Supplementary material for publication on line

New application of 1,2,4-triazole derivatives as antitubercular agents. Structure, in vitro screening and docking studies.

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X-ray structure determination

CIF for C1

data_shelx

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_audit_creation_method
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_shelx_SHELXL_version_number
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                              ?
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_chemical_formula_sum
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_chemical_formula_weight
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'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
'H' 'H' 0.0000 0.0000
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'F' 'F' 0.0171 0.0103
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_space_group_name_Hall	'-P 1'

_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.

28.0210

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loop_
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_cell_measurement_theta_max

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_reflns_special_details

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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 'ORTEP3 for Windows (Farrugia, 2012)' _computing_publication_material SHELXL-2014/7 (Sheldrick, 2014) and WINGX (Farrugia, 2012) : ? _refine_special_details _refine_ls_structure_factor_coef Fsqd _refine_ls_matrix_type full _refine_ls_weighting_scheme calc _refine_ls_weighting_details $w=1/[s^2(Fo^2)+(0.0698P)^2+0.1997P]$ 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_coef 2845 _refine_ls_number_reflns _refine_ls_number_parameters 175 _refine_ls_number_restraints 0 _refine_ls_R_factor_all 0.0802 _refine_ls_R_factor_gt 0.0559 _refine_ls_wR_factor_ref 0.1638 _refine_ls_wR_factor_gt 0.1416 _refine_ls_goodness_of_fit_ref 1.080 _refine_ls_restrained_S_all 1.080 _refine_ls_shift/su_max 0.000 _refine_ls_shift/su_mean 0.000

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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 S5 S 1.02937(11) -0.08188(11) 0.83396(6) 0.0529(3) Uani 1 1 d F42 F 0.8709(3) 0.2988(3) 0.56558(16) 0.0712(6) Uani 1 1 d N1 N 0.7815(4) 0.1184(3) 0.9481(2) 0.0489(6) Uani 1 1 d H1 H 0.835(5) 0.101(4) 1.008(3) 0.073 Uiso 1 1 d . U . . . N2 N 0.6229(4) 0.2223(3) 0.9409(2) 0.0514(6) Uani 1 1 d N4 N 0.7062(3) 0.1032(3) 0.79626(18) 0.0397(5) Uani 1 1 d N62 N 0.4161(4) 0.3240(3) 0.6953(2) 0.0544(6) Uani 1 1 d C3 C 0.5785(4) 0.2111(4) 0.8478(2) 0.0427(6) Uani 1 1 d C5 C 0.8395(4) 0.0452(4) 0.8613(2) 0.0417(6) Uani 1 1 d C41 C 0.6988(4) 0.0385(3) 0.7014(2) 0.0391(6) Uani 1 1 d C42 C 0.7849(4) 0.1372(4) 0.5867(2) 0.0430(6) Uani 1 1 d C43 C 0.7865(4) 0.0760(4) 0.4945(3) 0.0548(8) Uani 1 1 d H43 H 0.8438 0.1460 0.4171 0.082 Uiso 1 1 calc R U ... C44 C 0.7012(5) -0.0912(5) 0.5192(3) 0.0616(9) Uani 1 1 d H44 H 0.7010 -0.1353 0.4579 0.092 Uiso 1 1 calc R U . . . C45 C 0.6160(5) -0.1947(4) 0.6338(3) 0.0665(9) Uani 1 1 d H45 H 0.5590 -0.3078 0.6491 0.100 Uiso 1 1 calc R U . . . C46 C 0.6146(5) -0.1310(4) 0.7268(3) 0.0554(7) Uani 1 1 d H46 H 0.5584 -0.2009 0.8044 0.083 Uiso 1 1 calc R U . . . C61 C 0.4153(4) 0.3050(4) 0.8062(2) 0.0448(6) Uani 1 1 d C63 C 0.2672(5) 0.4080(4) 0.6589(3) 0.0639(9) Uani 1 1 d H63 H 0.2657 0.4234 0.5817 0.096 Uiso 1 1 calc R U . . . C64 C 0.1161(5) 0.4733(4) 0.7273(3) 0.0637(9) Uani 1 1 d H64 H 0.0134 0.5279 0.6981 0.096 Uiso 1 1 calc R U . . . C65 C 0.1199(5) 0.4563(5) 0.8396(3) 0.0678(9) Uani 1 1 d H65 H 0.0214 0.5023 0.8876 0.102 Uiso 1 1 calc R U . . . C66 C 0.2721(5) 0.3696(4) 0.8810(3) 0.0605(8) Uani 1 1 d H66 H 0.2776 0.3554 0.9573 0.091 Uiso 1 1 calc R U . . .

loop_

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_atom_site_aniso_U_13

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_atom_site_aniso_U_12
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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_

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C41 C46 1.384(4).? C42 C43 1.371(4) . ? C43 C44 1.372(4) . ? C43 H43 0.9300 . ? C44 C45 1.378(5).? C44 H44 0.9300 . ? C45 C46 1.392(4) . ? C45 H45 0.9300 . ? C46 H46 0.9300 . ? C61 C66 1.378(4).? C63 C64 1.369(5).? C63 H63 0.9300 . ? C64 C65 1.365(5).? C64 H64 0.9300 . ? C65 C66 1.384(5).? C65 H65 0.9300 . ? C66 H66 0.9300 . ?

loop_

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C45 C44 H44 119.6 . . ? C44 C45 C46 120.5(3) . . ? C44 C45 H45 119.8 . . ? C46 C45 H45 119.8 . . ? C41 C46 C45 118.5(3) . . ? C41 C46 H46 120.7 . . ? C45 C46 H46 120.7 . . ? N62 C61 C66 123.4(3) . . ? N62 C61 C3 117.0(2) . . ? C66 C61 C3 119.6(3) . . ? N62 C63 C64 124.4(3) . . ? N62 C63 H63 117.8 . . ? C64 C63 H63 117.8 . . ? C65 C64 C63 118.3(3) . . ? C65 C64 H64 120.9 . . ? C63 C64 H64 120.9 . . ? C64 C65 C66 119.1(3) . . ? C64 C65 H65 120.4 . . ? C66 C65 H65 120.4 . . ? C61 C66 C65 118.2(3) . . ? C61 C66 H66 120.9 . . ? C65 C66 H66 120.9 . . ? 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 C5 N1 N2 C3 1.1(3)? N1 N2 C3 N4 -0.3(3) ? N1 N2 C3 C61 -179.5(2)? C5 N4 C3 N2 -0.6(3)? C41 N4 C3 N2 172.4(2)? C5 N4 C3 C61 178.6(3)? C41 N4 C3 C61 -8.5(4) ? N2 N1 C5 N4 -1.4(3) ? N2 N1 C5 S5 177.3(2)? C3 N4 C5 N1 1.2(3) . . . ? C41 N4 C5 N1 -172.4(2) ? C3 N4 C5 S5 -177.6(2)? C41 N4 C5 S5 8.9(4) ? C5 N4 C41 C42 -99.2(3)? C3 N4 C41 C42 88.8(3)? C5 N4 C41 C46 76.6(3)? C3 N4 C41 C46 -95.4(3)?

C46 C41 C42 F42 -177.9(2)? N4 C41 C42 F42 -2.1(4)? C46 C41 C42 C43 1.8(4)? N4 C41 C42 C43 177.6(2)? F42 C42 C43 C44 178.6(3)? C41 C42 C43 C44 -1.1(4) ? C42 C43 C44 C45 0.1(5)? C43 C44 C45 C46 0.1(5)? C42 C41 C46 C45 -1.5(4) . . . ? N4 C41 C46 C45 -177.4(3)? C44 C45 C46 C41 0.6(5)? C63 N62 C61 C66 -1.0(4)? C63 N62 C61 C3 179.6(3)? N2 C3 C61 N62 162.3(3)? N4 C3 C61 N62 -16.7(4)? N2 C3 C61 C66 -17.2(4) ? N4 C3 C61 C66 163.8(3) ? C61 N62 C63 C64 -0.5(5)? N62 C63 C64 C65 2.0(5)? C63 C64 C65 C66 -1.9(5)? N62 C61 C66 C65 1.0(5)? C3 C61 C66 C65 -179.6(3)? C64 C65 C66 C61 0.5(5)? _refine_diff_density_max 0.697 _refine_diff_density_min -0.255 _refine_diff_density_rms 0.055 _shelx_res_file : shelx.res created by SHELXL-2014/7 TITL t1 in P-1 CELL 0.71073 7.0908 7.8326 12.4147 71.722 74.428 87.846 ZERR 2.00 0.0006 0.0007 0.0010 0.008 0.007 0.007 LATT 1 SFAC C H N F S UNIT 26 18 8 2 2 MERG 2 FMAP 2 PLAN -2 ACTA BOND \$H CONF L.S. 40 WGHT 0.069800 0.199700 FVAR 3.49246 MOLE 1 S5 5 1.029375 -0.081877 0.833960 11.00000 0.05789 0.06767 =

0.04357 -0.02557 -0.02374 0.01970 F42 4 0.870932 0.298845 0.565578 11.00000 0.08746 0.06599 = 0.05521 -0.00812 -0.02138 -0.01763 N1 3 0.781519 0.118388 0.948119 11.00000 0.05813 0.05730 = 0.03990 -0.02147 -0.02150 0.01295 H1 2 0.835365 0.100618 1.007969 11.00000 -1.50000 N2 3 0.622858 0.222322 0.940877 11.00000 0.05927 0.05741 = 0.04583 -0.02388 -0.02017 0.01279 N4 3 0.706198 0.103198 0.796261 11.00000 0.04626 0.04200 = 0.03462 -0.01348 -0.01588 0.00509 N62 3 0.416052 0.323985 0.695349 11.00000 0.06496 0.05571 = 0.05748 -0.02774 -0.03159 0.02165 C3 1 0.578480 0.211067 0.847799 11.00000 0.04933 0.04307 = 0.03878 -0.01606 -0.01376 0.00594 C5 1 0.839460 0.045242 0.861251 11.00000 0.04872 0.04419 = 0.03475 -0.01199 -0.01596 0.00125 C41 1 0.698826 0.038488 0.701409 11.00000 0.04342 0.04316 = 0.03708 -0.01593 -0.01783 0.00519 C42 1 0.784935 0.137151 0.586726 11.00000 0.04326 0.04699 = 0.04447 -0.01733 -0.01788 0.00195 C43 1 0.786498 0.076002 0.494457 11.00000 0.05178 0.07646 = 0.04378 -0.02492 -0.01922 0.00999 AFIX 43 H43 2 0.843831 0.146004 0.417109 11.00000 -1.50000 AFIX 0 C44 1 0.701230 -0.091247 0.519222 11.00000 0.06868 0.07558 = 0.06622 -0.04337 -0.03725 0.01883 AFIX 43 H44 2 0.700961 -0.135316 0.457875 11.00000 -1.50000 AFIX 0 C45 1 0.616026 -0.194675 0.633824 11.00000 0.08043 0.05177 = 0.08609 -0.03173 -0.04134 0.00176 AFIX 43 H45 2 0.559036 -0.307825 0.649135 11.00000 -1.50000 AFIX 0 C46 1 0.614630 -0.131044 0.726837 11.00000 0.06452 0.04868 = 0.05867 -0.01688 -0.02509 -0.00209 AFIX 43 H46 2 0.558361 -0.200904 0.804367 11.00000 -1.50000 AFIX 0 C61 1 0.415255 0.305046 0.806229 11.00000 0.04988 0.03930 = 0.04744 -0.01497 -0.01579 0.00522 C63 1 0.267226 0.408015 0.658871 11.00000 0.07601 0.05880 = 0.07765 -0.03196 -0.04471 0.02090 AFIX 43 H63 2 0.265733 0.423433 0.581719 11.00000 -1.50000 AFIX 0 C64 1 0.116059 0.473342 0.727318 11.00000 0.05924 0.04958 = 0.08681 -0.01773 -0.03308 0.01410 AFIX 43 H64 2 0.013450 0.527918 0.698085 11.00000 -1.50000

AFIX 0 C65 1 0.119857 0.456291 0.839576 11.00000 0.05961 0.05882 = $0.07603 \quad -0.01839 \quad -0.00901 \quad 0.01785$ AFIX 43 H65 2 0.021400 0.502327 0.887567 11.00000 -1.50000 AFIX 0 C66 1 0.272095 0.369626 0.881030 11.00000 0.06530 0.06014 = 0.05142 -0.01555 -0.01259 0.01669 AFIX 43 H66 2 0.277633 0.355350 0.957347 11.00000 -1.50000 AFIX 0 HKLF 4 REM t1 in P-1 REM R1 = 0.0559 for 2092 Fo > 4sig(Fo) and 0.0802 for all 2845 data REM 175 parameters refined using 0 restraints END WGHT 0.0651 0.2555 REM Highest difference peak 0.697, deepest hole -0.255, 1-sigma level 0.055 Q1 1 0.3191 0.3773 0.9534 11.00000 0.05 0.20 Q2 1 0.4976 0.2649 0.8239 11.00000 0.05 0.19 ;

CIF for C12

data_shelx

_audit_creation_method 'SHELXL-2014/7' _shelx_SHELXL_version_number '2014/7' _chemical_name_systematic 4-(4-methoxyphenyl)-3-(3-pyridyl)-1H-1,2,4-triazole-5-thione : ? _chemical_name_common 9 _chemical_melting_point 'C14 H12 N4 O S' _chemical_formula_moiety _chemical_formula_sum 'C14 H12 N4 O S' _chemical_formula_weight 284.34 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	9.3878	(19)
_cell_length_b	9.858(2	2)
_cell_length_c	14.600	(3)
_cell_angle_alpha	90	
_cell_angle_beta	90.59	(3)
_cell_angle_gamma	90	
_cell_volume	1351.1	(5)
_cell_formula_units_	Z 4	
_cell_measurement_t	emperature	296(2)
_cell_measurement_r	eflns_used	2157
_cell_measurement_t	heta_min	2.4820
_cell_measurement_t	heta_max	27.8000

_exptl_crystal_description needle colourles _exptl_crystal_colour _exptl_crystal_density_meas ? ? _exptl_crystal_density_method _exptl_crystal_density_diffrn 1.398 592 _exptl_crystal_F_000 _exptl_transmission_factor_min ? _exptl_transmission_factor_max ? _exptl_crystal_size_max 0.30 _exptl_crystal_size_mid 0.10

_exptl_crystal_size_min 0.10 _exptl_absorpt_coefficient_mu 0.240 _shelx_estimated_absorpt_T_min ? _shelx_estimated_absorpt_T_max ? _exptl_absorpt_correction_T_min 0.91656 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 ? _diffrn_ambient_temperature 296(2) _diffrn_radiation_wavelength 0.71073 _diffrn_radiation_type MoK\a 9 diffrn source _diffrn_measurement_device_type 'KM4 CCD four-circle diffractometer' _diffrn_measurement_method '\w scans' diffrn detector area resol mean ? 5862 _diffrn_reflns_number 0.0442 _diffrn_reflns_av_unetI/netI _diffrn_reflns_av_R_equivalents 0.0307 _diffrn_reflns_limit_h_min -12 12 _diffrn_reflns_limit_h_max -7 _diffrn_reflns_limit_k_min _diffrn_reflns_limit_k_max 12 -8 _diffrn_reflns_limit_l_min 19 _diffrn_reflns_limit_l_max 2.170 _diffrn_reflns_theta_min diffrn reflns theta max 29.234 _diffrn_reflns_theta_full 25.242 _diffrn_measured_fraction_theta_max 0.837 diffrn measured fraction theta full 1.000 _diffrn_reflns_Laue_measured_fraction_max 0.837 _diffrn_reflns_Laue_measured_fraction_full 1.000 _diffrn_reflns_point_group_measured_fraction_max 0.837 _diffrn_reflns_point_group_measured_fraction_full 1.000 reflns number total 3073 _reflns_number_gt 2286 _reflns_threshold_expression I > 2 (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 'ORTEP3 for Windows (Farrugia, 2012)' _computing_publication_material SHELXL-2014/7 (Sheldrick, 2014) and WINGX (Farrugia, 2012) ; refine special details ? _refine_ls_structure_factor_coef Fsqd _refine_ls_matrix_type full _refine_ls_weighting_scheme calc _refine_ls_weighting_details $w=1/[s^2(Fo^2)+(0.0479P)^2+0.4399P]$ 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_coef 3073 _refine_ls_number_reflns _refine_ls_number_parameters 184 refine ls number restraints 0 _refine_ls_R_factor_all 0.0661 _refine_ls_R_factor_gt 0.0423 _refine_ls_wR_factor_ref 0.1259

0.0958 _refine_ls_wR_factor_gt _refine_ls_goodness_of_fit_ref 1.113 _refine_ls_restrained_S_all 1.113 _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 S5 S 0.24115(7) 0.56484(6) 0.16270(4) 0.04008(19) Uani 1 1 d O47 O -0.0266(2) 0.99272(19) -0.14262(14) 0.0608(5) Uani 1 1 d N1 N 0.3579(2) 0.37414(18) 0.05440(12) 0.0333(4) Uani 1 1 d H1 H 0.369(3) 0.310(3) 0.0997(17) 0.050 Uiso 1 1 d . U . . . N2 N 0.3863(2) 0.34250(17) -0.03527(12) 0.0339(4) Uani 1 1 d N4 N 0.27969(18) 0.54222(16) -0.02347(11) 0.0281(4) Uani 1 1 d N63 N 0.3873(2) 0.34019(18) -0.32216(12) 0.0375(4) Uani 1 1 d C3 C 0.3377(2) 0.44692(19) -0.08155(13) 0.0280(4) Uani 1 1 d C5 C 0.2930(2) 0.4941(2) 0.06514(14) 0.0293(4) Uani 1 1 d C41 C 0.2016(2) 0.6621(2) -0.05053(13) 0.0291(4) Uani 1 1 d C42 C 0.2628(2) 0.7883(2) -0.04163(14) 0.0345(5) Uani 1 1 d H42 H 0.3530 0.7971 -0.0153 0.052 Uiso 1 1 calc R U . . . C43 C 0.1897(3) 0.9026(2) -0.07196(16) 0.0405(5) Uani 1 1 d H43 H 0.2306 0.9881 -0.0660 0.061 Uiso 1 1 calc R U . . . C44 C 0.0556(3) 0.8883(2) -0.11120(16) 0.0408(6) Uani 1 1 d C45 C -0.0042(3) 0.7602(3) -0.11995(17) 0.0459(6) Uani 1 1 d H45 H -0.0940 0.7507 -0.1466 0.069 Uiso 1 1 calc R U . . . C46 C 0.0678(2) 0.6474(2) -0.08960(16) 0.0397(5) Uani 1 1 d H46 H 0.0269 0.5618 -0.0953 0.060 Uiso 1 1 calc R U . . . C48 C 0.0307(3) 1.1257(3) -0.1398(2) 0.0671(9) Uani 1 1 d H481 H 0.1109 1.1311 -0.1799 0.101 Uiso 1 1 calc R U ... H482 H 0.0607 1.1466 -0.0783 0.101 Uiso 1 1 calc R U . . . H483 H -0.0408 1.1894 -0.1592 0.101 Uiso 1 1 calc R U ... C61 C 0.3524(2) 0.45924(19) -0.18097(13) 0.0280(4) Uani 1 1 d C62 C 0.3647(2) 0.3412(2) -0.23250(14) 0.0338(5) Uani 1 1 d H62 H 0.3565 0.2584 -0.2025 0.051 Uiso 1 1 calc R U . . . C64 C 0.3957(3) 0.4603(2) -0.36444(15) 0.0404(6) Uani 1 1 d H64 H 0.4110 0.4612 -0.4273 0.061 Uiso 1 1 calc R U . . . C65 C 0.3828(3) 0.5820(2) -0.32007(15) 0.0415(6) Uani 1 1 d

H65 H 0.3886 0.6632 -0.3523 0.062 Uiso 1 1 calc R U . . . C66 C 0.3611(3) 0.5820(2) -0.22687(15) 0.0361(5) Uani 1 1 d H66 H 0.3523 0.6634 -0.1952 0.054 Uiso 1 1 calc R U . . .

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 S5 0.0534(4) 0.0410(3) 0.0260(3) -0.0038(2) 0.0077(2) -0.0020(3) O47 0.0589(12) 0.0503(11) 0.0732(14) 0.0149(10) -0.0022(10) 0.0226(10) N1 0.0471(11) 0.0299(9) 0.0229(9) 0.0053(7) 0.0032(8) 0.0020(8) N2 0.0459(11) 0.0290(9) 0.0269(9) 0.0028(7) 0.0043(8) 0.0035(8) N4 0.0357(9) 0.0265(9) 0.0221(8) 0.0001(7) 0.0016(7) 0.0021(7) N63 0.0511(12) 0.0339(10) 0.0276(9) -0.0050(8) 0.0004(8) 0.0008(9) C3 0.0341(11) 0.0251(10) 0.0248(10) -0.0010(8) 0.0016(8) -0.0004(8) C5 0.0333(11) 0.0280(10) 0.0265(10) 0.0022(8) 0.0013(8) -0.0039(9) C41 0.0356(11) 0.0287(10) 0.0230(9) 0.0000(8) 0.0027(8) 0.0034(9) C42 0.0390(12) 0.0321(11) 0.0325(11) -0.0023(9) -0.0036(9) 0.0001(10) C43 0.0506(14) 0.0302(12) 0.0409(13) -0.0005(10) 0.0001(11) 0.0032(10) C44 0.0442(13) 0.0419(13) 0.0364(12) 0.0049(10) 0.0043(10) 0.0153(11) C45 0.0349(12) 0.0517(15) 0.0509(15) 0.0034(12) -0.0058(11) 0.0044(11) C46 0.0358(12) 0.0388(12) 0.0445(13) -0.0012(10) -0.0012(10) -0.0034(10) C48 0.086(2) 0.0478(17) 0.068(2) 0.0187(14) 0.0159(16) 0.0275(16) C61 0.0335(11) 0.0274(10) 0.0231(10) -0.0007(8) 0.0006(8) -0.0003(9) C62 0.0441(12) 0.0280(11) 0.0293(11) 0.0001(9) 0.0014(9) -0.0001(10) C64 0.0555(15) 0.0424(13) 0.0233(10) -0.0025(9) 0.0023(10) 0.0020(11) C65 0.0632(16) 0.0321(12) 0.0293(11) 0.0037(9) 0.0036(10) 0.0013(11) C66 0.0544(14) 0.0268(11) 0.0273(11) -0.0018(8) 0.0021(10) 0.0008(10)

_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 S5 C5 1.663(2) . ? O47 C44 1.363(3) . ? O47 C48 1.417(4) . ? N1 C5 1.340(3).? N1 N2 1.375(2) . ? N1 H1 0.92(3) . ? N2 C3 1.311(2) . ? N4 C3 1.381(2).? N4 C5 1.383(2) . ? N4 C41 1.443(2).? N63 C62 1.328(3).? N63 C64 1.338(3) . ? C3 C61 1.465(3) . ? C41 C42 1.376(3) . ? C41 C46 1.382(3) . ? C42 C43 1.389(3) . ? C42 H42 0.9300 . ? C43 C44 1.385(3).? C43 H43 0.9300 . ? C44 C45 1.387(3).? C45 C46 1.373(3) . ? C45 H45 0.9300 . ? C46 H46 0.9300 . ? C48 H481 0.9600 . ? C48 H482 0.9600 .? C48 H483 0.9600 . ? C61 C66 1.386(3) . ? C61 C62 1.391(3) . ? C62 H62 0.9300 . ? C64 C65 1.370(3) . ? C64 H64 0.9300 . ? C65 C66 1.378(3).? C65 H65 0.9300 . ? C66 H66 0.9300 . ? 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 C44 O47 C48 118.4(2) . . ? C5 N1 N2 113.87(17)..? C5 N1 H1 125.0(16) . . ? N2 N1 H1 120.3(16) . . ? C3 N2 N1 104.05(16) . . ? C3 N4 C5 107.99(16) . . ? C3 N4 C41 126.23(16) . . ? C5 N4 C41 125.29(16) . . ? C62 N63 C64 117.30(18) . . ? N2 C3 N4 110.79(17) . . ?

N2 C3 C61 122.69(18) . . ? N4 C3 C61 126.44(17) . . ? N1 C5 N4 103.30(17) . . ? N1 C5 S5 127.42(16) . . ? N4 C5 S5 129.29(16) . . ? C42 C41 C46 120.7(2) . . ? C42 C41 N4 120.29(18) . . ? C46 C41 N4 118.93(19) . . ? C41 C42 C43 119.9(2) . . ? C41 C42 H42 120.0 . . ? C43 C42 H42 120.0 . . ? C44 C43 C42 119.6(2) . . ? C44 C43 H43 120.2 . . ? C42 C43 H43 120.2 . . ? O47 C44 C43 124.9(2) . . ? O47 C44 C45 115.4(2) . . ? C43 C44 C45 119.7(2) . . ? C46 C45 C44 120.7(2) . . ? C46 C45 H45 119.6 . . ? C44 C45 H45 119.6 . . ? C45 C46 C41 119.4(2) . . ? C45 C46 H46 120.3 . . ? C41 C46 H46 120.3 . . ? O47 C48 H481 109.5 . . ? O47 C48 H482 109.5 . . ? H481 C48 H482 109.5 . . ? O47 C48 H483 109.5 . . ? H481 C48 H483 109.5 . . ? H482 C48 H483 109.5 . . ? C66 C61 C62 117.59(19) . . ? C66 C61 C3 123.93(18) . . ? C62 C61 C3 118.41(18) . . ? N63 C62 C61 123.65(19) . . ? N63 C62 H62 118.2 . . ? C61 C62 H62 118.2 . . ? N63 C64 C65 123.5(2) . . ? N63 C64 H64 118.3 . . ? C65 C64 H64 118.3 . . ? C64 C65 C66 118.8(2) . . ? C64 C65 H65 120.6 . . ? C66 C65 H65 120.6 . . ? C65 C66 C61 119.19(19) . . ? C65 C66 H66 120.4 . . ? C61 C66 H66 120.4 . . ?

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 C5 N1 N2 C3 0.3(2)? N1 N2 C3 N4 $-0.1(2) \dots ?$ N1 N2 C3 C61 176.84(18)? C5 N4 C3 N2 0.0(2)? C41 N4 C3 N2 -172.29(18)? C5 N4 C3 C61 -176.85(19)? C41 N4 C3 C61 10.9(3) ? N2 N1 C5 N4 -0.3(2) ? N2 N1 C5 S5 179.40(16)? C3 N4 C5 N1 0.2(2)? C41 N4 C5 N1 172.53(18)? C3 N4 C5 S5 -179.49(16)? C41 N4 C5 S5 -7.1(3) . . . ? C3 N4 C41 C42 -107.4(2)? C5 N4 C41 C42 81.6(3)? C3 N4 C41 C46 69.4(3)? C5 N4 C41 C46 -101.6(2)? C46 C41 C42 C43 0.1(3)? N4 C41 C42 C43 176.85(19)? C41 C42 C43 C44 -0.1(3)? C48 O47 C44 C43 3.4(4) ? C48 O47 C44 C45 -177.0(2)? C42 C43 C44 O47 179.4(2)? C42 C43 C44 C45 -0.1(4)? O47 C44 C45 C46 -179.2(2) ? C43 C44 C45 C46 0.4(4) ? C44 C45 C46 C41 -0.4(4) ? C42 C41 C46 C45 0.2(3)? N4 C41 C46 C45 -176.6(2) ? N2 C3 C61 C66 -151.1(2)? N4 C3 C61 C66 25.4(3) ? N2 C3 C61 C62 26.0(3)? N4 C3 C61 C62 -157.5(2) . . . ? C64 N63 C62 C61 -1.4(3)? C66 C61 C62 N63 1.6(3) . . . ? C3 C61 C62 N63 -175.7(2)? C62 N63 C64 C65 0.3(4)? N63 C64 C65 C66 0.5(4) . . . ? C64 C65 C66 C61 -0.2(4) . . . ? C62 C61 C66 C65 -0.8(3) ? C3 C61 C66 C65 176.4(2) ?

_refine_diff_density_max 0.206 _refine_diff_density_min -0.252 _refine_diff_density_rms 0.055

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_shelx_res_file
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:
  shelx.res created by SHELXL-2014/7
TITL T15 in P2(1)/c
CELL 0.71073 9.3878 9.8576 14.6003 90.000 90.586 90.000
ZERR 4.00 0.0019 0.0020 0.0029 0.000 0.030 0.000
LATT 1
SYMM - X, 1/2 + Y, 1/2 - Z
SFAC C H N O S
UNIT 56 48 16 4 4
MERG 2
FMAP 2
PLAN -2
ACTA
BOND $H
CONF
L.S. 40
WGHT 0.047900 0.439900
FVAR
       1.59215
MOLE 1
S5 5 0.241152 0.564845 0.162700 11.00000 0.05339 0.04096 =
    0.02604 -0.00377 0.00773 -0.00204
O47 4 -0.026586 0.992715 -0.142623 11.00000 0.05889 0.05028 =
    0.07318 0.01492 -0.00225 0.02259
N1 3 0.357904 0.374142 0.054397 11.00000 0.04710 0.02993 =
    0.02291 0.00527 0.00323 0.00204
H1 2 0.369350 0.309782 0.099680 11.00000 -1.50000
N2 3 0.386303 0.342496 -0.035267 11.00000 0.04595 0.02904 =
    0.02690 0.00284 0.00432 0.00353
N4 3 0.279690 0.542218 -0.023475 11.00000 0.03566 0.02649 =
    0.02208 0.00014 0.00163 0.00210
N63 3 0.387341 0.340187 -0.322163 11.00000 0.05106 0.03387 =
    0.02761 -0.00500 0.00038 0.00076
C3 1 0.337663 0.446921 -0.081553 11.00000 0.03413 0.02508 =
    0.02476 -0.00100 0.00160 -0.00037
C5 1 0.292966 0.494086 0.065143 11.00000 0.03331 0.02804 =
    0.02653 0.00218 0.00134 -0.00394
C41 1 0.201632 0.662100 -0.050534 11.00000 0.03558 0.02868 =
    0.02301 0.00005 0.00271 0.00335
C42 1 0.262786 0.788329 -0.041629 11.00000 0.03895 0.03207 =
    0.03247 -0.00233 -0.00359 0.00007
AFIX 43
H42 2 0.352954 0.797135 -0.015326 11.00000 -1.50000
AFIX 0
C43 1 0.189724 0.902553 -0.071958 11.00000 0.05058 0.03017 =
    0.04085 -0.00051 0.00005 0.00319
AFIX 43
H43 2 0.230592 0.988077 -0.065969 11.00000 -1.50000
AFIX 0
```

C44 1 0.055558 0.888310 -0.111198 11.00000 0.04418 0.04186 = 0.03640 0.00489 0.00432 0.01531 C45 1 -0.004195 0.760236 -0.119947 11.00000 0.03493 0.05173 = 0.05091 0.00339 -0.00576 0.00438 AFIX 43 H45 2 -0.094001 0.750701 -0.146650 11.00000 -1.50000 AFIX 0 C46 1 0.067824 0.647373 -0.089604 11.00000 0.03575 0.03879 = 0.04446 -0.00124 -0.00119 -0.00336 AFIX 43 H46 2 0.026942 0.561813 -0.095292 11.00000 -1.50000 AFIX 0 C48 1 0.030683 1.125674 -0.139791 11.00000 0.08629 0.04775 = 0.06765 0.01866 0.01585 0.02747 **AFIX 133** H481 2 0.110898 1.131086 -0.179885 11.00000 -1.50000 H482 2 0.060705 1.146565 -0.078342 11.00000 -1.50000 H483 2 -0.040756 1.189446 -0.159228 11.00000 -1.50000 AFIX 0 C61 1 0.352409 0.459245 -0.180969 11.00000 0.03346 0.02739 = 0.02313 -0.00070 0.00062 -0.00035 C62 1 0.364652 0.341192 -0.232504 11.00000 0.04409 0.02796 = 0.02932 0.00007 0.00141 -0.00009 AFIX 43 H62 2 0.356498 0.258362 -0.202517 11.00000 -1.50000 AFIX 0 C64 1 0.395672 0.460308 -0.364436 11.00000 0.05551 0.04241 = 0.02329 -0.00254 0.00229 0.00200 AFIX 43 H64 2 0.411014 0.461188 -0.427261 11.00000 -1.50000 AFIX 0 C65 1 0.382798 0.582040 -0.320073 11.00000 0.06324 0.03215 = 0.02927 0.00372 0.00358 0.00126 AFIX 43 H65 2 0.388619 0.663233 -0.352256 11.00000 -1.50000 AFIX 0 C66 1 0.361063 0.582008 -0.226874 11.00000 0.05436 0.02684 = 0.02725 -0.00178 0.00210 0.00077 AFIX 43 H66 2 0.352308 0.663354 -0.195163 11.00000 -1.50000 AFIX 0 HKLF 4 REM T15 in P2(1)/cREM R1 = 0.0423 for 2286 Fo > 4sig(Fo) and 0.0661 for all 3073 data REM 184 parameters refined using 0 restraints END WGHT 0.0547 0.2965

```
REM Highest difference peak 0.206, deepest hole -0.252, 1-sigma level 0.055
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Q2 1 0.1945 0.9996 -0.0164 11.00000 0.05 0.19
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CIF for C13

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_shelx_space_group_comment

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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.

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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.
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loop_

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_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_

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C45 H45 0.9300 . ? C46 H46 0.9300 . ? C61 C66 1.370(4) . ? C61 C62 1.380(4).? C62 H62 0.9300 . ? C64 C65 1.374(5).? C64 H64 0.9300 . ? C65 C66 1.377(5).? C65 H65 0.9300 . ? C66 H66 0.9300 . ? 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 C5 N1 N2 113.6(2) . . ? C5 N1 H1 124.0(17) . . ? N2 N1 H1 122.4(17) . . ? C3 N2 N1 104.8(2) . . ? C5 N4 C3 108.5(2) . . ? C5 N4 C41 126.1(2) . . ? C3 N4 C41 125.4(2) . . ? C64 N63 C62 116.8(3) . . ? N2 C3 N4 110.1(2) . . ? N2 C3 C31 125.0(2) . . ? N4 C3 C31 124.8(2) . . ? N1 C5 N4 103.0(2) . . ? N1 C5 S5 129.1(2) . . ? N4 C5 S5 127.8(2) . . ? C3 C31 C61 114.8(2) . . ? C3 C31 H311 108.6 . . ? C61 C31 H311 108.6 . . ? C3 C31 H312 108.6 . . ? C61 C31 H312 108.6 . . ? H311 C31 H312 107.5 . . ? C46 C41 C42 119.8(2) . . ? C46 C41 N4 119.5(2) . . ? C42 C41 N4 120.7(2) . . ? C43 C42 C41 120.1(3) . . ? C43 C42 Cl42 119.3(2) . . ? C41 C42 Cl42 120.7(2) . . ? C44 C43 C42 119.1(3) . . ? C44 C43 H43 120.4 . . ? C42 C43 H43 120.4 . . ? C43 C44 C45 121.6(3) . . ? C43 C44 Cl44 119.5(2) . . ? C45 C44 Cl44 118.8(3) . . ?

C44 C45 C46 118.6(3) . . ? C44 C45 H45 120.7 . . ? C46 C45 H45 120.7 . . ? C41 C46 C45 120.8(3) . . ? C41 C46 H46 119.6 . . ? C45 C46 H46 119.6 . . ? C66 C61 C62 117.0(3) . . ? C66 C61 C31 122.9(3) . . ? C62 C61 C31 120.0(3) . . ? N63 C62 C61 124.3(3) . . ? N63 C62 H62 117.8 . . ? C61 C62 H62 117.8 . . ? N63 C64 C65 123.9(4) . . ? N63 C64 H64 118.0 . . ? C65 C64 H64 118.0 . . ? C64 C65 C66 117.6(4) . . ? C64 C65 H65 121.2 . . ? C66 C65 H65 121.2 . . ? C61 C66 C65 120.3(3) . . ? C61 C66 H66 119.8 . . ? C65 C66 H66 119.8 . . ? 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 C5 N1 N2 C3 0.7(3)? N1 N2 C3 N4 0.6(3) ? N1 N2 C3 C31 -175.9(3) ? C5 N4 C3 N2 -1.6(3) ? C41 N4 C3 N2 179.0(2)? C5 N4 C3 C31 174.9(3)? C41 N4 C3 C31 -4.5(4) ? N2 N1 C5 N4 -1.6(3) ? N2 N1 C5 S5 175.3(2)? C3 N4 C5 N1 1.9(3)? C41 N4 C5 N1 -178.7(2)? C3 N4 C5 S5 -175.0(2) ? C41 N4 C5 S5 4.4(4) ? N2 C3 C31 C61 -118.1(3)? N4 C3 C31 C61 65.9(4) ? C5 N4 C41 C46 -108.0(3)? C3 N4 C41 C46 71.3(4) ? C5 N4 C41 C42 73.5(4)?

C3 N4 C41 C42 -107.2(3) ? C46 C41 C42 C43 0.8(4)? N4 C41 C42 C43 179.4(2)? C46 C41 C42 Cl42 -178.1(2)? N4 C41 C42 Cl42 0.5(4) ? C41 C42 C43 C44 -0.7(4) ? Cl42 C42 C43 C44 178.2(2)? C42 C43 C44 C45 -0.1(5) ? C42 C43 C44 Cl44 178.0(2)? C43 C44 C45 C46 0.8(5) ? Cl44 C44 C45 C46 -177.4(2) . . . ? C42 C41 C46 C45 -0.1(5)? N4 C41 C46 C45 -178.7(3)? C44 C45 C46 C41 -0.7(5) ? C3 C31 C61 C66 44.0(4) ? C3 C31 C61 C62 -138.6(3)? C64 N63 C62 C61 0.1(5)? C66 C61 C62 N63 0.5(5)? C31 C61 C62 N63 -177.0(3)? C62 N63 C64 C65 -0.7(5) ? N63 C64 C65 C66 0.6(6) ? C62 C61 C66 C65 -0.6(4) ? C31 C61 C66 C65 176.8(3)? C64 C65 C66 C61 0.1(5)? _refine_diff_density_max 0.263 _refine_diff_density_min -0.250 _refine_diff_density_rms 0.072 _shelx_res_file ; shelx.res created by SHELXL-2014/7 TITL T17 in P-1 CELL 0.71073 8.4234 8.8584 11.2513 81.356 83.534 66.032 ZERR 2.00 0.0010 0.0009 0.0020 0.013 0.012 0.011 LATT 1 SFAC C H N S CL UNIT 28 20 8 2 4 MERG 0 FMAP 2 ACTA BOND \$H CONF L.S. 40 PLAN -2 WGHT 0.058200 EXTI 0.018621 BASF 0.43793

FVAR 5.87660 CL42 5 0.008987 0.277750 0.114641 11.00000 0.06378 0.04856 = 0.04694 -0.00870 -0.01592 -0.01844 CL44 5 0.260172 -0.211735 0.474562 11.00000 0.08883 0.03452 = 0.07105 0.00877 -0.00268 -0.01920 S5 4 -0.292806 0.592551 0.309938 11.00000 0.03376 0.06467 = 0.06436 -0.01031 -0.00279 -0.02114 N1 3 -0.081959 0.748820 0.214471 11.00000 0.03070 0.02521 = 0.06097 -0.00794 -0.00583 -0.00290 H1 2 -0.177457 0.858046 0.192663 11.00000 -1.50000 N2 3 0.091303 0.724038 0.196216 11.00000 0.03547 0.02612 = 0.06631 -0.00853 -0.00173 -0.01063 N4 3 0.060391 0.494564 0.275511 11.00000 0.02801 0.02387 = 0.04315 -0.00519 -0.00611 -0.00974 N63 3 0.676106 0.072723 0.145731 11.00000 0.04444 0.03081 = 0.08333 -0.00553 -0.00574 -0.00034 C3 1 0.175794 0.568754 0.234591 11.00000 0.03129 0.02702 = 0.04883 -0.00972 -0.00419 -0.01168 C5 1 -0.107077 0.613449 0.264356 11.00000 0.03242 0.03237 = 0.04381 -0.01091 -0.00373 -0.01014 C31 1 0.367567 0.485272 0.240914 11.00000 0.03272 0.03089 = 0.06995 -0.00993 -0.00717 -0.01192 AFIX 23 H311 2 0.394293 0.442206 0.324017 11.00000 -1.50000 H312 2 0.416930 0.567624 0.216549 11.00000 -1.50000 AFIX 0 C41 1 0.108110 0.322973 0.322657 11.00000 0.03510 0.02879 = 0.03901 -0.00571 -0.00211 -0.01468 C42 1 0.089052 0.212459 0.256222 11.00000 0.03493 0.03303 = 0.04039 -0.00680 -0.00280 -0.01281 C43 1 0.137148 0.047155 0.302315 11.00000 0.05148 0.03697 = 0.04935 -0.01269 0.00288 -0.02187 AFIX 43 H43 2 0.123435 -0.027359 0.258482 11.00000 -1.50000 AFIX 0 C44 1 0.205359 -0.005386 0.413579 11.00000 0.05303 0.02966 = 0.05064 0.00252 -0.00028 -0.01312 C45 1 0.227310 0.102517 0.480386 11.00000 0.06702 0.04690 = 0.04213 0.00245 -0.01296 -0.01573 AFIX 43 H45 2 0.274991 0.064972 0.555250 11.00000 -1.50000 AFIX 0 C46 1 0.176887 0.267666 0.433626 11.00000 0.05409 0.03855 = 0.04473 -0.00801 -0.01087 -0.01826 AFIX 43 H46 2 0.189740 0.342179 0.477922 11.00000 -1.50000 AFIX 0 C61 1 0.454652 0.344724 0.164084 11.00000 0.02405 0.02670 = 0.05332 -0.00005 -0.00230 -0.00980 C62 1 0.586891 0.200032 0.208447 11.00000 0.03981 0.03386 =

0.06124 -0.00142 -0.00672 -0.00387 AFIX 43 H62 2 0.615414 0.191068 0.287419 11.00000 -1.50000 AFIX 0 C64 1 0.633451 0.088206 0.033415 11.00000 0.06068 0.04568 = 0.07234 -0.02078 0.01103 -0.02037 AFIX 43 H64 2 0.695308 0.000801 -0.012128 11.00000 -1.50000 AFIX 0 C65 1 0.503254 0.225714 -0.019816 11.00000 0.06577 0.04936 = 0.05778 -0.00474 -0.00235 -0.02559 AFIX 43 H65 2 0.476729 0.231186 -0.098796 11.00000 -1.50000 AFIX 0 C66 1 0.413493 0.355082 0.047964 11.00000 0.04295 0.03096 = 0.06434 0.00706 -0.01250 -0.01212 AFIX 43 H66 2 0.324488 0.449954 0.014741 11.00000 -1.50000 AFIX 0 HKLF 5 REM T17 in P-1 REM R1 = 0.0397 for 3212 Fo > 4sig(Fo) and 0.0666 for all 4996 data REM 195 parameters refined using 0 restraints END WGHT 0.0582 0.0000 REM Highest difference peak 0.263, deepest hole -0.250, 1-sigma level 0.072 Q1 1 -0.4070 0.5775 0.2774 11.00000 0.05 0.26 $Q2 \quad 1 \quad 0.5058 \quad 0.1623 \quad 0.0052 \quad 11.00000 \quad 0.05 \quad 0.23$;





Figure 1S. The molecular packing of C1, C12 and C13. Dashed lines indicate intermolecular hydrogen bonds; C1 i = -x, 2-y, -z; C12: i = x, 3/2-y, -1/2+z; C13: i = -1+x, 1+y, z.

MS analysis for new compounds

Table 1S.

LP.	Compound formula, thoretical and measured monoisotopic	Theoretical monoisotopic mass of protonated ion and acquired LC/QTOF HRMS spectra		
	mass, and mass difference in			
D12				
D12	C14F12N4OSCI2 Calculated monoisotonic mass:	x10 ⁵ Cpd 1: C14 H12 Cl2 N4 O S: +ESI Scan (rt: 1.617-1.667, 1.851-2.150 min, 23 scans) Frag=350.0V B13M.d S.		
		3-		
	354.0109	2.8-355.0196		
	Measured monoisotonic mass:	2.4-		
	254 0122	2-		
	554.0122	1.8-357.0167		
	Mass error: 3.84 ppm	14-		
		0.8		
		0.4-358.0191 359.0136		
		Counts vs. Mass-to-Charge (m/z)		
B14	C14H12ON4SCI2	x10 ^s Cpd 1: C14 H12 Cl2 N4 O S: +ESI Scan (rt: 2.049-2.066, 2.199-2.449 min, 18 scans) Frag=350.0V B14M.d 1- 355.0193 357.0165		
	Calculated monoisotopic mass:	0.8-		
	354.0109	0.6-		
	Massured manaisatonic mass:	0.4 - 356.0219 359.0135		
		0.2- 360.0159 262.0115		
	554.012	0 352 353 354 355 356 357 358 359 360 361 362 363 364 365 366		
	Mass error: 3.13 ppm	Counts vs. Mass-to-Charge (m/z)		
C13	C14H10N4SCl2	x10 ⁵ Cpd 1: C14 H10 Cl2 N4 S: +ESI Scan (rt: 1.962, 2.178-2.212 min, 4 scans) Frag=350.0V C13M.d Subtract		
	Calculated monoisotopic mass:	8 - 337.0082		
	336.0003	6		
	Measured monoisotopic mass:	4		
	336.0017	2-338.0106 340.0084 341.0063		
	550.0017	0-1		
	Mass error: 3.96 ppm	Counts vs. Mass-to-Charge (m/z)		
C14	C14H10N4SCl2	Theoretical monoisotopic mass of [M+H] ⁺ : 337.0076		
	Calculated monoisotopic mass:	x10 ⁵ Cpd 1: C14 H10 Cl2 N4 S: +ESI Scan (rt: 2.141-2.175, 2.308-2.541 min, 18 scans) Frag=350.0V C14M.d Subt		
	336.0003	3.75 - 337.0079		
		3.0		
	Measured monoisotopic mass:	3-		
	336.0006	2.5 339.0050		
		2-		
	Mass error: 0.78 ppm	1.5		
		0.5 338.0106 340.0075 341.0021		
		0.25 342 0044 343.0013 344.0029		
		336 337 338 339 340 341 342 343 344 345 Counts vs. Mass-to-Charge (m/z)		



Figure 2S. A view of the molecules C1–C14 with the vectors of dipole moments in the lowenergy conformation calculated at the DFT/B3LYP/6-311++G(d,p) level.



Fig. 3S. The contour maps of molecular electrostatic potential for compounds **C1–C14** in the plane of 1,2,4-triazole ring; the red lines represent regions with the negative electrostatic potential and yellow lines represent regions with the positive electrostatic potential. *Molecular docking*



Fig. 4S. The crystal structure of the cytochrome P450 CYP121 of Mycobacterium tuberculosis obtained from expression system *Escherichia coli* in complex with 1-[(4-chlorophenyl)methyl]-4-(3-imidazol-1-ylpropyl)piperazin-2-one (9KE) and heme.



Fig. 5S. The overlay of molecules **C4** and **C14** in conformations observed in binding site of the cytochrome P450 CYP121 9red) and to their starting conformations for the docking process obtained as a result of energy minimization and energy optimization at DFT level (yellow).

Spectral data for new compounds:

Compound B13





Compound C13







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