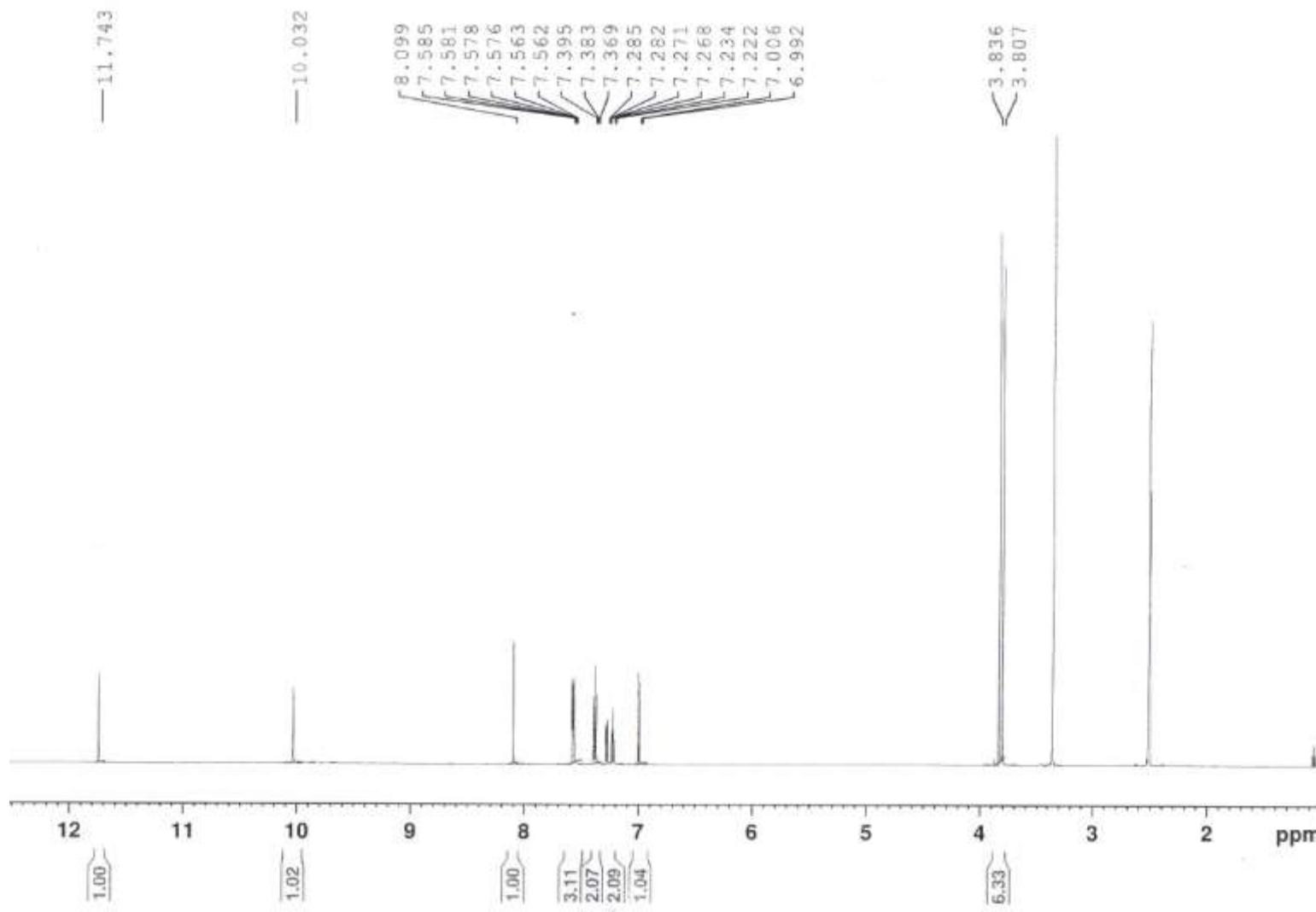
T7	 $M=329,41$	<p>C17H19N3O2S Calculated monoisotopic mass: 329.1198 Measured monoisotopic mass: 329.1202 Mass error: 1.31 ppm</p>	
T8	 $M=338,25$	<p>C15H13N3SCl2 Calculated monoisotopic mass: 337.0207 Measured monoisotopic mass: 337.0211 Mass error: 1.23 ppm</p>	<p>Theoretical monoisotopic mass of $[M+H]^+$: 338.0280</p> 
T9	 $M=368,68$	<p>C14H11N3SClBr Calculated monoisotopic mass: 366.9546 Measured monoisotopic mass: 366.9545 Mass error: -0.03 ppm</p>	<p>Theoretical monoisotopic mass of $[M+H]^+$: 367.9618</p>

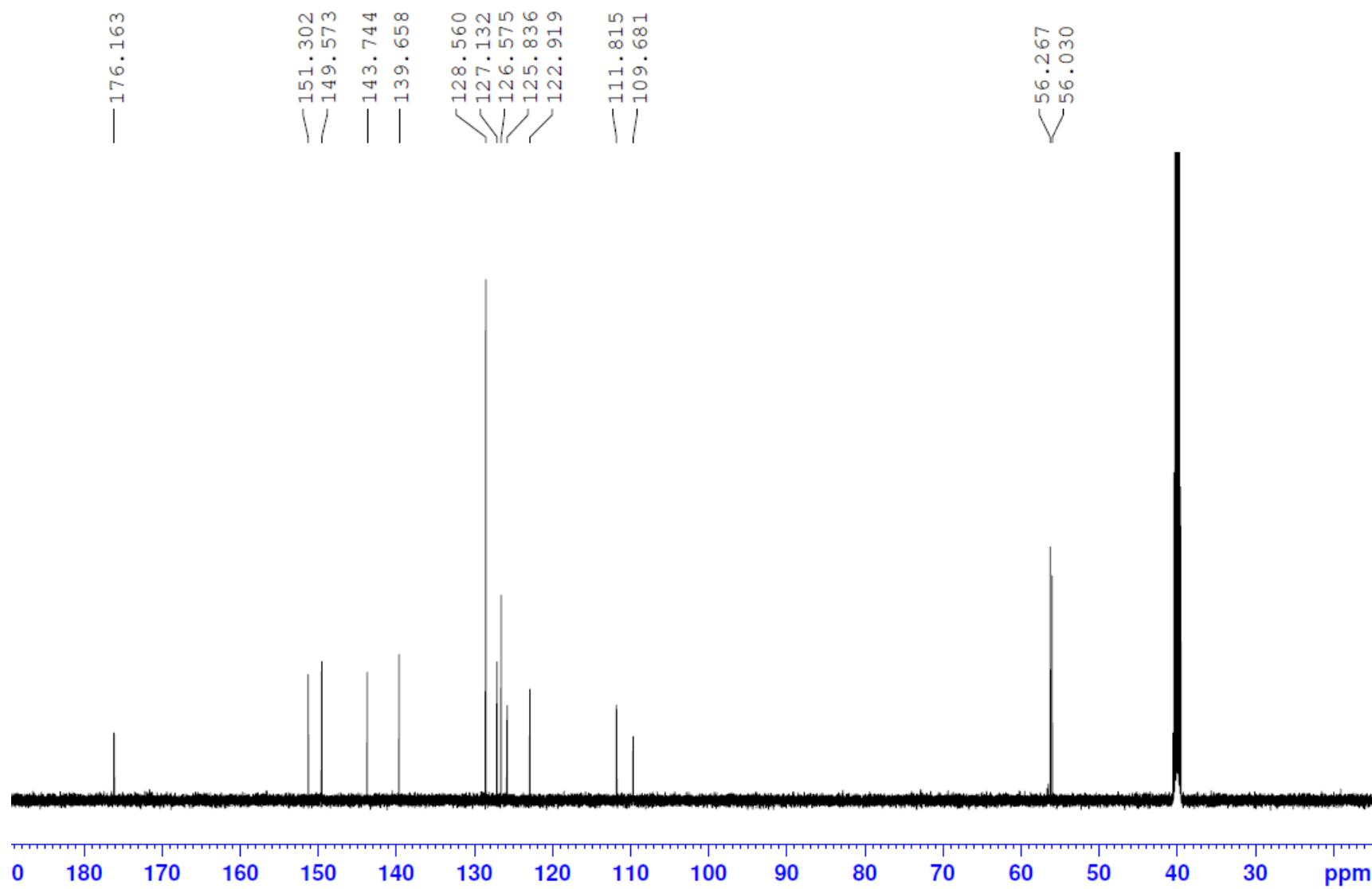
T10	 M=349,84	<p>C₁₆H₁₆N₃O₂SCl</p> <p>Calculated monoisotopic mass: 349.0652</p> <p>Measured monoisotopic mass: 349.0653</p> <p>Mass error: 0.35 ppm</p>	
T11	 M=303,80	<p>C₁₅H₁₄N₃SCl</p> <p>Calculated monoisotopic mass: 303.0597</p> <p>Measured monoisotopic mass: 303.0601</p> <p>Mass error: 1.31 ppm</p>	

T12	 $M=358.67$	<p>C14H10N3SCl3</p> <p>Calculated monoisotopic mass: 356.9661</p> <p>Measured monoisotopic mass: 356.9654</p> <p>Mass error: -1.83 ppm</p>	<p>Theoretical monoisotopic mass of $[M+H]^+$: 357.9734</p> 
T13	 $M=368.68$	<p>C14H11N3SClBr</p> <p>Calculated monoisotopic mass: 366.9546</p> <p>Measured monoisotopic mass: 366.9551</p> <p>Mass error: 1.46 ppm</p>	<p>Theoretical monoisotopic mass of $[M+H]^+$: 367.9618</p> 

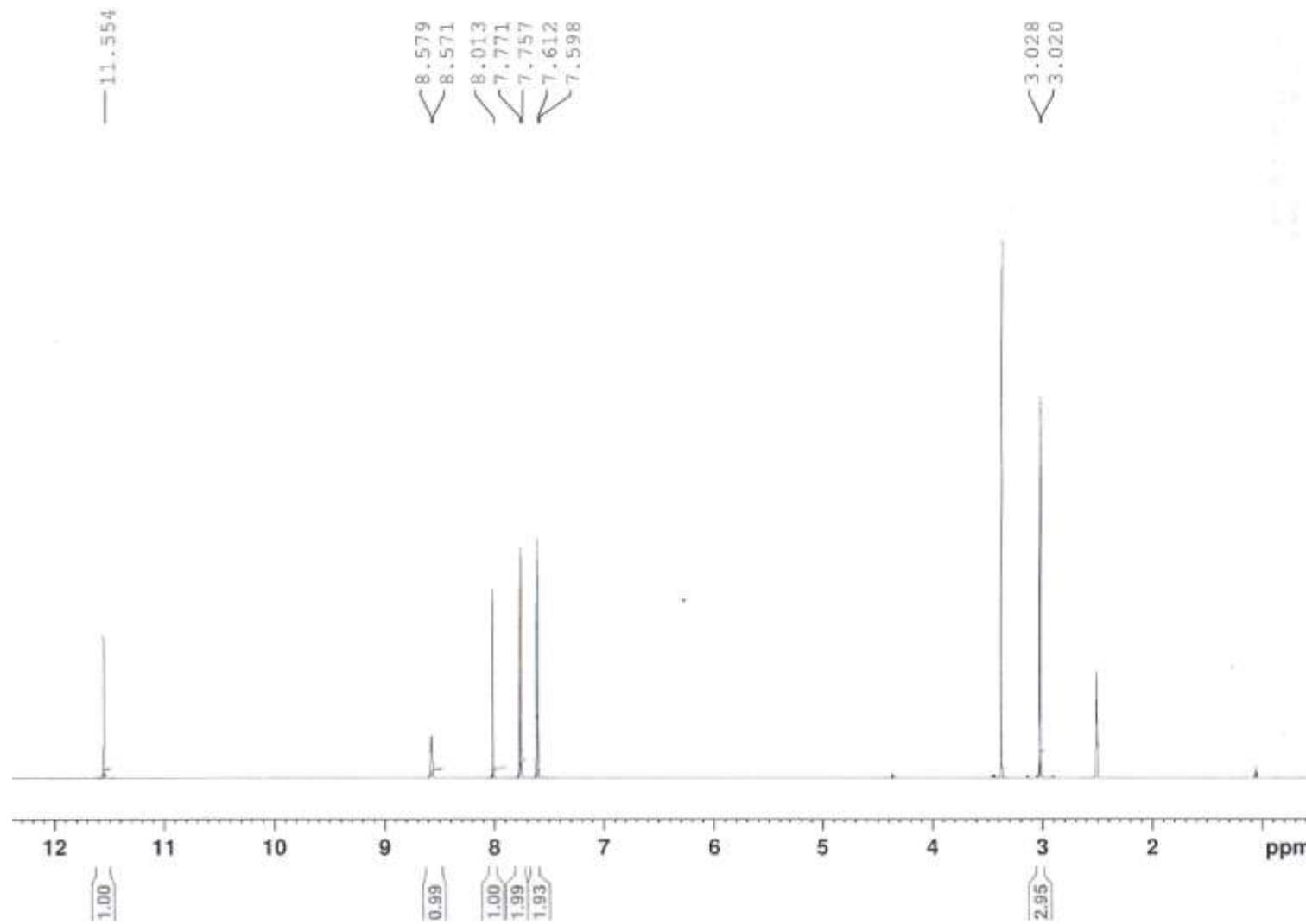
T17	<p>M=358,67</p>	<p>C14H10N3SCl3 Calculated monoisotopic mass: 356.9661 Measured monoisotopic mass: 356.9664 Mass error: 0.96 ppm</p>	<p>Theoretical monoisotopic mass of [M+H]⁺: 357.9734</p> <p>x10⁶ Cpd 1: C14 H10 Cl3 N3 S: +ESI Scan (rt: 5.144-5.161 min, 2 scans) Frag=350.0V T17.d Subtract</p> <table border="1"> <thead> <tr> <th>m/z</th> <th>Counts (approx.)</th> </tr> </thead> <tbody> <tr><td>357.9735</td><td>1.6</td></tr> <tr><td>358.9763</td><td>0.3</td></tr> <tr><td>359.9710</td><td>1.7</td></tr> <tr><td>360.9737</td><td>0.3</td></tr> <tr><td>361.9682</td><td>0.6</td></tr> <tr><td>362.9704</td><td>0.1</td></tr> <tr><td>363.9650</td><td>0.1</td></tr> <tr><td>364.9677</td><td>0.1</td></tr> <tr><td>365.9612</td><td>0.1</td></tr> </tbody> </table> <p>Counts vs. Mass-to-Charge (m/z)</p>	m/z	Counts (approx.)	357.9735	1.6	358.9763	0.3	359.9710	1.7	360.9737	0.3	361.9682	0.6	362.9704	0.1	363.9650	0.1	364.9677	0.1	365.9612	0.1
m/z	Counts (approx.)																						
357.9735	1.6																						
358.9763	0.3																						
359.9710	1.7																						
360.9737	0.3																						
361.9682	0.6																						
362.9704	0.1																						
363.9650	0.1																						
364.9677	0.1																						
365.9612	0.1																						

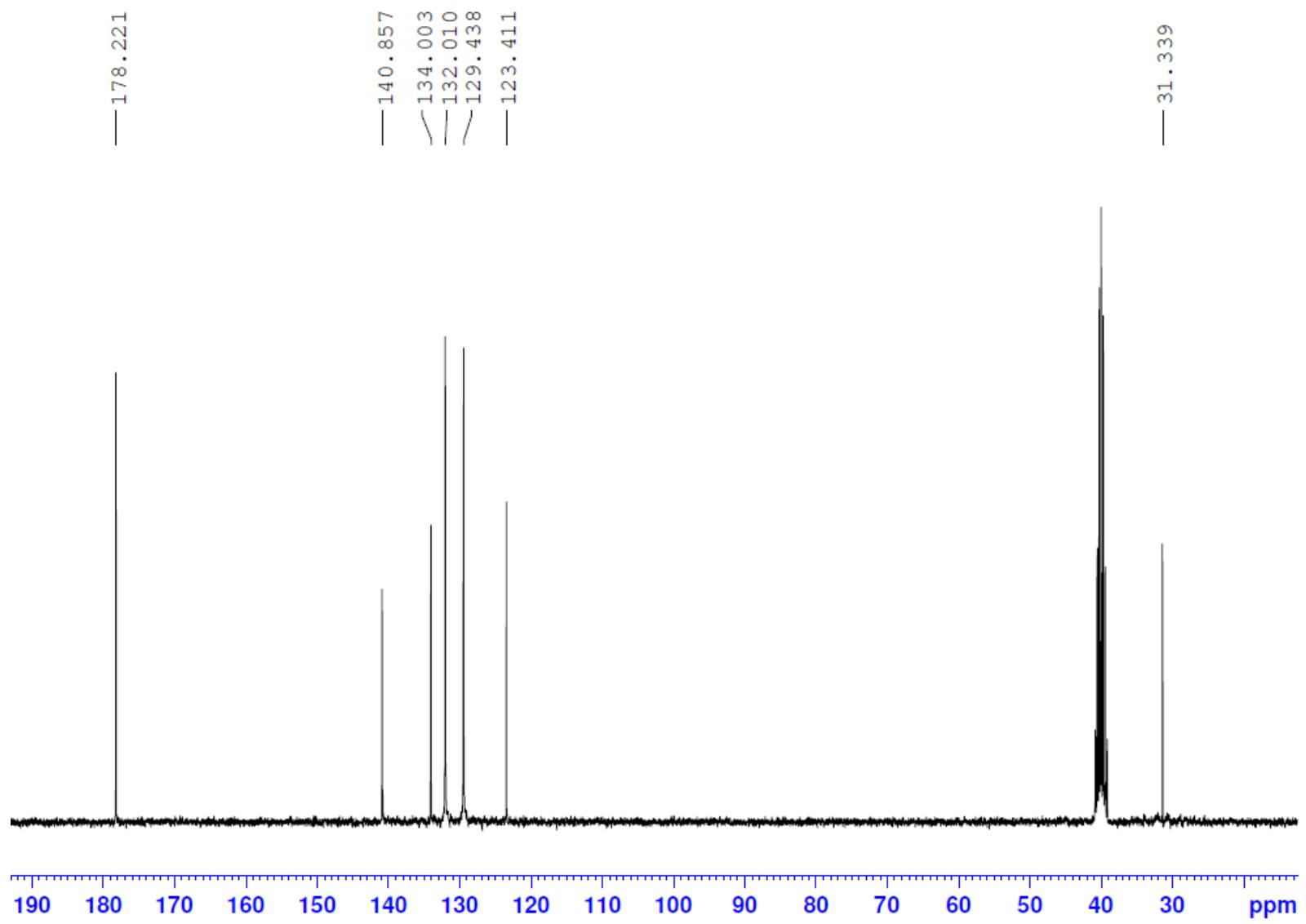
T1

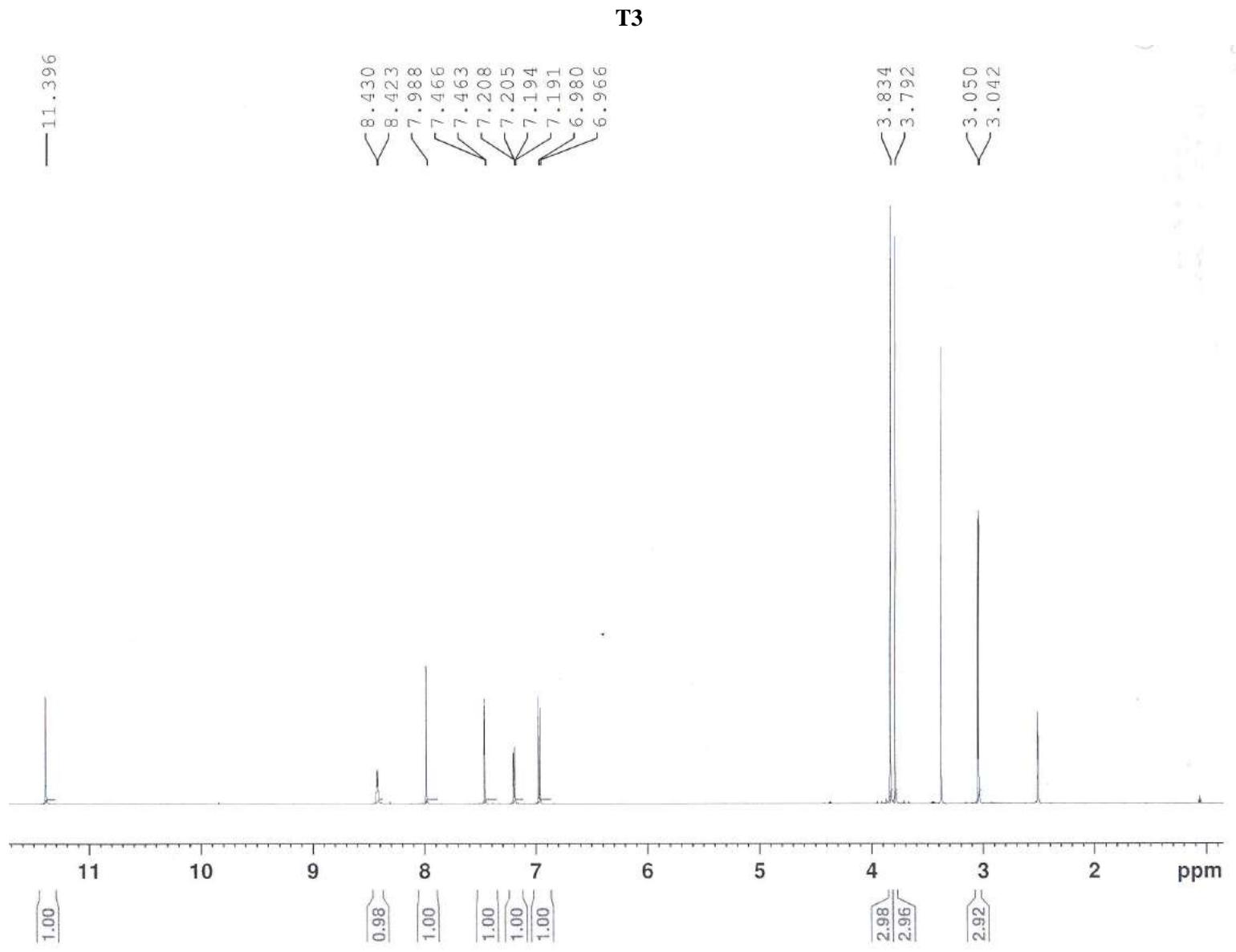


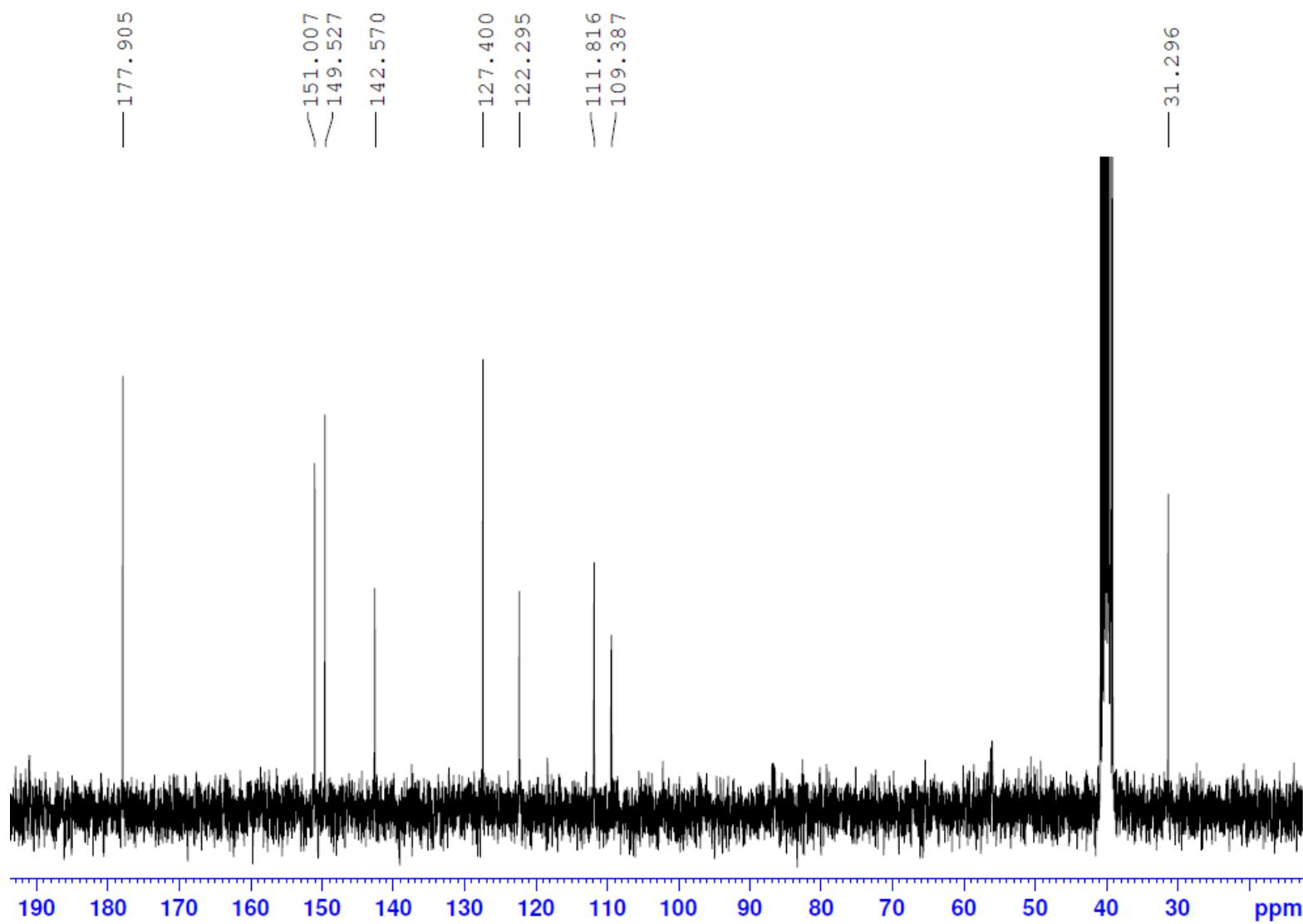


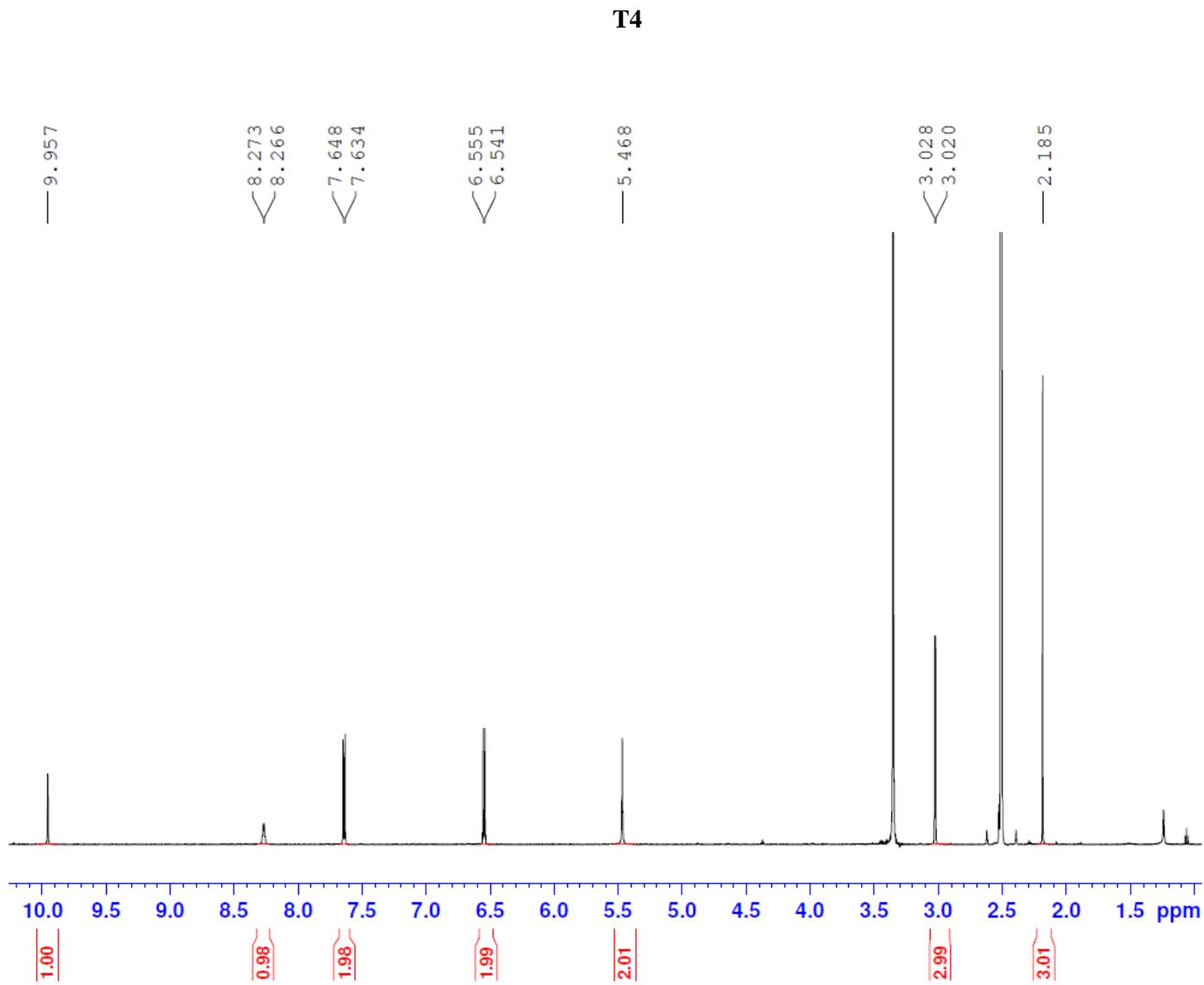
T2

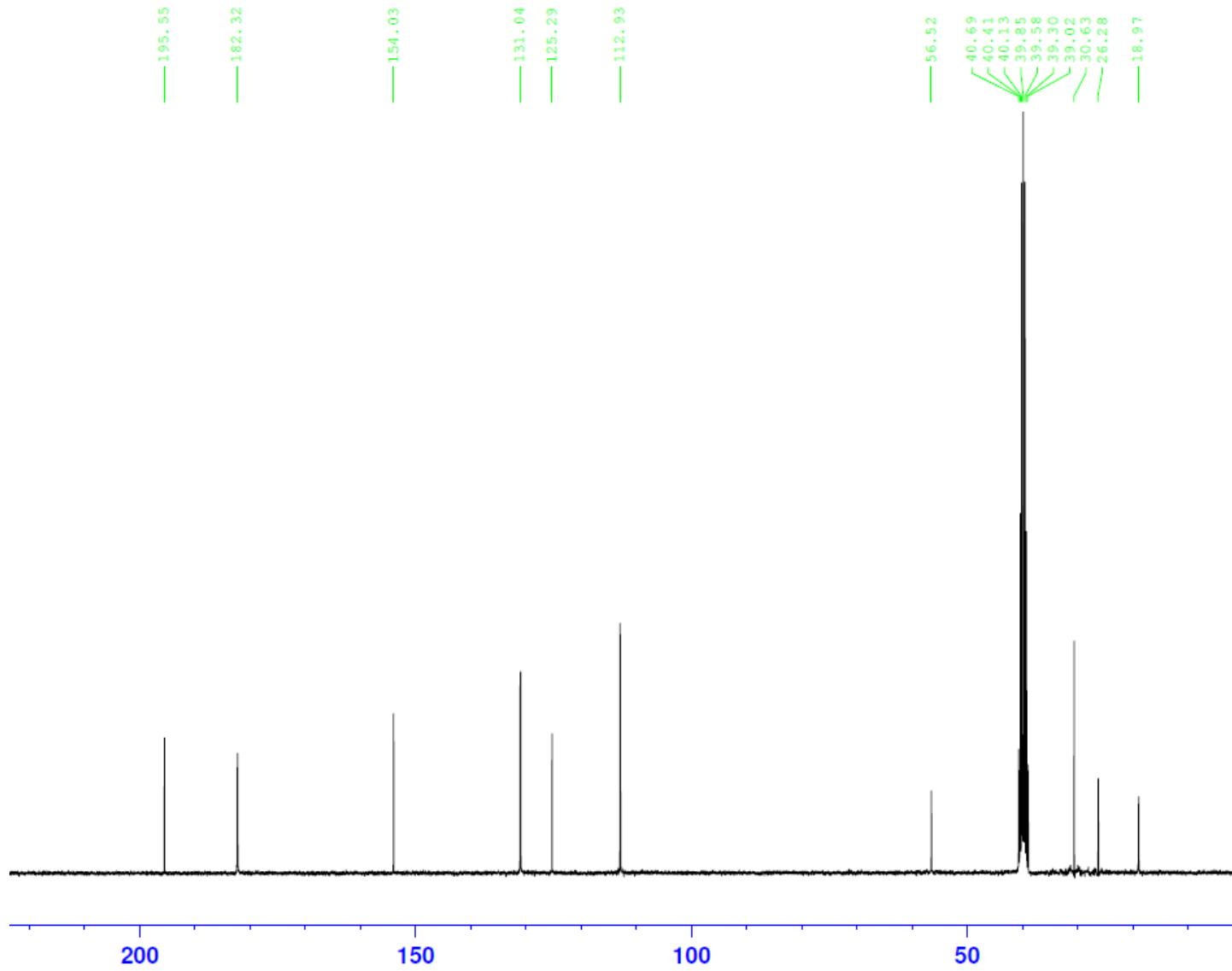




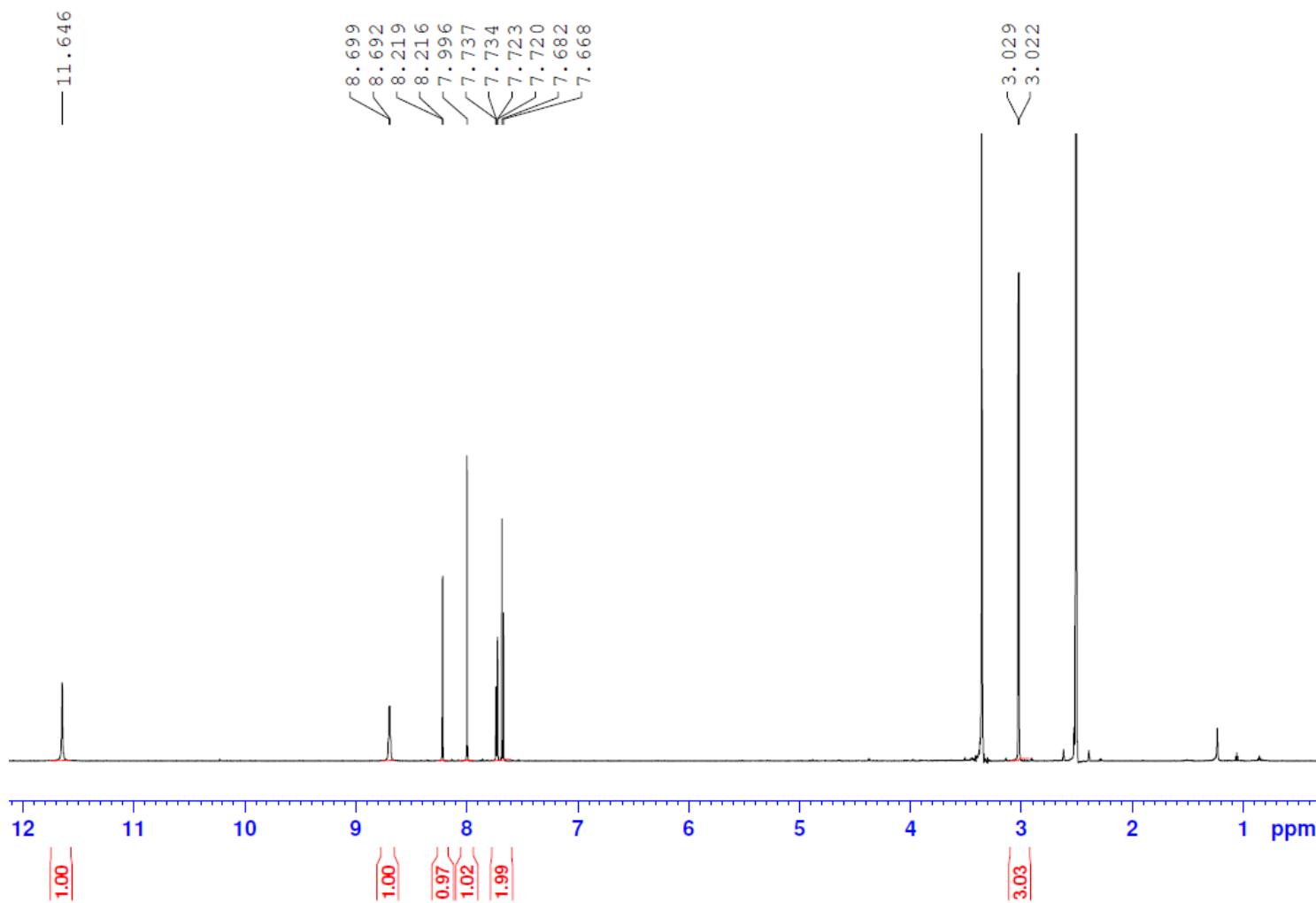


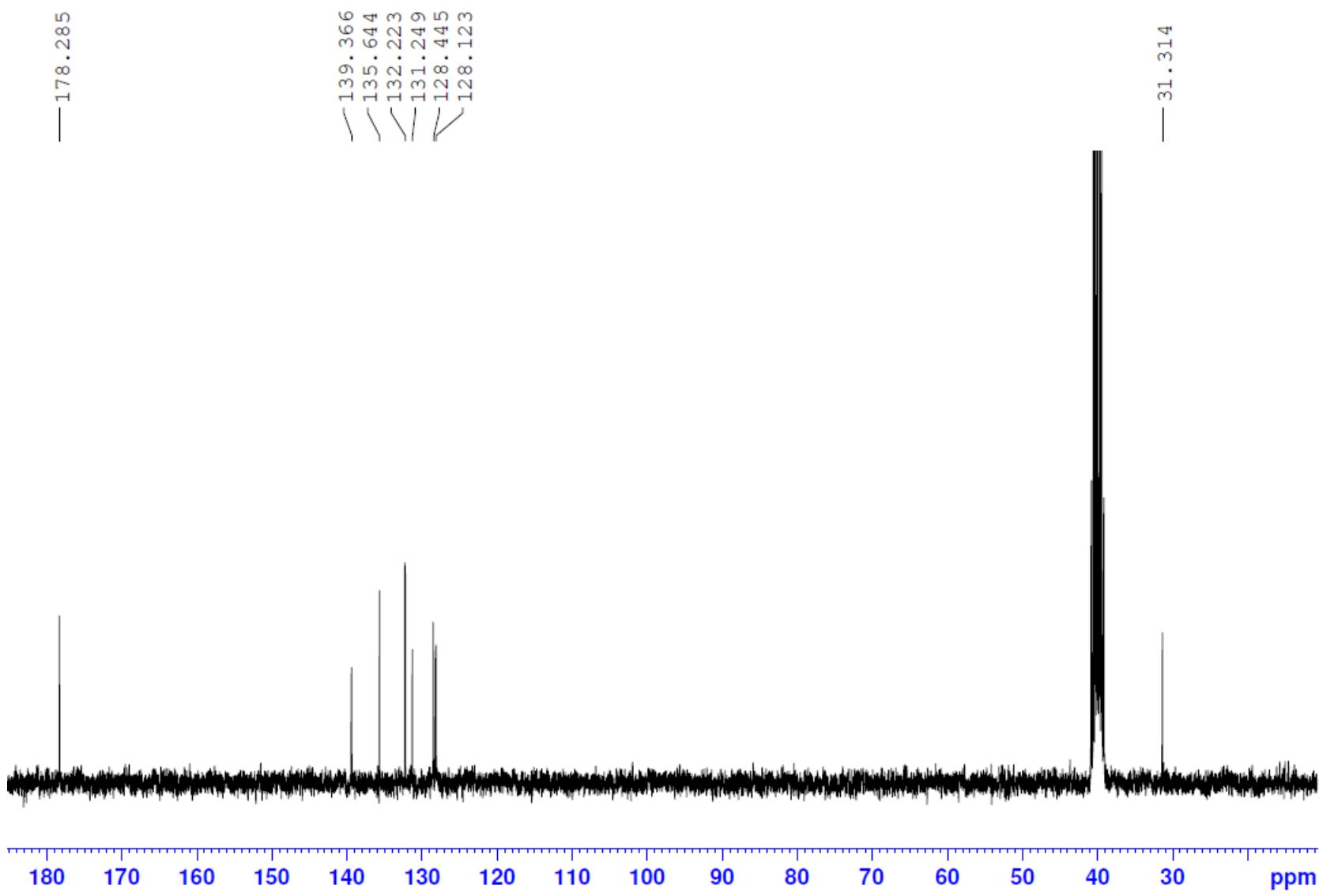


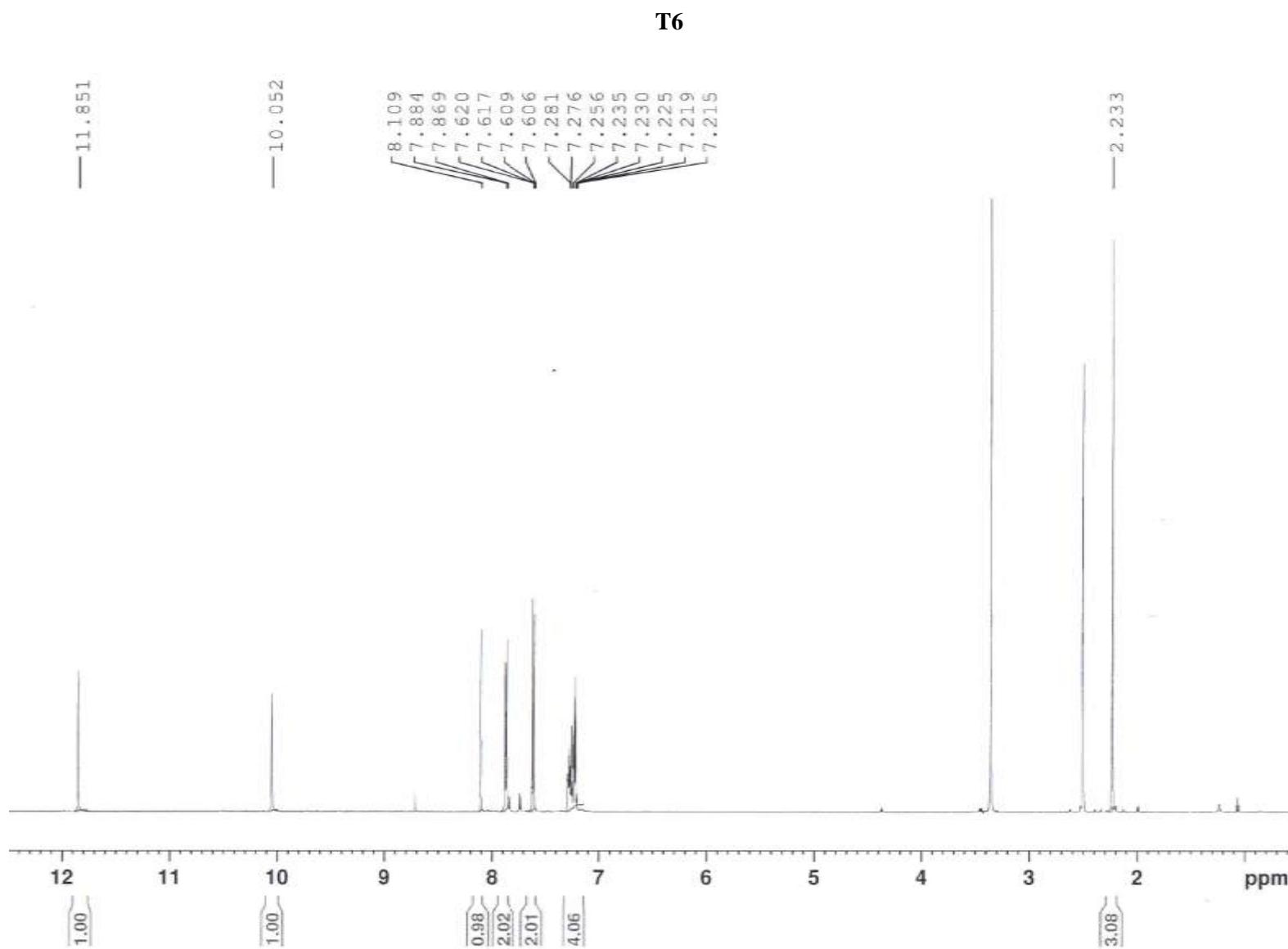


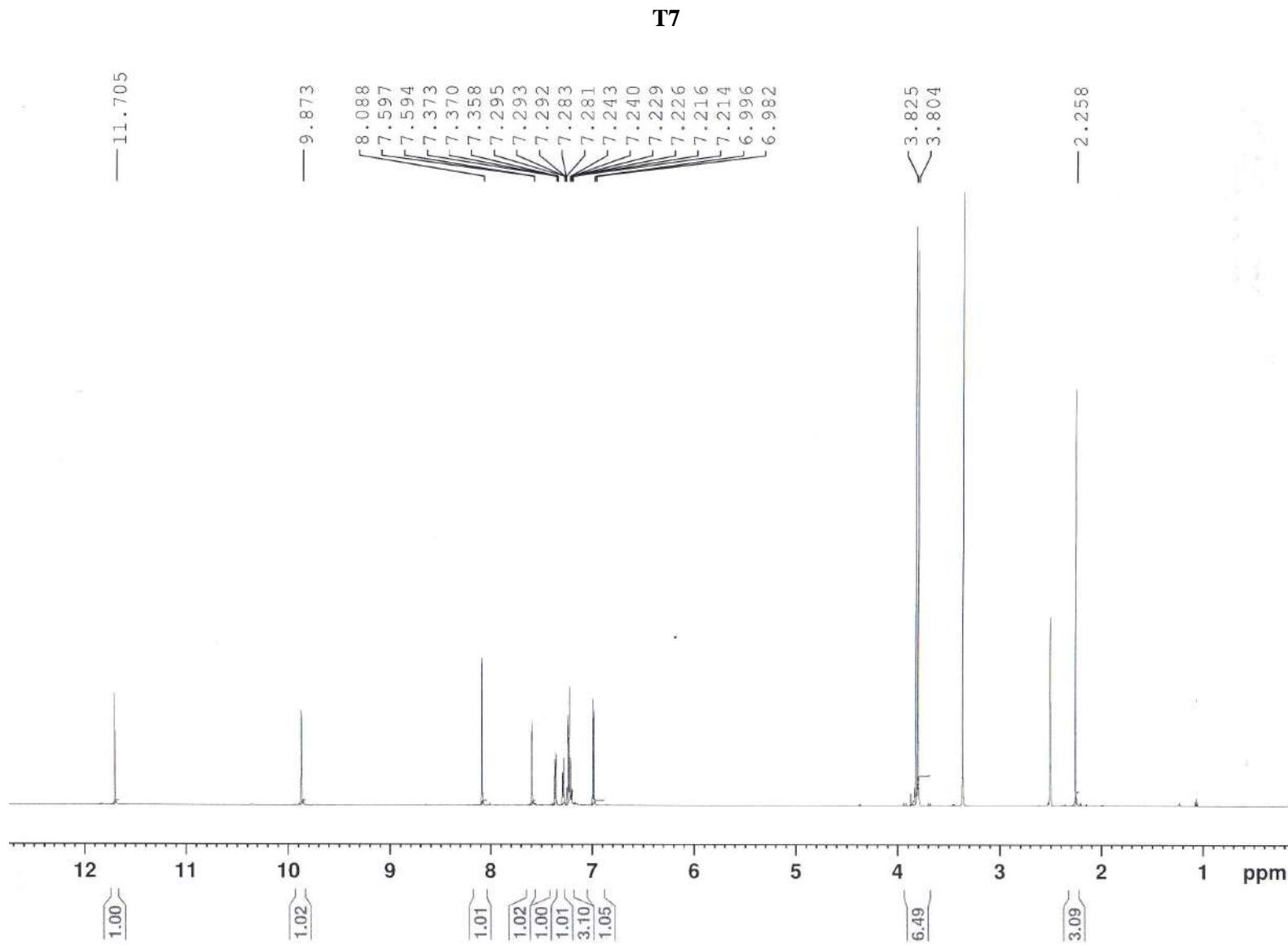


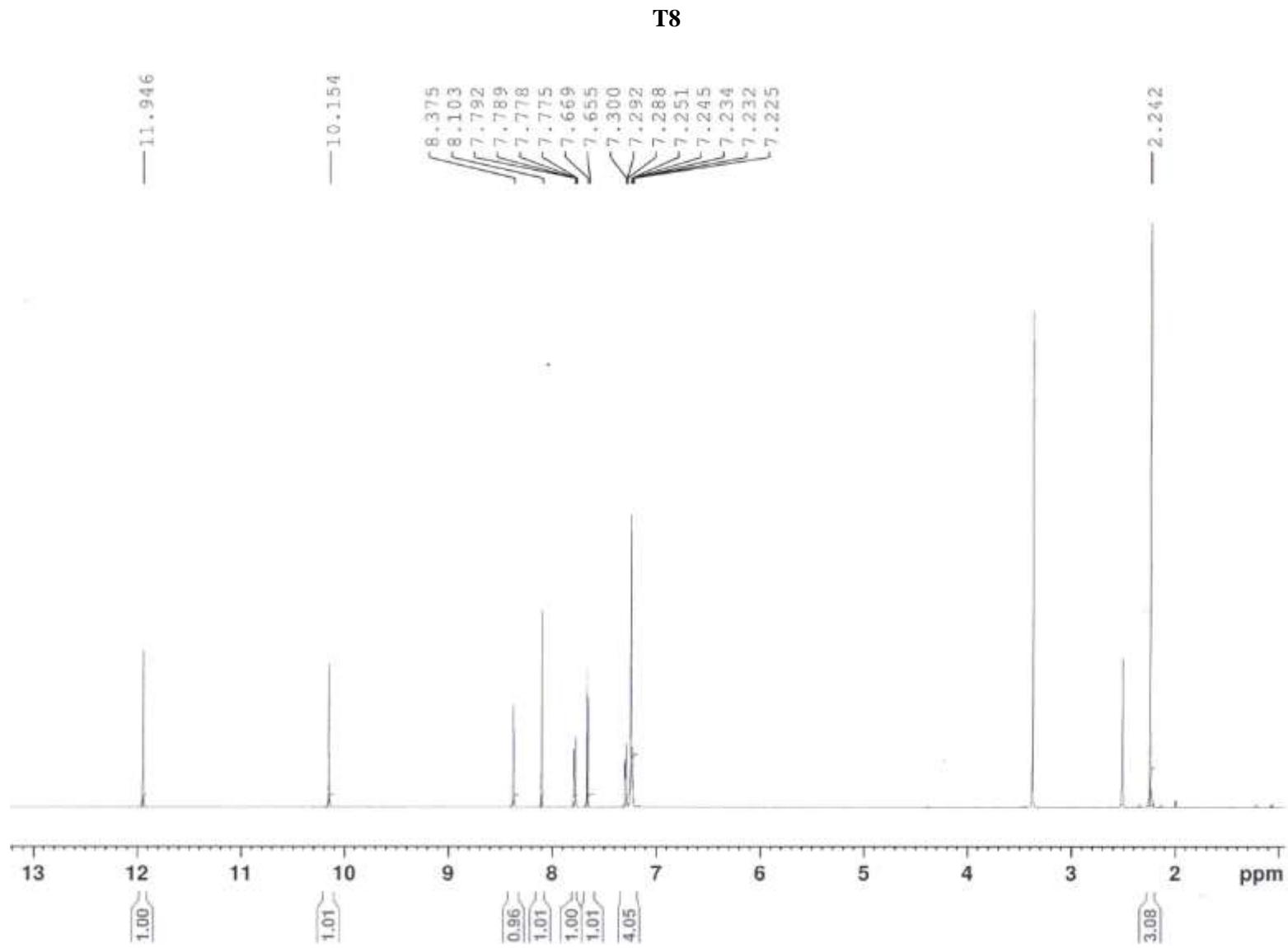
T5



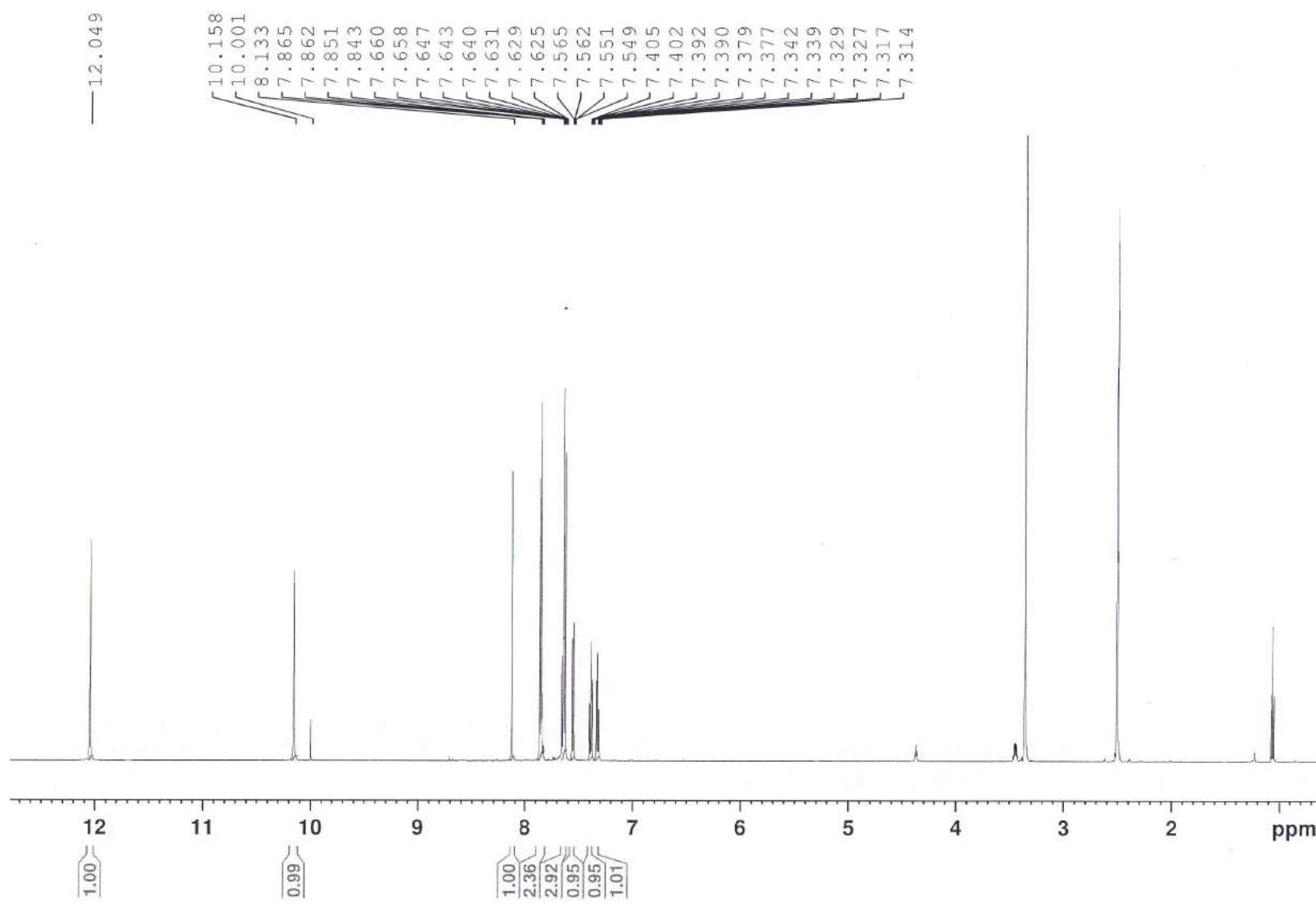


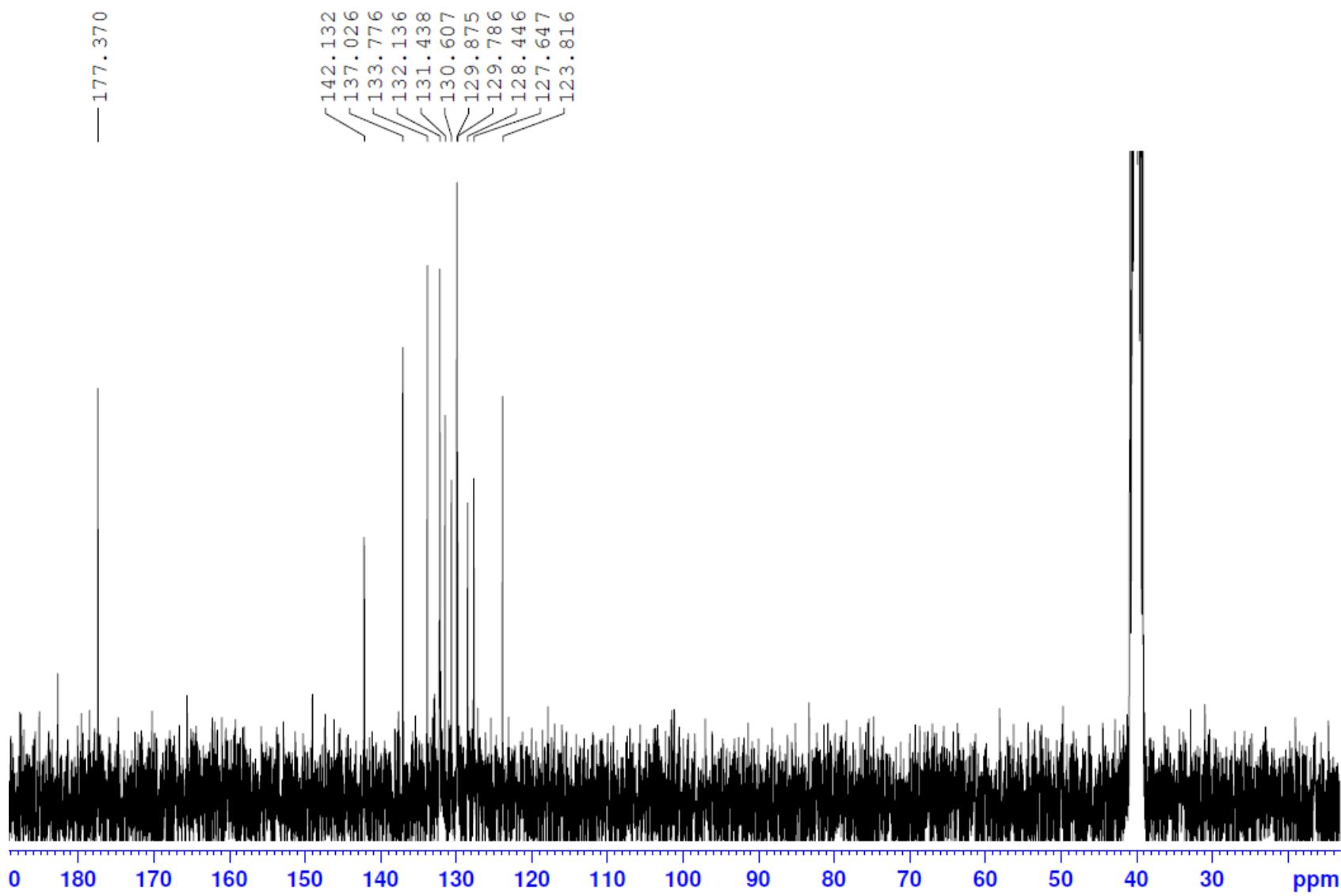




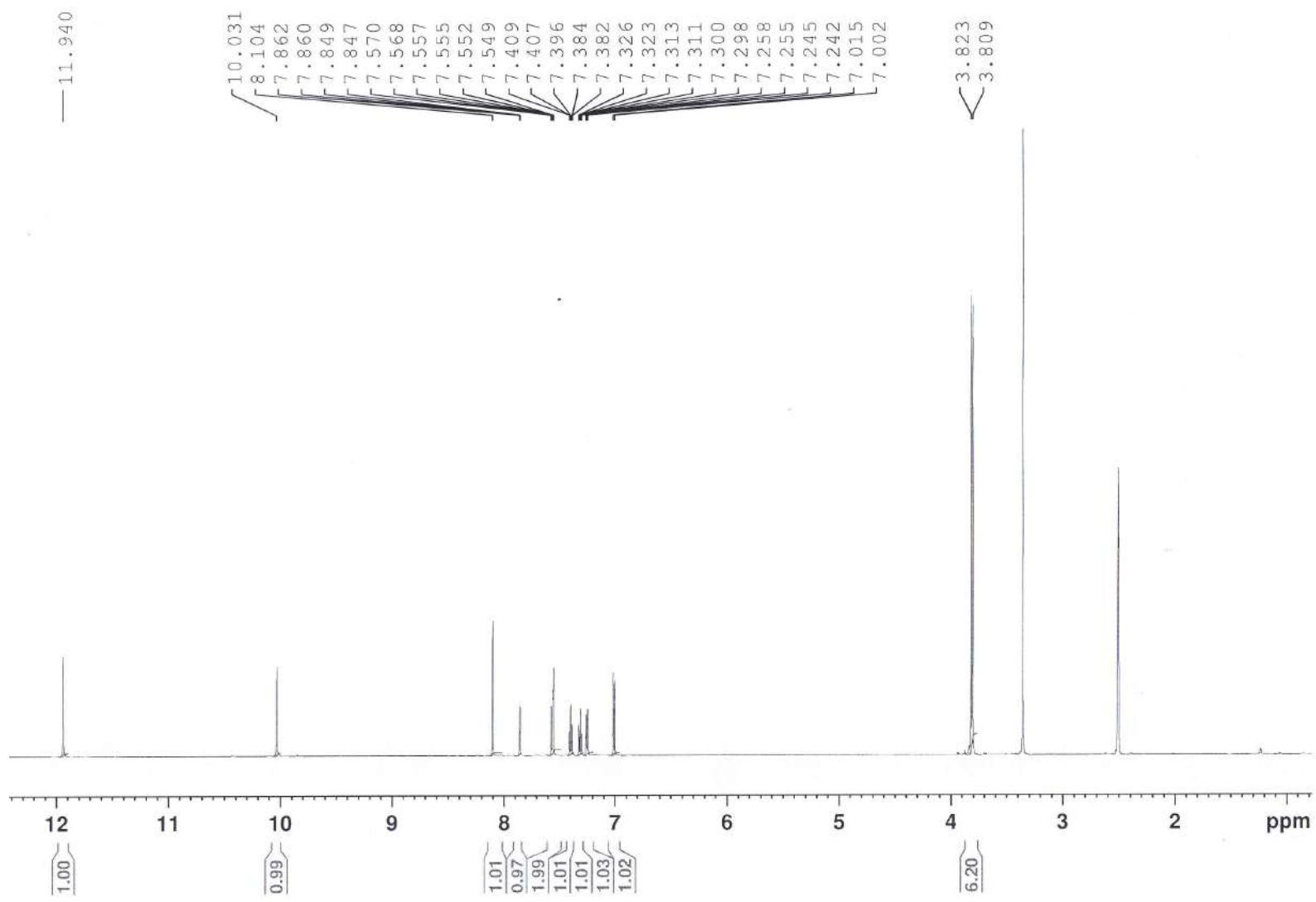


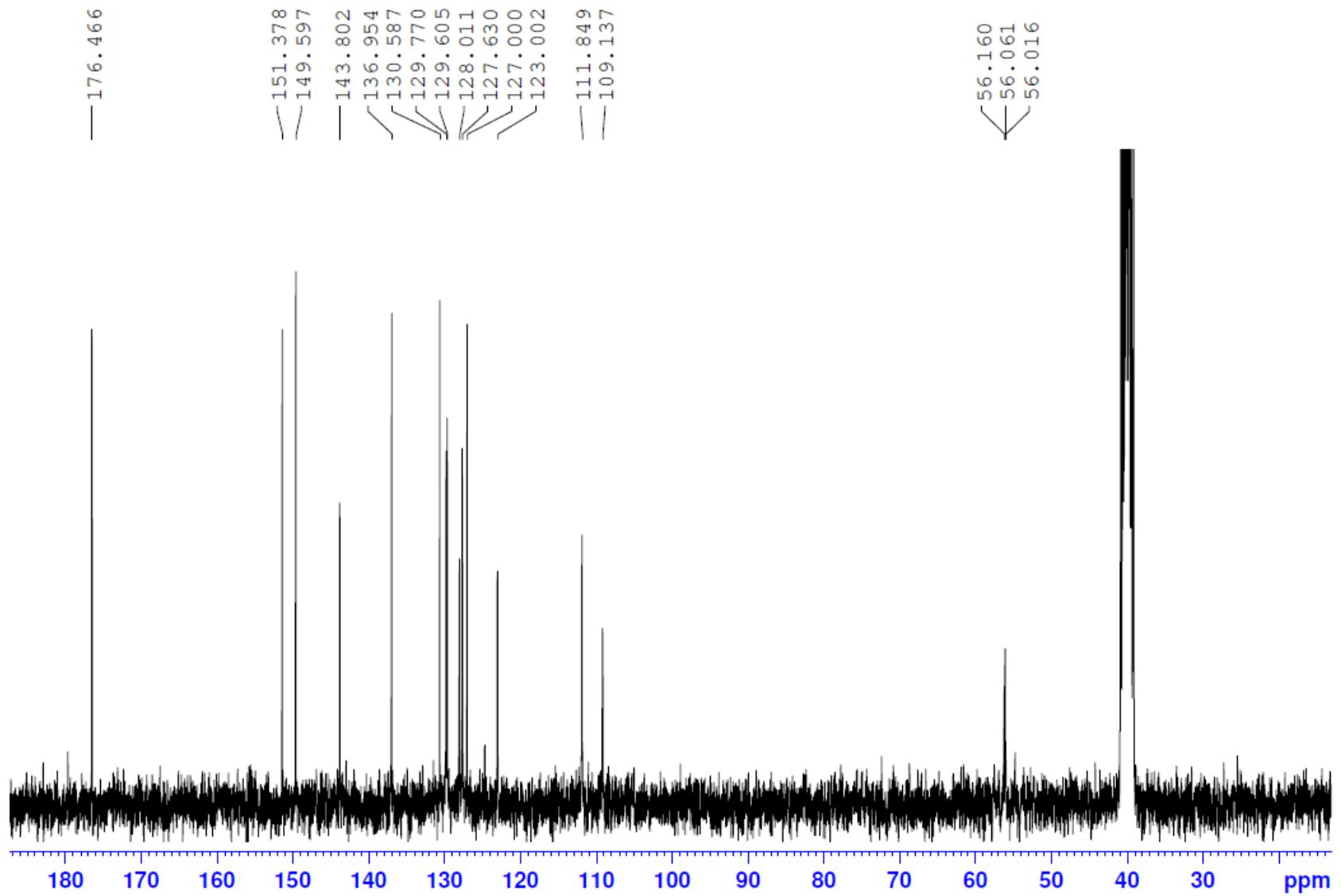
T9



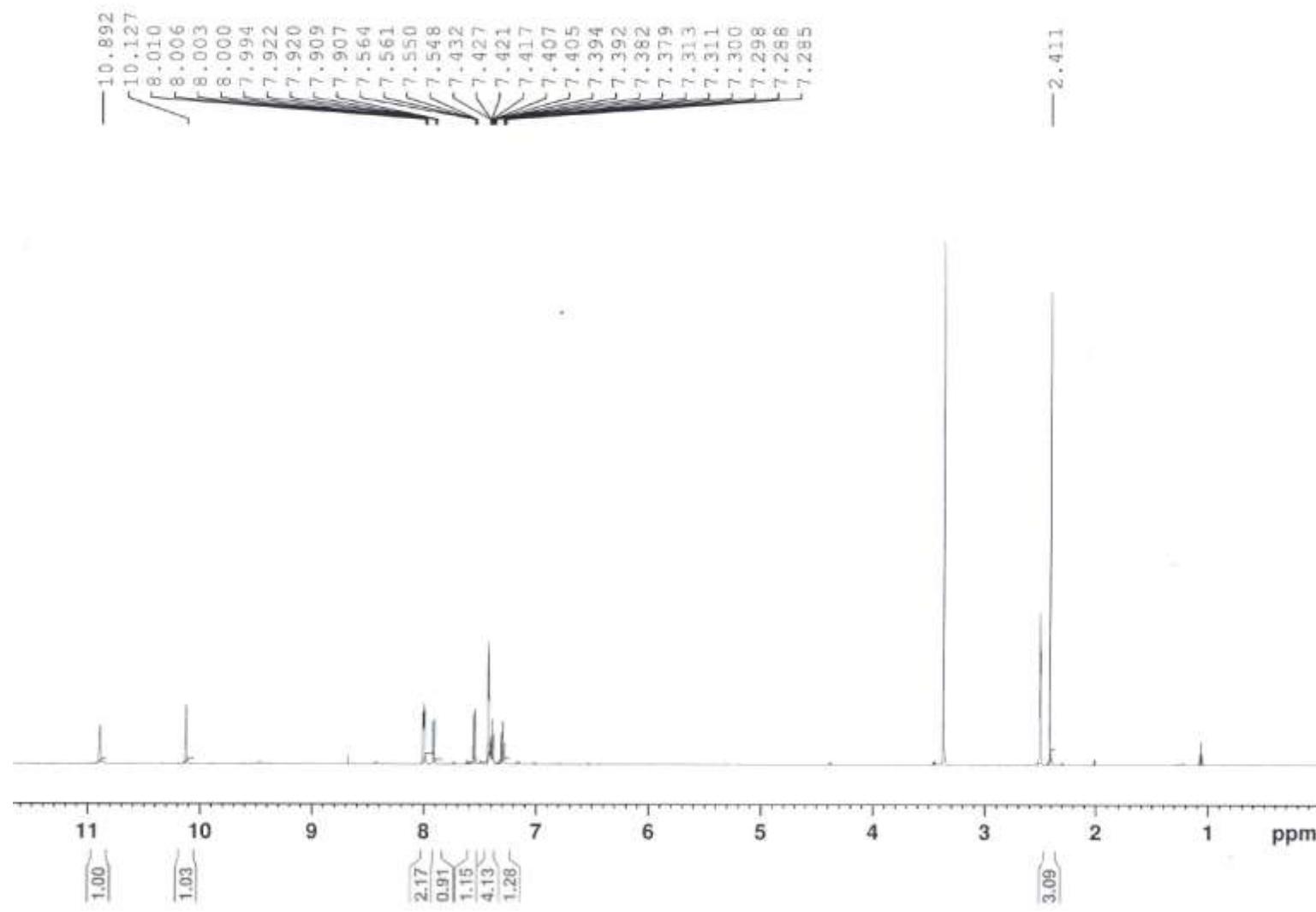


T10

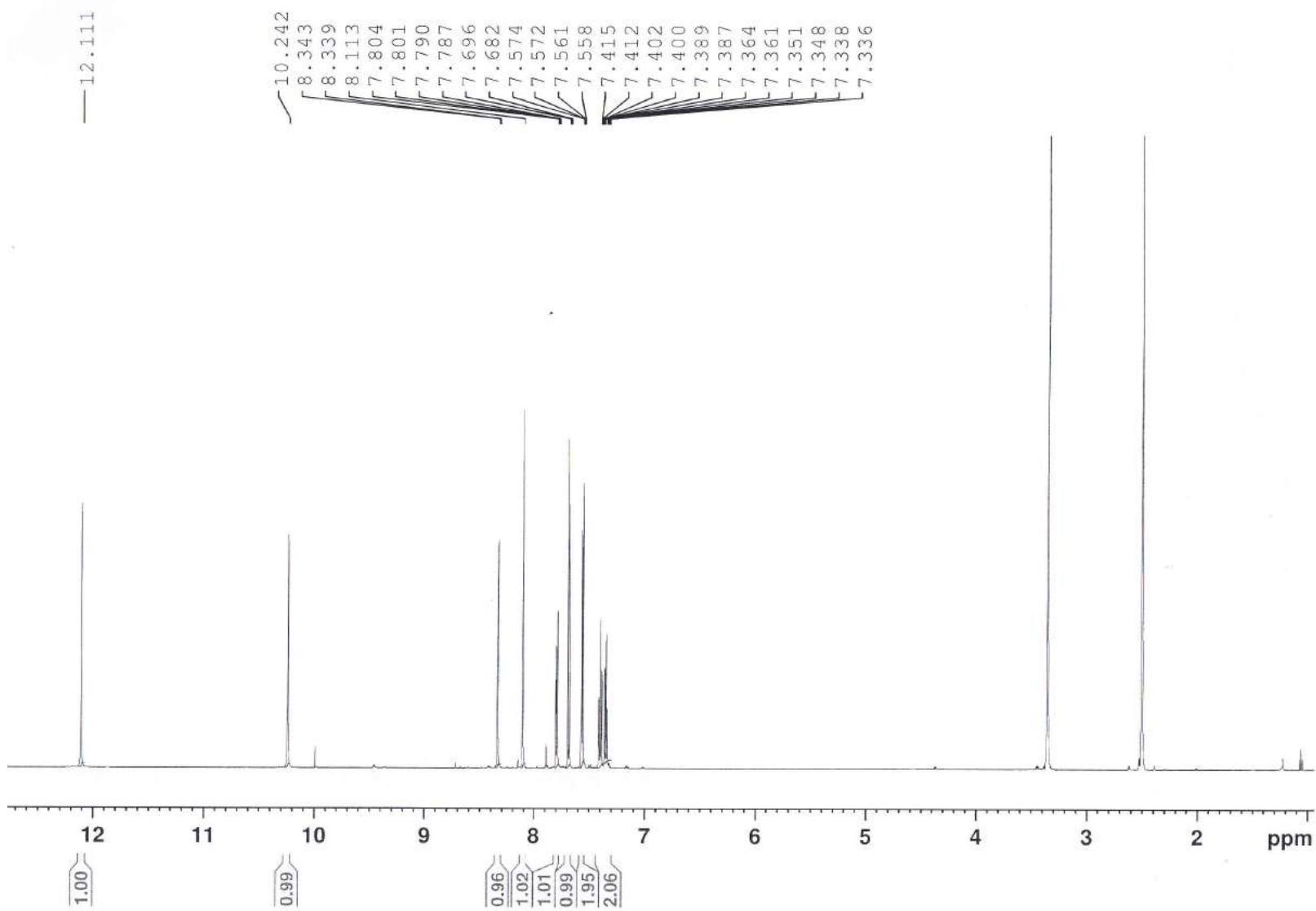


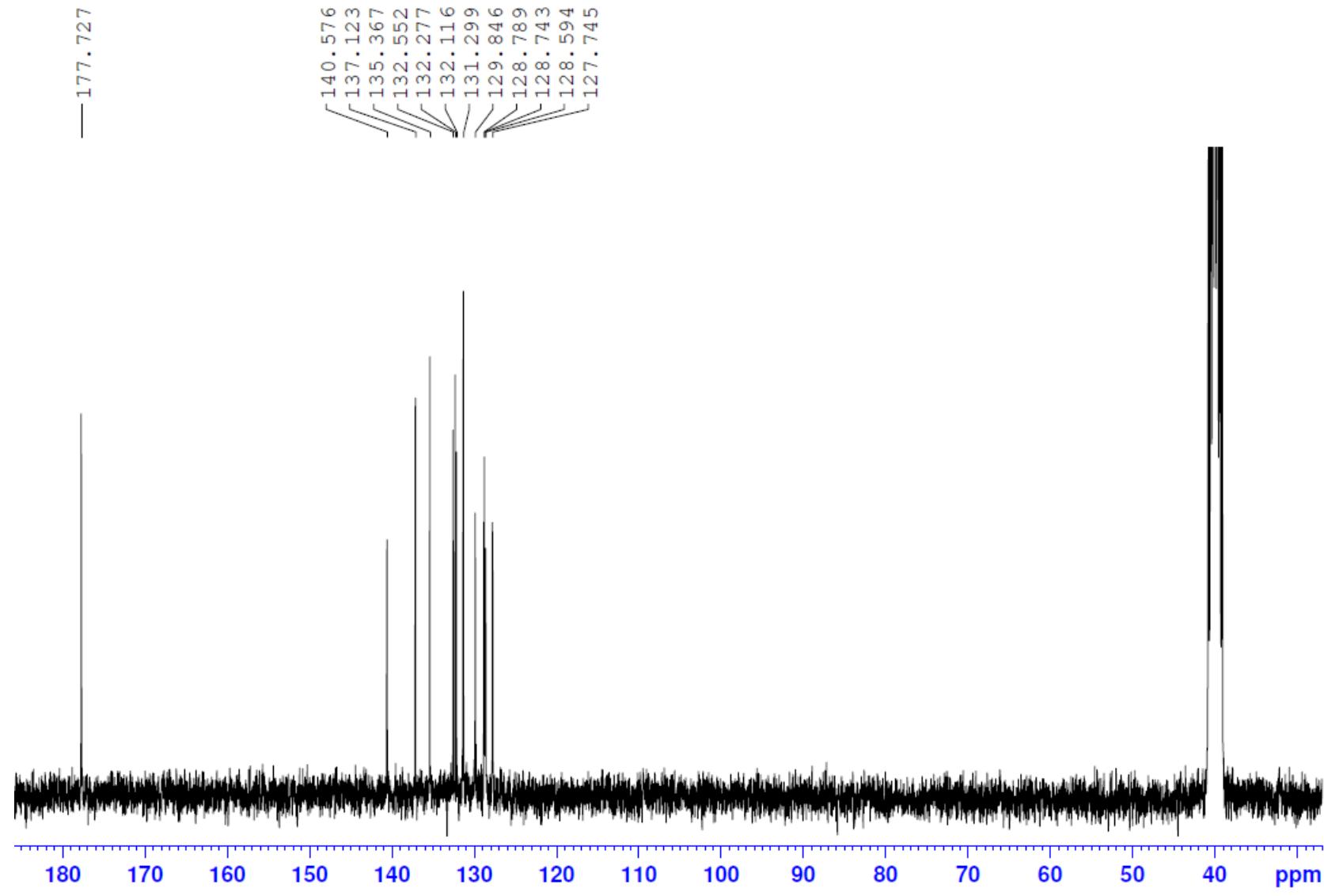


T11

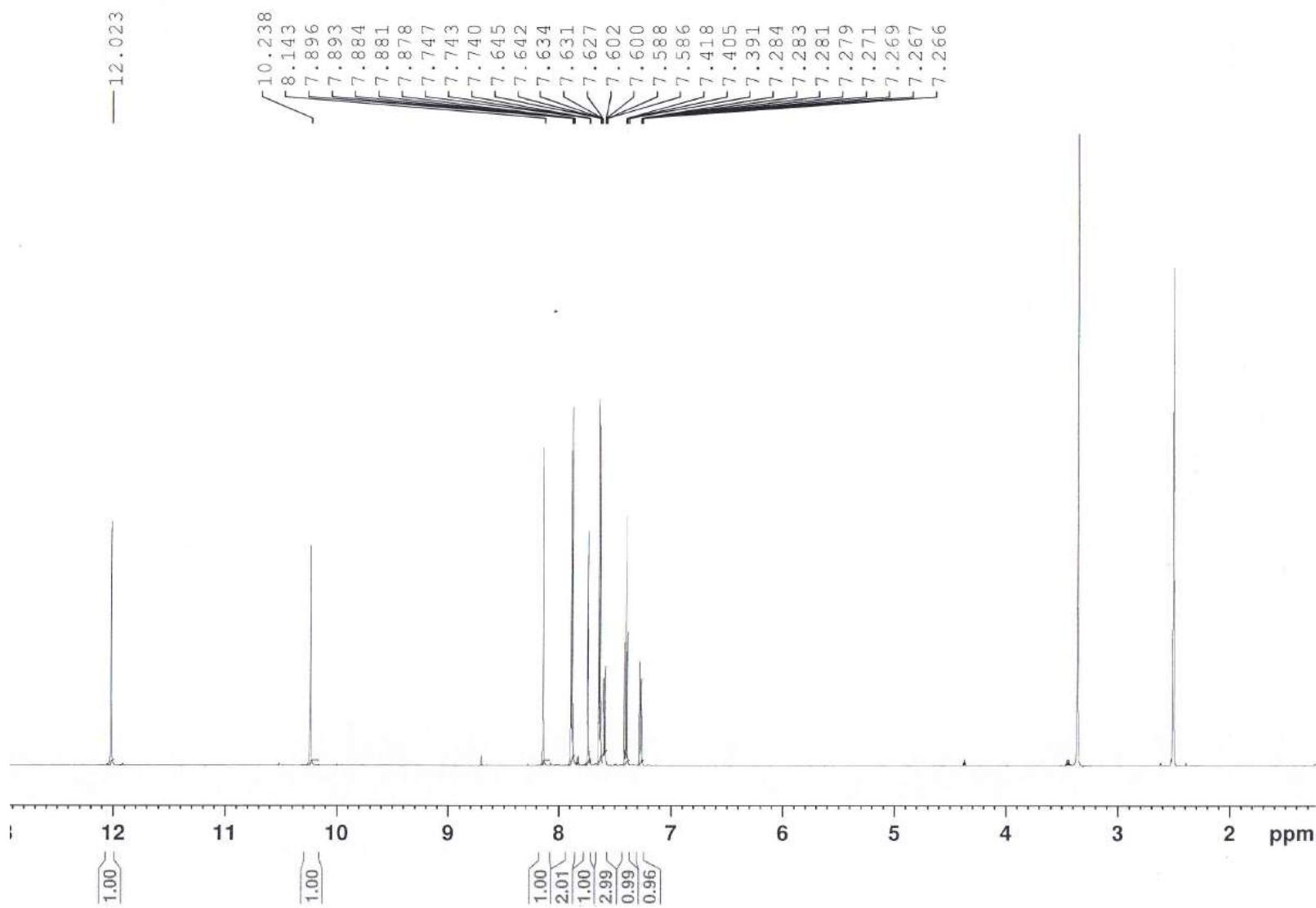


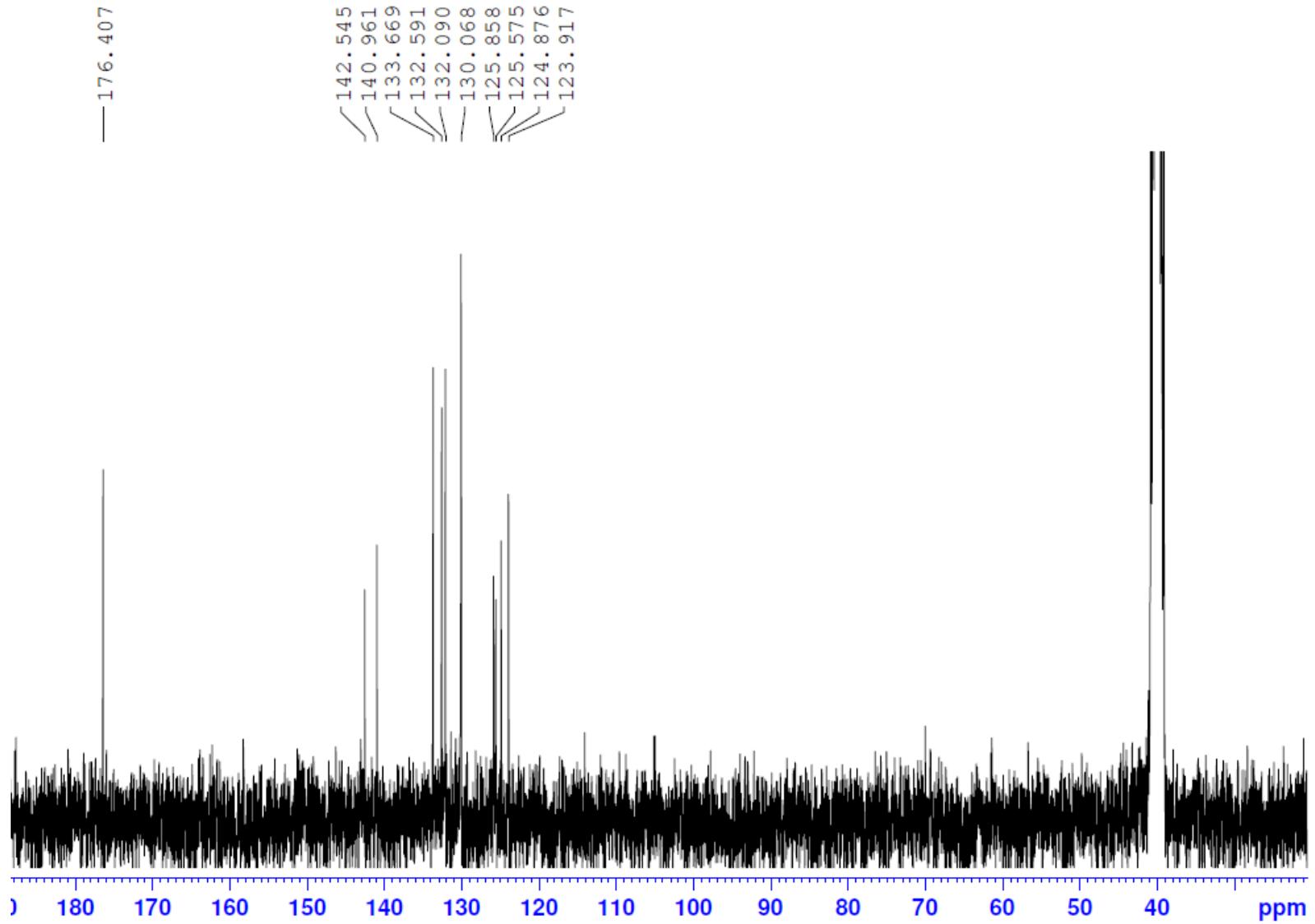
T12

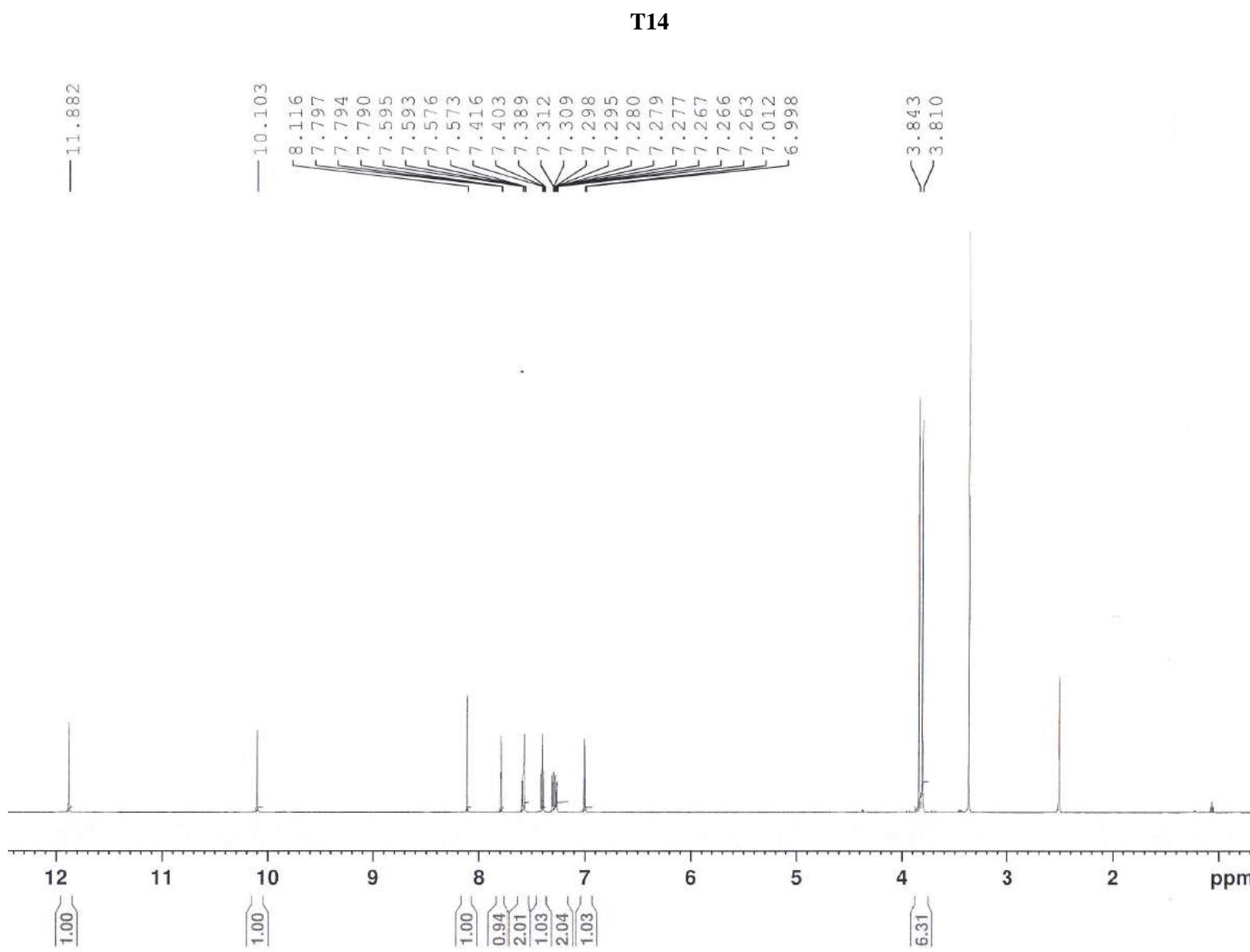


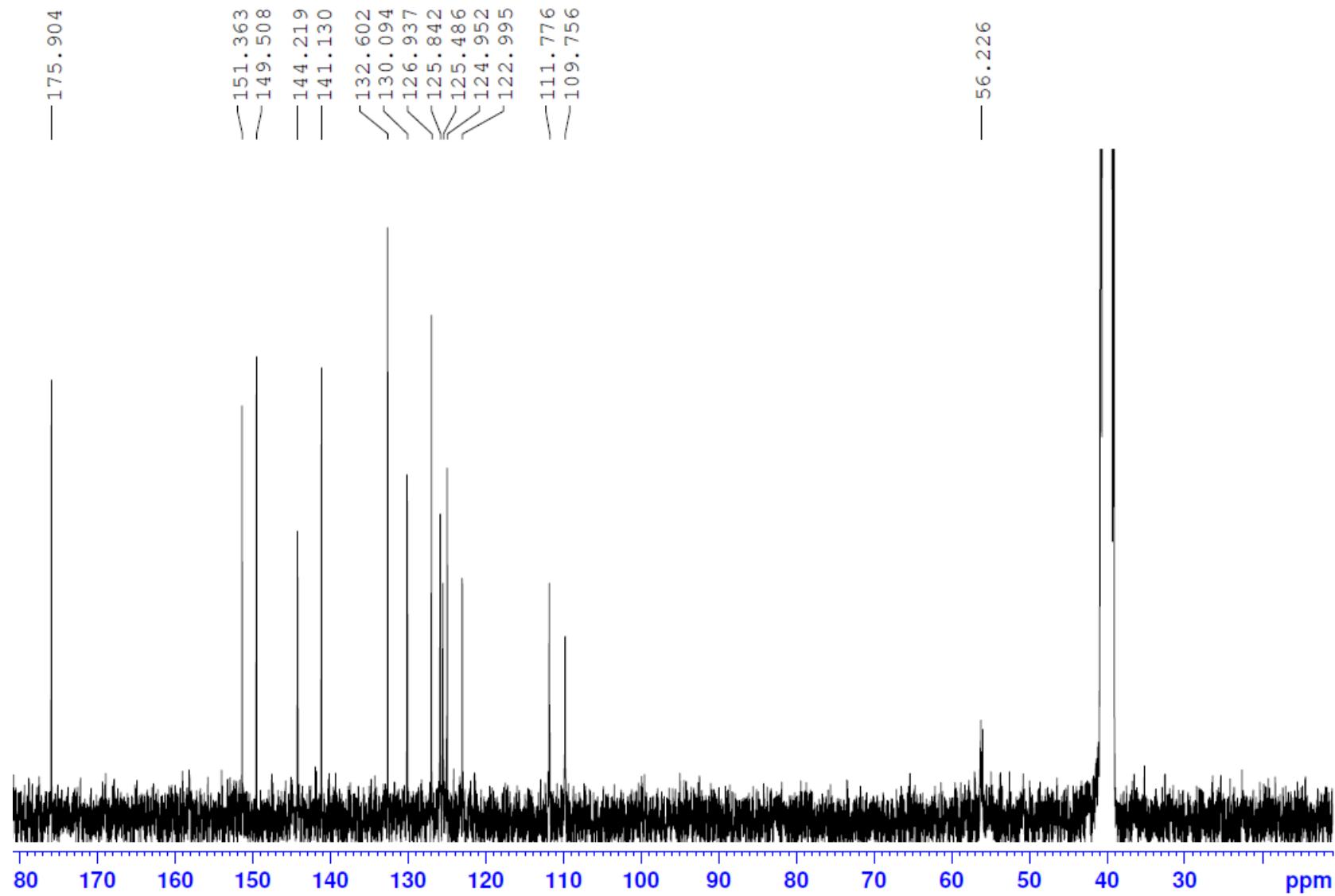


T13

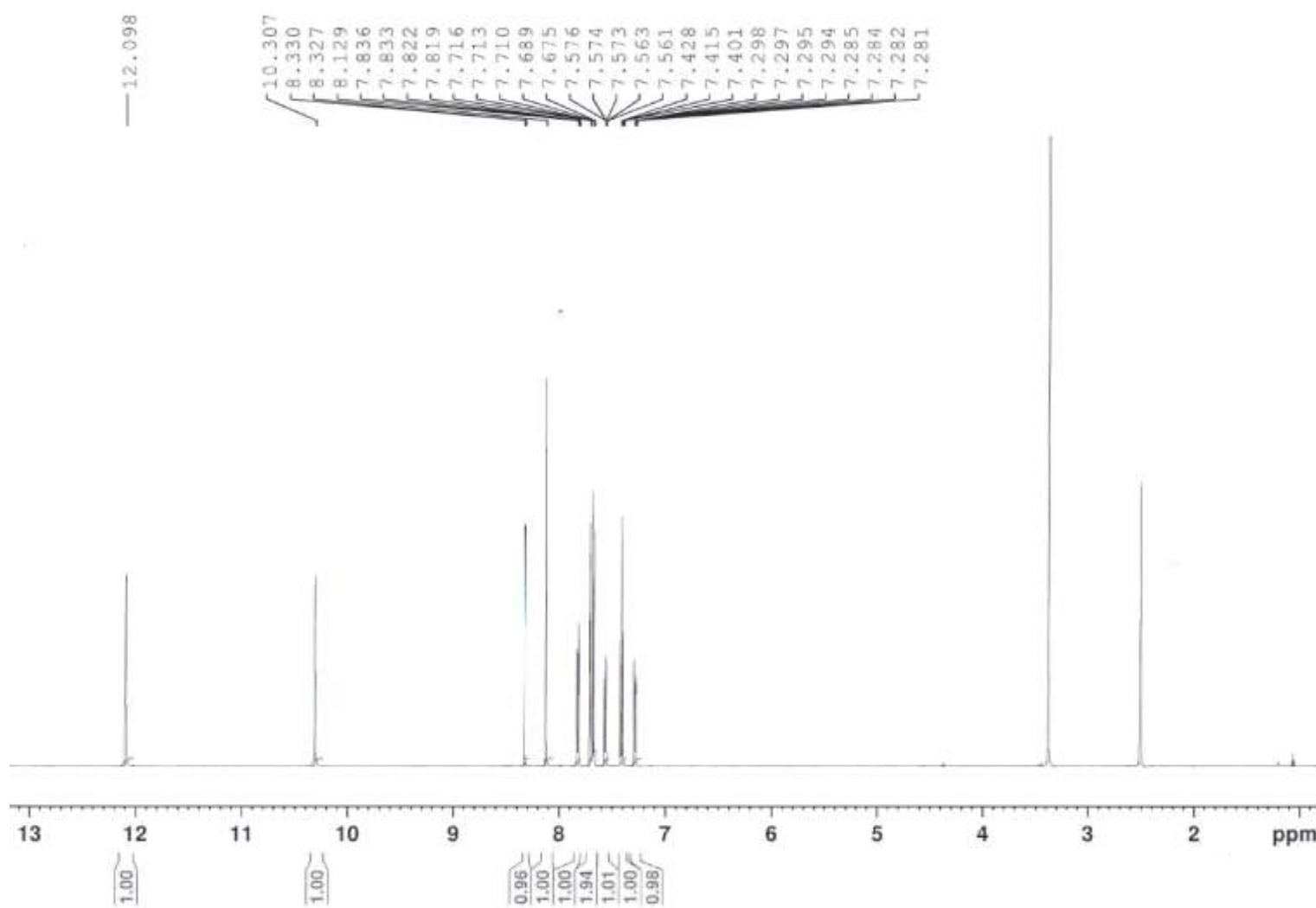




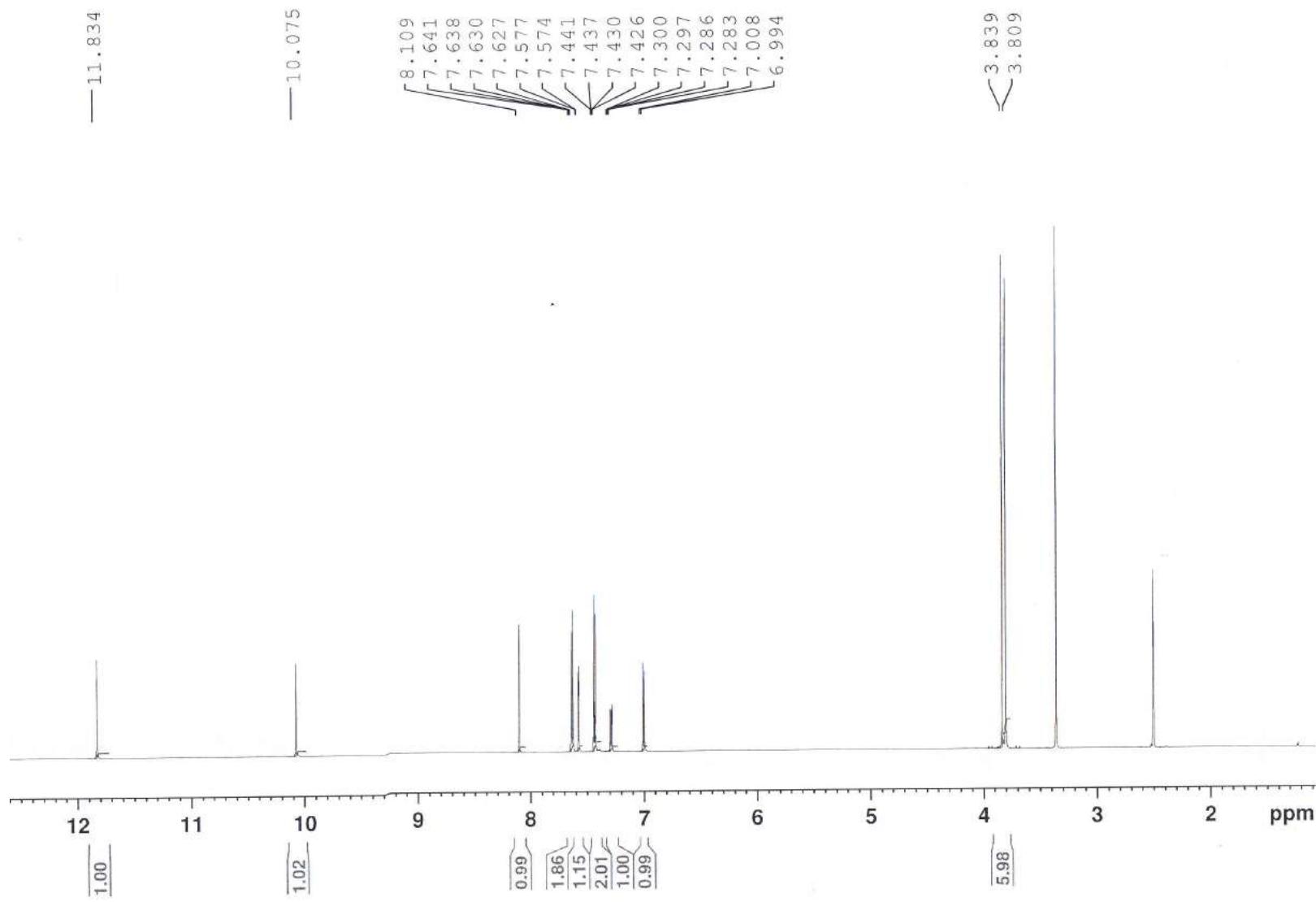


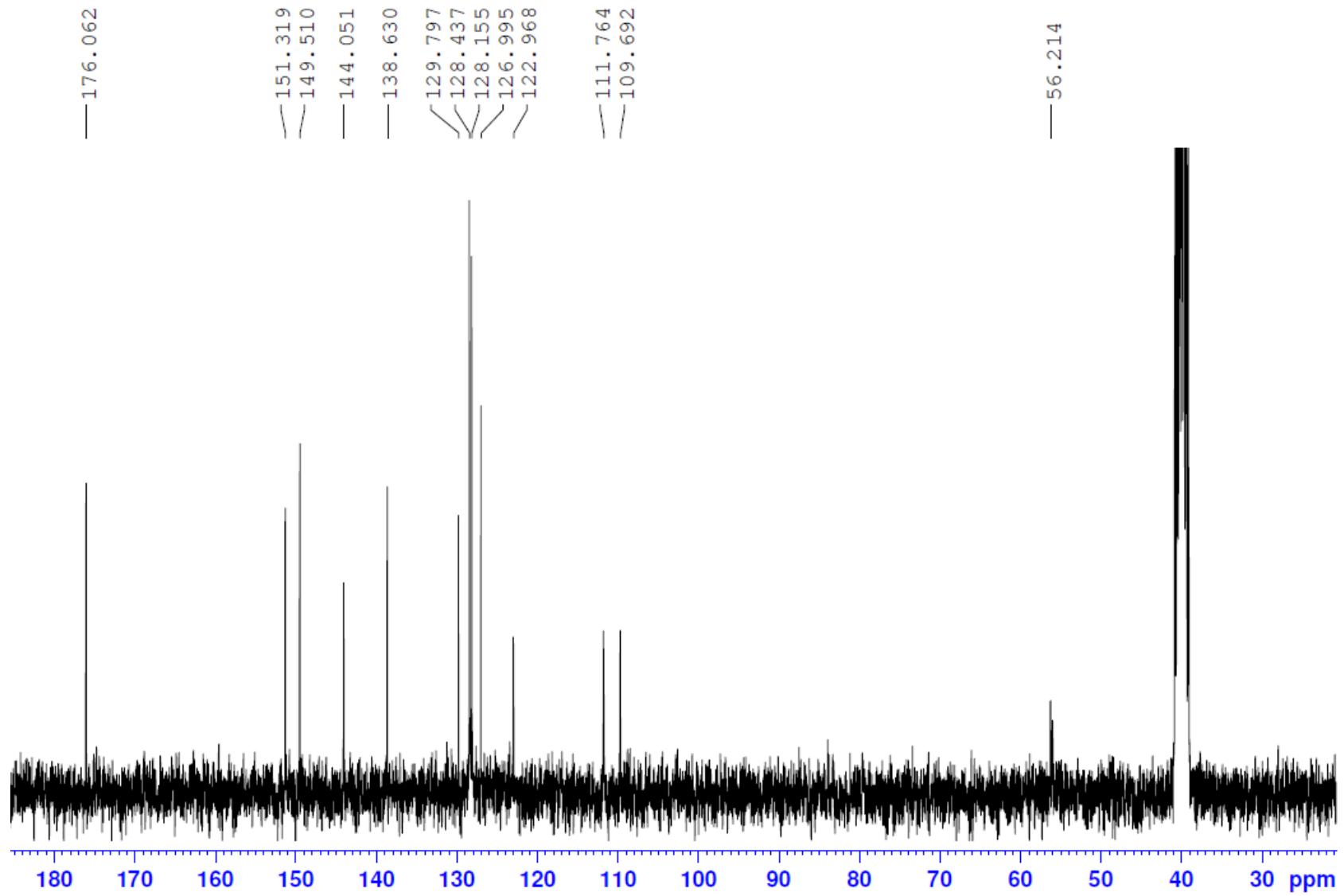


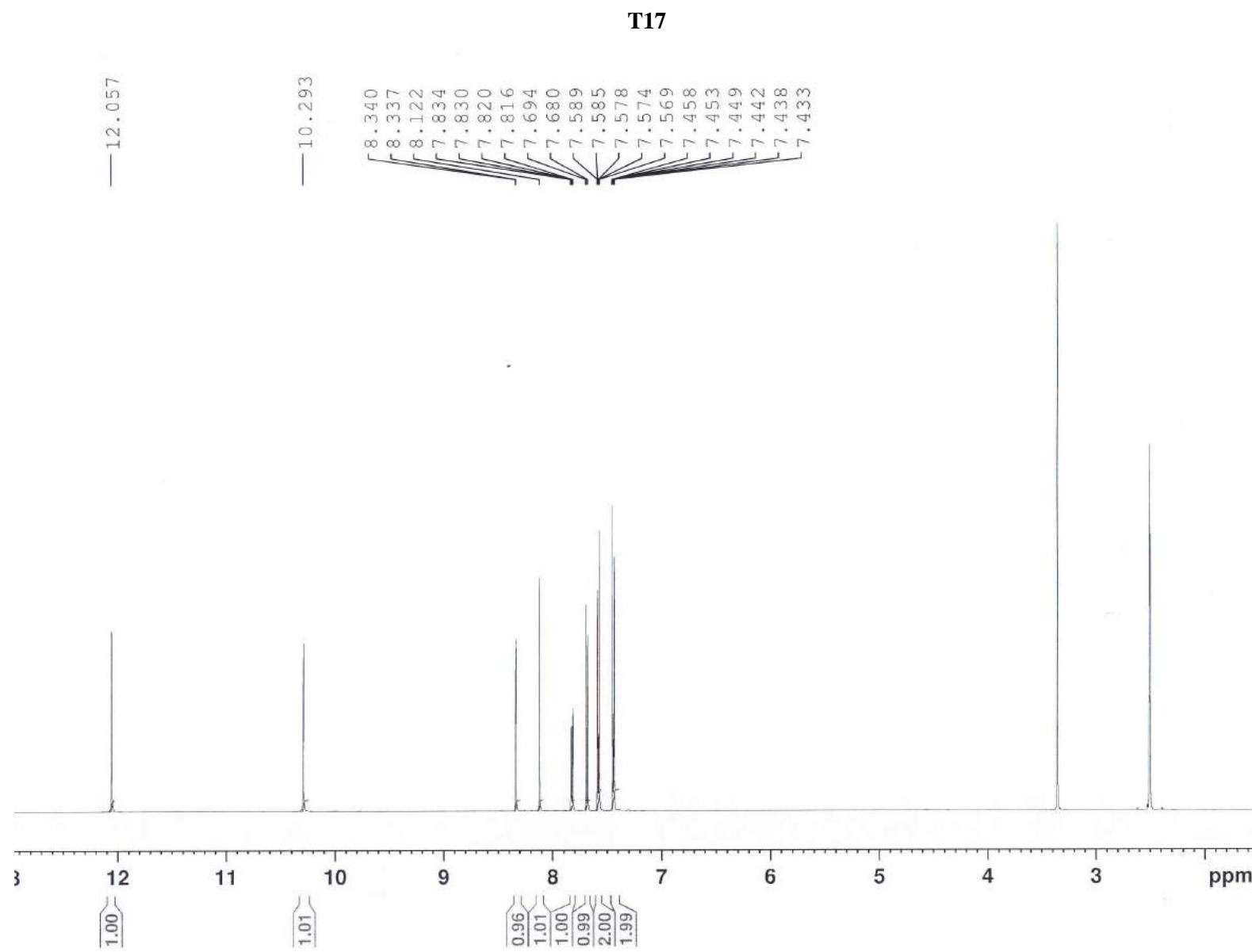
T15

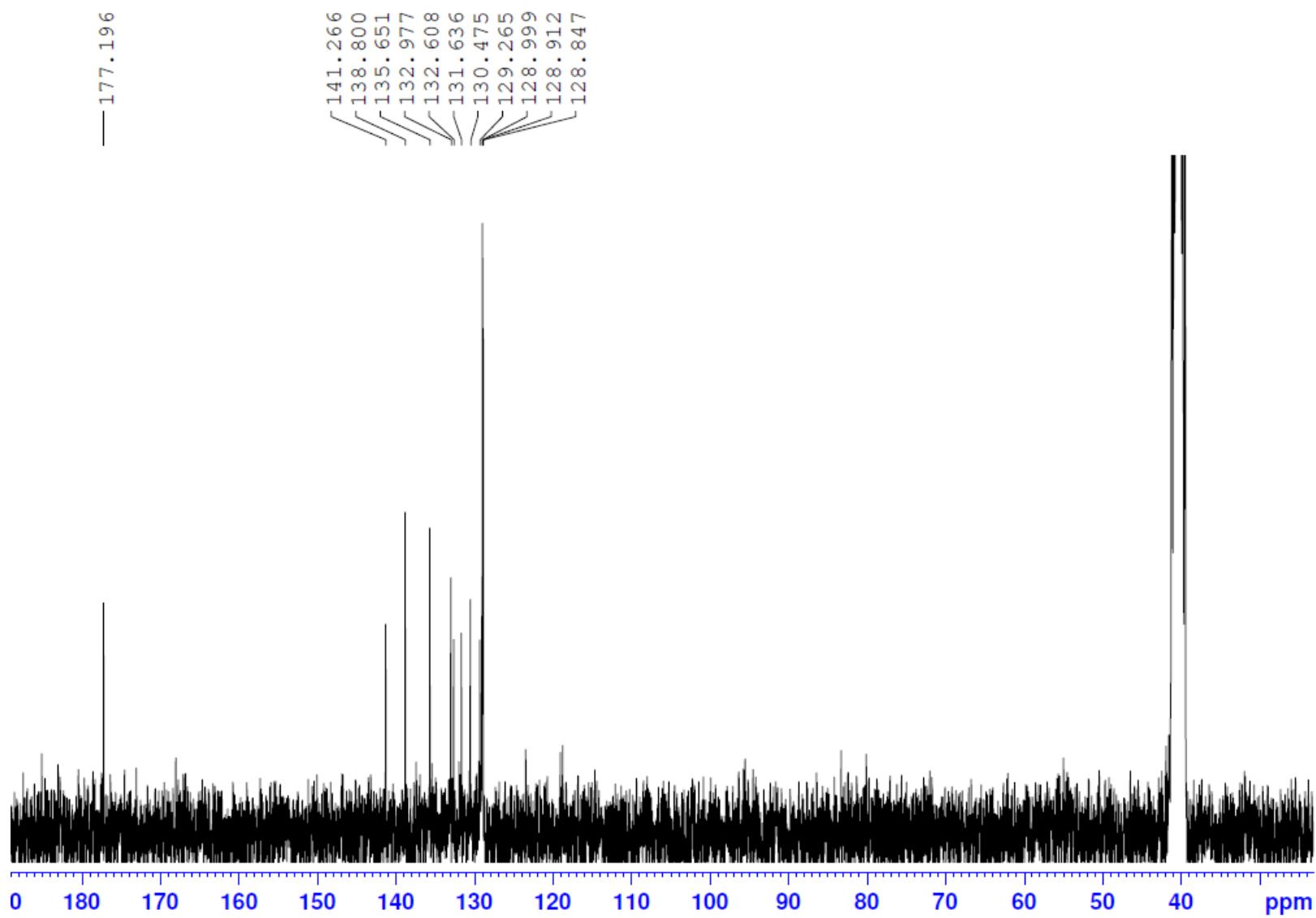


T16

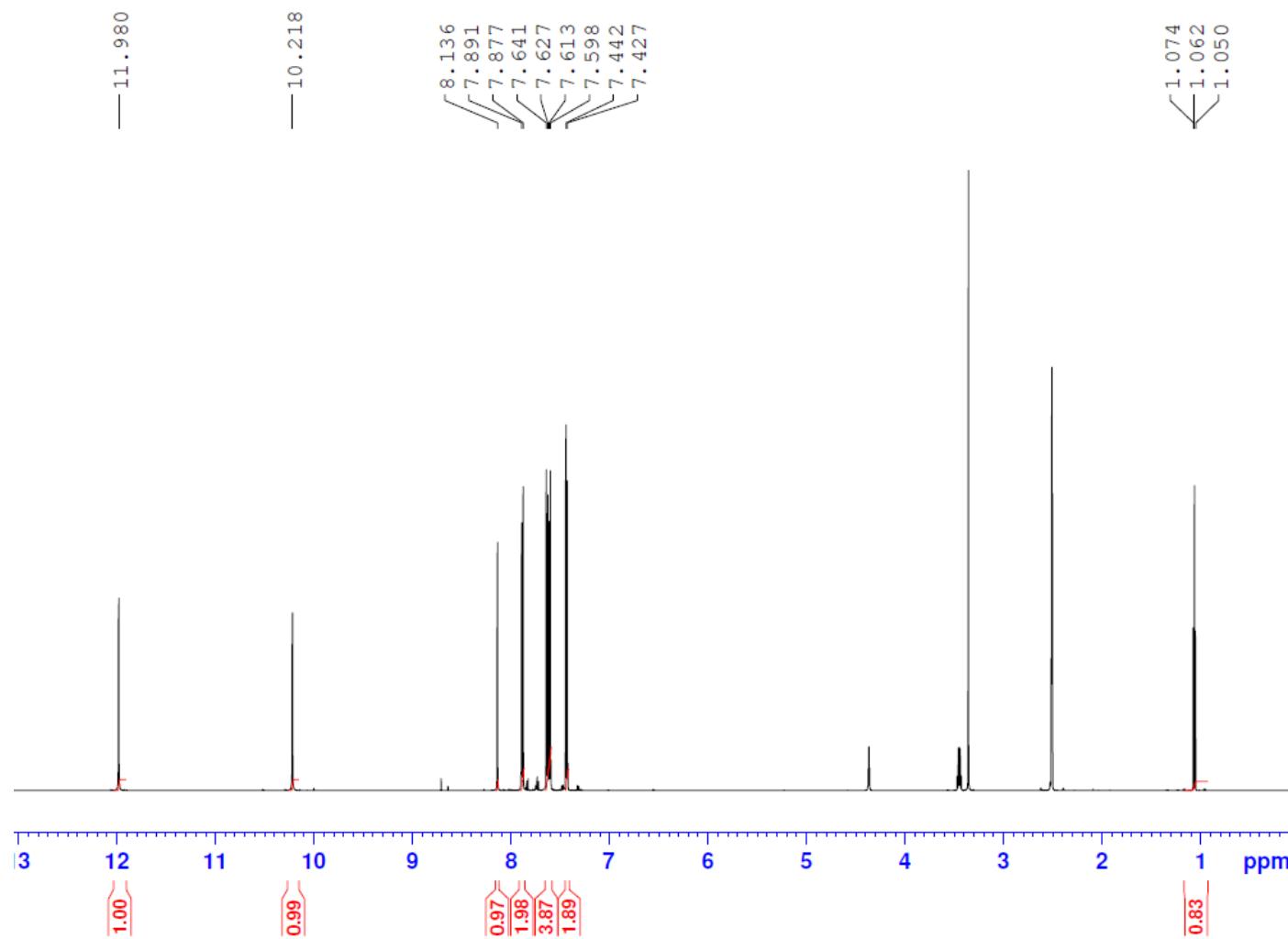


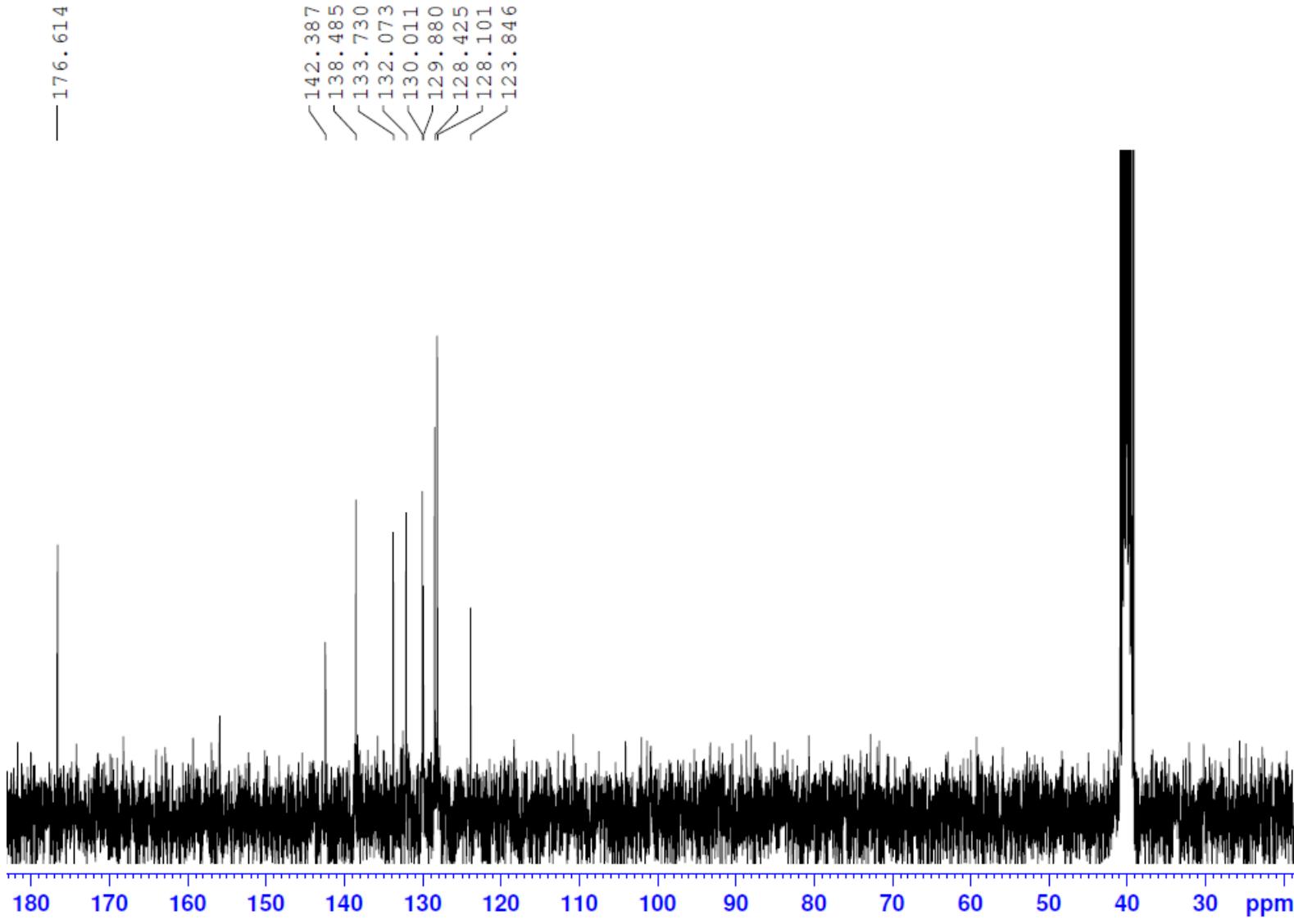






T18





T19

