

Supplementary Information File

A synthetic approach towards drugs modification: 2-hydroxy-1-naphthaldehyde based imine-zwitterion preparation, single-crystal study, Hirshfeld surface analysis, and computational investigation

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Contents:

- 1. Table S1: Enrichment ratio for the pair of chemical species in DSPIN. Enrichment ratio is not calculated for the pairs which random contact less than 0.99.**
- 2. Table S2: Enrichment ratio for the pair of chemical species in ACPIN. Enrichment ratio is not calculated for the pairs which random contact less than 0.99.**
- 3. Table S3: Interaction energies between molecular pairs in DSPIN.**
- 4. Table S4: Interaction energies between molecular pairs in ACPIN.**
- 5. checkCIF/PLATON report of DSPIN.**
- 6. checkCIF/PLATON report of ACPIN.**
- 7. Table S5: Selected structural parameters for the compounds DSPIN and ACPIN, calculated using the B3LYP/6-311+G(d,p) approach with the implicit solvent effects from ethanol.**
- 8. Table S6. Second-order perturbation theory analysis results for the compounds DSPIN and ACPIN, calculated using the B3LYP/6-311+G(d,p) approach with the implicit solvent effects from ethanol.**

Table S1: Enrichment ratio for the pair of chemical species in **DSPIN**. Enrichment ratio is not calculated for the pairs which random contact less than 0.99.

Contact %	Atom	H	C	N	O	S
	H	31.9	22.1	6.6	20	5.6
	C	22.1	7.1	1.2	3.5	0
	N	6.6	1.2	0	0.2	0
	O	20	3.5	0.2	0	1.8
	S	5.6	0	0	1.8	0
Surface%		59.05	20.5	4	12.75	3.7
Random contacts %	Atom	H	C	N	O	S
	H	34.87				
	C	24.21	4.20			
	N	4.72	1.64	0.16		
	O	15.06	5.23	1.02	1.63	
	S	4.37	1.52	0.30	0.94	0.14
Enrichment ratio	Atom	H	C	N	O	S
	H	0.91				
	C	0.91	1.69			
	N	1.40	0.73			
	O	1.33	0.67	0.20	0.00	
	S	1.28	0.00			

Table S2: Enrichment ratio for the pair of chemical species in ACPIN. Enrichment ratio is not calculated for the pairs which random contact less than 0.99.

Contact %	Atom	H	C	N	O	Cl
	H	45.1	19.5	6.7	5.2	11.4
	C	19.5	6.4	1.3	1.7	0.3
	N	6.7	1.3	0	0.3	0.1
	O	5.2	1.7	0.3	0.4	0.1
	Cl	11.4	0.3	0.1	0.1	1.5
Surface%		66.5	17.8	4.2	4.05	7.45
Random contacts %	Atom	H	C	N	O	Cl
	H	44.22				
	C	23.67	3.17			
	N	5.59	1.50	0.18		
	O	5.39	1.44	0.34	0.16	
	Cl	9.91	2.65	0.63	0.60	0.56
Enrichment ratio	Atom	H	C	N	O	Cl
	H	1.02				
	C	0.82	2.02			
	N	1.20	0.87			
	O	0.97	1.18			
	Cl	1.15	0.11			

Table S3: Interaction energies between molecular pairs in DSPIN.

Crystal	Atoms	Surface	Energies							
Interaction Energies (kJ/mol)										
R is the distance between molecular centroids (mean atomic position) in Å.										
Total energies, only reported for two benchmarked energy models, are the sum of the four energy components, scaled appropriately (see the scale factor table below)										

	N	Symop	R	Electron Density	E_ele	E_pol	E_dis	E_rep	E_tot	
	1	-x, -y, -z	7.34	B3LYP/6-31G(d,p)	-8.1	-3.6	-66.8	35.7	-47.3	
	2	-x, y+1/2, -z+1/2	16.26	B3LYP/6-31G(d,p)	-3.0	-0.5	-9.4	0.0	-11.7	
	2	-x, y+1/2, -z+1/2	9.29	B3LYP/6-31G(d,p)	-6.6	-3.2	-17.4	9.8	-18.5	
	1	x, y, z	7.84	B3LYP/6-31G(d,p)	-8.2	-3.5	-8.9	2.4	-17.5	
	0	-x, -y, -z	13.10	B3LYP/6-31G(d,p)	7.4	-1.7	-5.1	0.0	2.1	
	1	-x, -y, -z	6.32	B3LYP/6-31G(d,p)	-19.9	-7.5	-99.0	56.8	-77.7	
	2	x, -y+1/2, z+1/2	10.73	B3LYP/6-31G(d,p)	-15.8	-5.0	-16.5	23.9	-20.1	
	0	-x, y+1/2, -z+1/2	7.93	B3LYP/6-31G(d,p)	-21.3	-8.9	-33.4	29.0	-40.2	
	2	x, -y+1/2, z+1/2	11.34	B3LYP/6-31G(d,p)	-2.5	-1.5	-20.1	11.2	-14.3	
	0	-x, -y, -z	11.78	B3LYP/6-31G(d,p)	-121.0	-29.5	-30.7	99.1	-115.3	

Scale factors for benchmarked energy models										
See Mackenzie et al. IUCrJ (2017)										

Energy Model	k_ele	k_pol	k_disp	k_rep						
CE-HF ... HF/3-21G electron densities	1.019	0.651	0.901	0.811						
CE-B3LYP ... B3LYP/6-31G(d,p) electron densities	1.057	0.740	0.871	0.618						

Table S4: Interaction energies between molecular pairs in ACPIN.

Interaction Energies (kJ/mol)

R is the distance between molecular centroids (mean atomic position) in Å.

Total energies, only reported for two benchmarked energy models, are the sum of the four energy components, scaled appropriately (see the scale factor table below)

	N	Symop	R	Electron Density	E_ele	E_pol	E_dis	E_rep	E_tot
	1	-x, -y, -z	18.35	B3LYP/6-31G(d,p)	-0.7	-0.3	-11.3	0.0	-10.8
	2	x, y, z	4.54	B3LYP/6-31G(d,p)	1.2	-4.4	-91.7	43.7	-54.9
	1	-x, -y, -z	6.34	B3LYP/6-31G(d,p)	-11.4	-2.4	-52.1	28.7	-41.5
	2	x, y, z	18.31	B3LYP/6-31G(d,p)	-1.0	-0.1	-4.5	0.0	-5.0
	1	-x, -y, -z	7.26	B3LYP/6-31G(d,p)	-91.3	-21.3	-41.7	94.2	-90.4
	2	x, y, z	17.67	B3LYP/6-31G(d,p)	2.6	-0.2	-7.6	0.0	-4.0
	1	-x, -y, -z	6.25	B3LYP/6-31G(d,p)	-19.3	-7.1	-47.7	20.9	-54.3
	1	-x, -y, -z	18.64	B3LYP/6-31G(d,p)	0.0	-0.1	-7.9	0.0	-7.0
	1	-x, -y, -z	7.29	B3LYP/6-31G(d,p)	-6.7	-0.9	-47.7	29.1	-31.3

Scale factors for benchmarked energy models

See Mackenzie et al. IUCrJ (2017)

Energy Model	k_ele	k_pol	k_disp	k_rep
CE-HF ... HF/3-21G electron densities	1.019	0.651	0.901	0.811
CE-B3LYP ... B3LYP/6-31G(d,p) electron densities	1.057	0.740	0.871	0.618

checkCIF/PLATON report of DSPIN

Structure factors have been supplied for datablock(s) napsuly_0m

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

No syntax errors found. CIF dictionary Interpreting this report

Datablock: DSPIN.

Bond precision: C-C = 0.0034 A Wavelength=0.71073

Cell: a=7.8389(8) b=13.8151(16) c=17.198(2)
 alpha=90 beta=94.032(4) gamma=90
 Temperature: 296 K

	Calculated	Reported
Volume	1857.9(4)	1857.9(4)
Space group	P 21/c	P 21/c
Hall group	-P 2ybc	-P 2ybc
Moiety formula	C20 H15 N3 O3 S2	C20 H15 N3 O3 S2
Sum formula	C20 H15 N3 O3 S2	C20 H15 N3 O3 S2
Mr	409.47	409.47
Dx, g cm ⁻³	1.464	1.464
Z	4	4
Mu (mm ⁻¹)	0.314	0.314
F000	848.0	848.0
F000'	849.32	
h, k, lmax	10, 17, 21	9, 17, 21
Nref	4049	4040
Tmin, Tmax	0.904, 0.930	0.869, 0.894
Tmin'	0.904	

Correction method= # Reported T Limits: Tmin=0.869 Tmax=0.894

AbsCorr = MULTI-SCAN

Data completeness= 0.998

Theta (max)= 26.999

R(reflections)= 0.0437 (2548)

wR2 (reflections)=

0.1087 (4040)

S = 1.018

Npar= 259

The following ALERTS were generated. Each ALERT has the format

test-name_ALERT_alert-type_alert-level.

Click on the hyperlinks for more details of the test.

Alert level C

PLAT230_ALERT_2_C Hirshfeld Test Diff for	S1	--O2	.	6.6 s.u.
PLAT334_ALERT_2_C Small <C-C> Benzene Dist.	C12	-C17	.	1.37 Ang.
PLAT905_ALERT_3_C Negative K value in the Analysis of Variance ... -1.284 Report				
PLAT911_ALERT_3_C Missing FCF Refl				
Between Thmin & STh/L=	0.600		2	Report

Alert level G

PLAT333_ALERT_2_G Large Aver C6-Ring C-C Dist C1		-C10	.	1.42 Ang.
PLAT432_ALERT_2_G Short Inter X...Y Contact O1		..C19	.	3.00 Ang.
		x,1/2-y,-1/2+z =		4_565 Check
PLAT480_ALERT_4_G Long H...A H-Bond Reported H3A		..S1	.	2.89 Ang.
PLAT480_ALERT_4_G Long H...A H-Bond Reported H11		..O3	.	2.64 Ang.
PLAT910_ALERT_3_G Missing # of FCF Reflection(s) Below Theta(Min).				1 Note
PLAT912_ALERT_4_G Missing # of FCF Reflections Above STh/L= 0.600				6 Note
PLAT933_ALERT_2_G Number of HKL-OMIT Records in Embedded .res File				1 Note
PLAT941_ALERT_3_G Average HKL Measurement Multiplicity				3.8 Low
PLAT967_ALERT_5_G Note: Two-Theta Cutoff Value in Embedded .res ..				54.0 Degree
PLAT978_ALERT_2_G Number C-C Bonds with Positive Residual Density.				1 Info

0 **ALERT level A** = Most likely a serious problem - resolve or explain

0 **ALERT level B** = A potentially serious problem, consider carefully

4 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight

10 **ALERT level G** = General information/check it is not something unexpected

0 ALERT type 1 CIF construction/syntax error, inconsistent or missing data

6 ALERT type 2 Indicator that the structure model may be wrong or deficient
4 ALERT type 3 Indicator that the structure quality may be low
3 ALERT type 4 Improvement, methodology, query or suggestion
1 ALERT type 5 Informative message, check

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

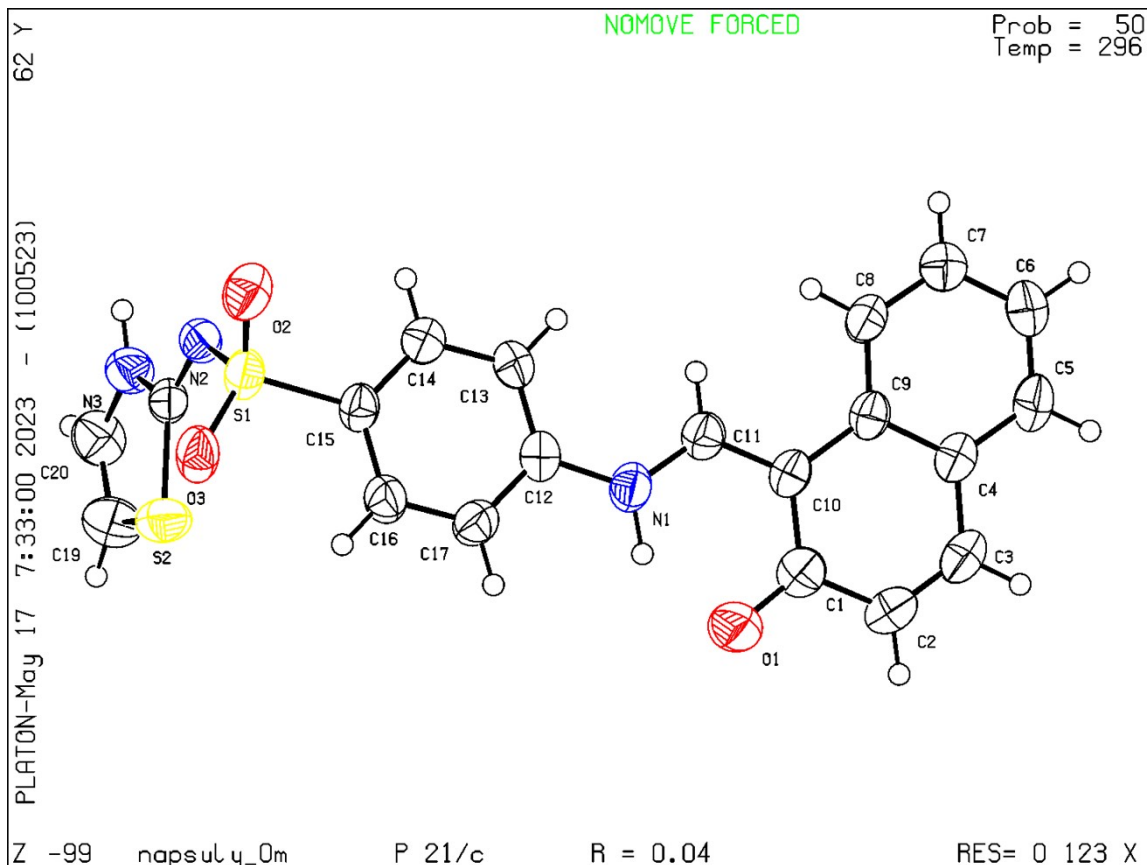
Publication of your CIF in IUCr journals

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

Publication of your CIF in other journals

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

Datablock napsuly_0m - ellipsoid plot

**checkCIF/PLATON report of ACPIN.**

Structure factors have been supplied for datablock(s) pymtvan_twnew

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No syntax errors found. CIF dictionary Interpreting this report

Datablock: ACPIN

Bond precision: C-C = 0.0057 A

Wavelength=0.71073

Cell: a=4.5436(6) b=12.5255(16) c=17.673(2)
 alpha=88.308(8) beta=89.195(8) gamma=82.611(9)
 Temperature: 296 K

	Calculated	Reported
Volume	997.0(2)	996.9(2)
Space group	P -1	P -1
Hall group	-P 1	-P 1
Moiety formula	C23 H19 Cl N4 O	C23 H19 Cl N4 O
Sum formula	C23 H19 Cl N4 O	C23 H19 Cl N4 O
Mr	402.87	402.87
Dx, g cm ⁻³	1.342	1.324
Z	2	2
Mu (mm ⁻¹)	0.214	0.214
F000	420.0	420.0
F000'	420.44	
h, k, lmax	5, 16, 22	5, 16, 22
Nref	4543	4351
Tmin, Tmax	0.940, 0.958	0.869, 0.894
Tmin'	0.914	

Correction method= # Reported T Limits: Tmin=0.869 Tmax=0.894

AbsCorr = MULTI-SCAN

Data completeness= 0.958

Theta(max)= 27.448

R(reflections)= 0.0763(2397)

wR2(reflections)=
0.2358(4351)

S = 1.139

Npar= 264

The following ALERTS were generated. Each ALERT has the format

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Click on the hyperlinks for more details of the test.

■ Alert level C

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PLAT046_ALERT_1_C	Reported Z, MW and D(calc) are Inconsistent		1.342 Check
PLAT242_ALERT_2_C	Low 'MainMol' Ueq as Compared to Neighbors of		C21 Check
PLAT340_ALERT_3_C	Low Bond Precision on C-C Bonds	0.00574 Ang.	
PLAT906_ALERT_3_C	Large K Value in the Analysis of Variance		29.015 Check
PLAT906_ALERT_3_C	Large K Value in the Analysis of Variance		2.364 Check
PLAT911_ALERT_3_C	Missing FCF Refl Between Thmin & STh/L=	0.600	121 Report

■ Alert level G

PLAT007_ALERT_5_G	Number of Unrefined Donor-H Atoms		3 Report
PLAT333_ALERT_2_G	Large Aver C6-Ring C-C Dist C1	-C10	1.43 Ang.
PLAT870_ALERT_4_G	ALERTS Related to Twinning Effects Suppressed ..		! Info
PLAT910_ALERT_3_G	Missing # of FCF Reflection(s) Below Theta(Min).		1 Note
PLAT912_ALERT_4_G	Missing # of FCF Reflections Above STh/L= 0.600		70 Note
PLAT931_ALERT_5_G	CIFcalcFCF Twin Law (0 0 1)	Est.d BASF	0.18 Check
PLAT941_ALERT_3_G	Average HKL Measurement Multiplicity		1.0 Low

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2 ALERT type 5 Informative message, check

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

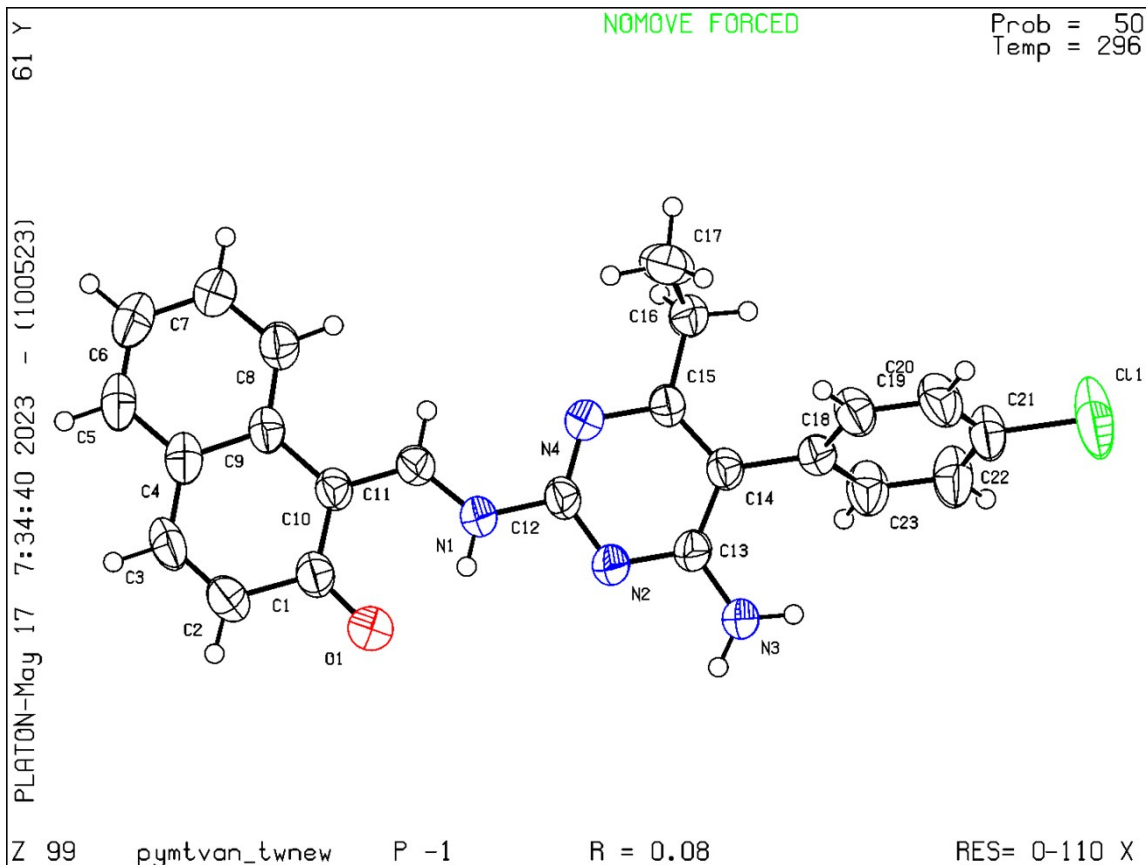
Publication of your CIF in IUCr journals

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

Publication of your CIF in other journals

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

Datablock pymtvan_twnew - ellipsoid plot

**CIF of DSPIN without hkl data**

data_global

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PROCESSING SUMMARY (IUCr Office Use only)

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Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.

Farrugia, L. J. (2012). *J. Appl. Cryst.* **45**, 849--854.

Sheldrick, G. M. (2008). *Acta Cryst.* A **64**, 112--122.

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They are only intended as comments.

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_cell_angle_gamma    90
_cell_volume         1857.9(4)
_cell_formula_units_Z  4
_cell_measurement_reflns_used  2548
_cell_measurement_theta_min  2.374
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_cell_measurement_theta_max      27.000
_cell_measurement_temperature     296(2)
_exptl_crystal_description       prism
_exptl_crystal_colour            'red'
_exptl_crystal_size_max         0.32
_exptl_crystal_size_mid         0.28
_exptl_crystal_size_min         0.23
_exptl_crystal_density_diffn     1.464
_exptl_crystal_density_meas     ?
_exptl_crystal_density_method   'not measured'
_exptl_crystal_F_000            848
_exptl_absorpt_coefficient_mu    0.314
_exptl_absorpt_correction_type   multi-scan
_exptl_absorpt_process_details   '(SADABS; Bruker, 2007)'
_exptl_absorpt_correction_T_min 0.869
_exptl_absorpt_correction_T_max 0.894
_exptl_special_details
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_diffrn_ambient_temperature     296(2)
_diffrn_radiation_type          MoK\alpha
_diffrn_radiation_wavelength    0.71073
_diffrn_radiation_source        'fine-focus sealed tube'
_diffrn_radiation_monochromator  graphite
_diffrn_measurement_device_type  'Bruker Kappa APEXII CCD'
_diffrn_measurement_method      \w
_diffrn_detector_area_resol_mean 7.828
_diffrn_reflns_number           15333
_diffrn_reflns_av_unetl/netl    0.0473
_diffrn_reflns_av_R_equivalents 0.0404
_diffrn_reflns_limit_h_min      -9
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_diffrn_reflns_limit_h_max    9
_diffrn_reflns_limit_k_min   -17
_diffrn_reflns_limit_k_max    15
_diffrn_reflns_limit_l_min   -21
_diffrn_reflns_limit_l_max    21
_diffrn_reflns_theta_min     2.374
_diffrn_reflns_theta_max     26.999
_diffrn_reflns_theta_full    25.242
_diffrn_measured_fraction_theta_max 0.998
_diffrn_measured_fraction_theta_full 0.999
_diffrn_reflns_Laue_measured_fraction_max 0.998
_diffrn_reflns_Laue_measured_fraction_full 0.999
_diffrn_reflns_point_group_measured_fraction_max 0.998
_diffrn_reflns_point_group_measured_fraction_full 0.999
_diffrn_standards_number    0
_diffrn_standards_interval_count  ?
_diffrn_standards_interval_time  ?
_diffrn_standards_decay_%   ?

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Refinement of $\langle F \rangle^2$ against ALL reflections. The weighted $\langle R \rangle$ -factor $\langle wR \rangle$ and goodness of fit $\langle S \rangle$ are based on $\langle F \rangle^2$, conventional $\langle R \rangle$ -factors $\langle R \rangle$ are based on $\langle F \rangle$, with $\langle F \rangle$ set to zero for negative $\langle F \rangle^2$. The threshold expression of $\langle F \rangle^2 > \sqrt{\langle F \rangle^2}$ is used only for calculating $\langle R \rangle$ -factors(gt) etc. and is not relevant to the choice of reflections for refinement.

$\langle R \rangle$ -factors based on $\langle F \rangle^2$ are statistically about twice as large as those based on $\langle F \rangle$, and $\langle R \rangle$ - factors based on ALL data will be even larger.

```
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```

```

_reflns_number_total    4040
_reflns_number_gt       2548

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_refine_ls_structure_factor_coef  Fsqd
_refine_ls_matrix_type  full
_refine_ls_R_factor_all  0.0852
_refine_ls_R_factor_gt  0.0437
_refine_ls_wR_factor_ref  0.1087
_refine_ls_wR_factor_gt  0.0916
_refine_ls_goodness_of_fit_ref  1.018
_refine_ls_restrained_S_all  1.018
_refine_ls_number_reflns  4040
_refine_ls_number_parameters  259
_refine_ls_number_restraints  0
_refine_ls_extinction_method  none
_refine_ls_extinction_coef  .
_refine_ls_hydrogen_treatment  mixed
_refine_ls_weighting_scheme  calc
_refine_ls_weighting_details
'w=1/[\s^2^(Fo^2^)+(0.0429P)^2^+0.3132P] where P=(Fo^2^+2Fc^2^)/3'
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_atom_sites_solution_primary  direct
_atom_sites_solution_secondary  difmap
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_refine_ls_shift/su_mean  0.000
_computing_data_collection  'APEX2 (Bruker, 2007)'
_computing_cell_refinement  'SAINT (Bruker, 2007)'
_computing_data_reduction  'SAINT (Bruker, 2007)'
_computing_structure_solution  'SHELXT2014 (Sheldrick, 2008)'
_computing_structure_refinement  'SHELXL-2019/2 (Sheldrick, 2015)'
_computing_molecular_graphics
'ORTEP-3 for Windows (Farrugia, 2012) and PLATON (Spek, 2009)'
_computing_publication_material
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WinGX (Farrugia, 2012) and PLATON (Spek, 2009)

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_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

S1 S 0.62806(8) 0.37379(5) 0.35983(3) 0.04756(19) Uani 1 1 d

S2 S 0.27019(9) 0.27664(5) 0.41242(4) 0.0608(2) Uani 1 1 d

O1 O 0.1217(2) 0.41542(13) -0.06501(11) 0.0680(5) Uani 1 1 d

O2 O 0.7889(2) 0.42300(13) 0.36955(9) 0.0608(5) Uani 1 1 d

O3 O 0.6259(2) 0.27012(12) 0.36427(9) 0.0600(5) Uani 1 1 d

N1 N 0.3001(2) 0.48144(15) 0.05142(11) 0.0486(5) Uani 1 1 d

H1 H 0.238(3) 0.4297(17) 0.0229(14) 0.058 Uiso 1 1 d . U . . .

N2 N 0.5153(2) 0.42218(13) 0.42388(10) 0.0448(5) Uani 1 1 d

N3 N 0.2837(3) 0.42242(15) 0.49937(12) 0.0508(5) Uani 1 1 d

H3A H 0.322(3) 0.4750(18) 0.5249(14) 0.061 Uiso 1 1 d . U . . .

C1 C 0.1231(3) 0.49923(19) -0.09520(14) 0.0509(6) Uani 1 1 d

C2 C 0.0363(3) 0.5141(2) -0.17026(14) 0.0572(7) Uani 1 1 d

H2 H -0.019101 0.462165 -0.195502 0.069 Uiso 1 1 calc R U . . .

C3 C 0.0327(3) 0.6003(2) -0.20484(14) 0.0566(7) Uani 1 1 d

H3 H -0.025917 0.606851 -0.253469 0.068 Uiso 1 1 calc R U . . .
C4 C 0.1162(3) 0.68335(18) -0.16976(13) 0.0479(6) Uani 1 1 d
C5 C 0.1148(3) 0.7723(2) -0.20827(14) 0.0652(8) Uani 1 1 d
H5 H 0.055085 0.778511 -0.256678 0.078 Uiso 1 1 calc R U . . .
C6 C 0.1997(4) 0.8502(2) -0.17602(15) 0.0700(8) Uani 1 1 d
H6 H 0.197231 0.909269 -0.201978 0.084 Uiso 1 1 calc R U . . .
C7 C 0.2892(3) 0.8406(2) -0.10451(14) 0.0630(7) Uani 1 1 d
H7 H 0.348124 0.893463 -0.082612 0.076 Uiso 1 1 calc R U . . .
C8 C 0.2926(3) 0.75469(18) -0.06548(13) 0.0528(6) Uani 1 1 d
H8 H 0.353845 0.750163 -0.017294 0.063 Uiso 1 1 calc R U . . .
C9 C 0.2064(3) 0.67346(17) -0.09614(12) 0.0424(5) Uani 1 1 d
C10 C 0.2088(3) 0.57993(17) -0.05741(12) 0.0422(5) Uani 1 1 d
C11 C 0.2919(3) 0.56653(18) 0.01579(12) 0.0460(6) Uani 1 1 d
H11 H 0.343968 0.619554 0.040893 0.055 Uiso 1 1 calc R U . . .
C12 C 0.3795(3) 0.46035(17) 0.12503(12) 0.0433(5) Uani 1 1 d
C13 C 0.4622(3) 0.52816(18) 0.17266(13) 0.0536(6) Uani 1 1 d
H13 H 0.467179 0.592349 0.156691 0.064 Uiso 1 1 calc R U . . .
C14 C 0.5374(3) 0.50080(18) 0.24386(13) 0.0542(6) Uani 1 1 d
H14 H 0.594324 0.546593 0.275698 0.065 Uiso 1 1 calc R U . . .
C15 C 0.5292(3) 0.40645(17) 0.26843(12) 0.0422(5) Uani 1 1 d
C16 C 0.4473(3) 0.33821(17) 0.22109(13) 0.0494(6) Uani 1 1 d
H16 H 0.442650 0.274010 0.237083 0.059 Uiso 1 1 calc R U . . .
C17 C 0.3723(3) 0.36588(18) 0.14980(13) 0.0501(6) Uani 1 1 d
H17 H 0.315809 0.319999 0.117875 0.060 Uiso 1 1 calc R U . . .
C18 C 0.3718(3) 0.38159(16) 0.44415(12) 0.0431(5) Uani 1 1 d
C19 C 0.1106(3) 0.2962(2) 0.47421(17) 0.0673(8) Uani 1 1 d
H19 H 0.016556 0.255785 0.477583 0.081 Uiso 1 1 calc R U . . .
C20 C 0.1373(3) 0.3750(2) 0.51617(16) 0.0615(7) Uani 1 1 d
H20 H 0.064756 0.396229 0.553157 0.074 Uiso 1 1 calc R U . . .

loop_

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_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

S1 0.0609(4) 0.0475(4) 0.0340(3) 0.0018(3) 0.0011(3) 0.0097(3)
S2 0.0707(5) 0.0502(4) 0.0613(4) -0.0118(3) 0.0032(3) -0.0108(3)
O1 0.0838(13) 0.0467(12) 0.0711(12) -0.0001(9) -0.0100(10) 0.0007(10)
O2 0.0509(10) 0.0817(13) 0.0494(10) 0.0020(9) 0.0002(8) 0.0001(9)
O3 0.0929(13) 0.0448(11) 0.0420(9) 0.0018(8) 0.0019(8) 0.0207(9)
N1 0.0544(12) 0.0511(14) 0.0393(10) 0.0037(9) -0.0027(9) 0.0033(10)
N2 0.0608(12) 0.0377(11) 0.0364(10) -0.0015(8) 0.0079(9) -0.0008(10)
N3 0.0611(13) 0.0444(13) 0.0479(12) -0.0040(10) 0.0103(10) -0.0030(11)
C1 0.0504(14) 0.0508(17) 0.0512(14) -0.0031(12) 0.0011(11) 0.0082(12)
C2 0.0574(16) 0.0615(18) 0.0513(15) -0.0113(13) -0.0067(12) -0.0043(13)
C3 0.0542(15) 0.075(2) 0.0393(13) -0.0036(13) -0.0071(11) 0.0000(14)
C4 0.0484(14) 0.0577(17) 0.0369(11) 0.0007(11) -0.0011(10) 0.0016(12)
C5 0.0754(18) 0.074(2) 0.0436(14) 0.0155(14) -0.0134(13) -0.0016(16)
C6 0.091(2) 0.0604(19) 0.0562(16) 0.0208(14) -0.0119(15) -0.0061(16)
C7 0.084(2) 0.0512(17) 0.0523(15) 0.0027(13) -0.0073(14) -0.0059(14)
C8 0.0682(17) 0.0525(16) 0.0361(12) -0.0005(11) -0.0083(11) 0.0005(13)
C9 0.0437(13) 0.0497(15) 0.0340(11) -0.0001(10) 0.0043(9) 0.0066(11)
C10 0.0416(13) 0.0501(15) 0.0347(11) -0.0019(10) 0.0021(9) 0.0055(11)
C11 0.0495(14) 0.0483(16) 0.0401(12) -0.0002(11) 0.0032(10) 0.0033(11)
C12 0.0436(13) 0.0490(15) 0.0377(12) 0.0057(11) 0.0051(10) 0.0073(11)
C13 0.0730(17) 0.0409(15) 0.0457(13) 0.0098(11) -0.0040(12) -0.0028(13)
C14 0.0713(17) 0.0461(16) 0.0439(13) 0.0012(11) -0.0059(12) -0.0030(13)
C15 0.0488(13) 0.0446(15) 0.0336(11) 0.0010(10) 0.0047(10) 0.0079(11)
C16 0.0659(16) 0.0398(14) 0.0422(12) 0.0031(11) 0.0018(11) 0.0056(12)
C17 0.0598(16) 0.0470(16) 0.0424(13) -0.0051(11) -0.0032(11) 0.0016(12)
C18 0.0574(15) 0.0375(13) 0.0340(11) 0.0023(10) 0.0000(10) 0.0030(12)
C19 0.0589(17) 0.0621(19) 0.081(2) -0.0010(16) 0.0080(14) -0.0108(14)
C20 0.0615(18) 0.0620(19) 0.0628(16) 0.0005(14) 0.0160(13) 0.0012(15)

_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.

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_geom_bond_site_symmetry_2

_geom_bond_publ_flag

S1 O2 1.4315(17) . ?

S1 O3 1.4344(17) . ?

S1 N2 1.6056(18) . ?

S1 C15 1.762(2) . ?

S2 C19 1.719(3) . ?

S2 C18 1.724(2) . ?

O1 C1 1.269(3) . ?

N1 C11 1.325(3) . ?

N1 C12 1.402(3) . ?

N1 H1 0.98(2) . ?

N2 C18 1.325(3) . ?

N3 C18 1.338(3) . ?

N3 C20 1.370(3) . ?

N3 H3A 0.89(2) . ?

C1 C2 1.431(3) . ?

C1 C10 1.434(3) . ?

C2 C3 1.331(3) . ?
C2 H2 0.9300 . ?
C3 C4 1.433(3) . ?
C3 H3 0.9300 . ?
C4 C5 1.396(3) . ?
C4 C9 1.413(3) . ?
C5 C6 1.362(4) . ?
C5 H5 0.9300 . ?
C6 C7 1.379(3) . ?
C6 H6 0.9300 . ?
C7 C8 1.362(3) . ?
C7 H7 0.9300 . ?
C8 C9 1.394(3) . ?
C8 H8 0.9300 . ?
C9 C10 1.453(3) . ?
C10 C11 1.388(3) . ?
C11 H11 0.9300 . ?
C12 C17 1.375(3) . ?
C12 C13 1.376(3) . ?
C13 C14 1.374(3) . ?
C13 H13 0.9300 . ?
C14 C15 1.373(3) . ?
C14 H14 0.9300 . ?
C15 C16 1.375(3) . ?
C16 C17 1.376(3) . ?
C16 H16 0.9300 . ?
C17 H17 0.9300 . ?
C19 C20 1.315(3) . ?
C19 H19 0.9300 . ?
C20 H20 0.9300 . ?

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O2 S1 O3 118.78(11) .. ?
O2 S1 N2 104.17(10) .. ?
O3 S1 N2 111.73(10) .. ?
O2 S1 C15 108.12(11) .. ?
O3 S1 C15 107.28(10) .. ?
N2 S1 C15 106.06(10) .. ?
C19 S2 C18 90.71(13) .. ?
C11 N1 C12 127.3(2) .. ?
C11 N1 H1 114.3(14) .. ?
C12 N1 H1 118.3(14) .. ?
C18 N2 S1 121.50(16) .. ?
C18 N3 C20 115.5(2) .. ?
C18 N3 H3A 121.7(16) .. ?
C20 N3 H3A 122.8(16) .. ?
O1 C1 C2 118.8(2) .. ?
O1 C1 C10 122.8(2) .. ?
C2 C1 C10 118.4(2) .. ?
C3 C2 C1 121.7(2) .. ?
C3 C2 H2 119.2 .. ?
C1 C2 H2 119.2 .. ?
C2 C3 C4 122.3(2) .. ?
C2 C3 H3 118.9 .. ?
C4 C3 H3 118.9 .. ?
C5 C4 C9 119.8(2) .. ?
C5 C4 C3 121.2(2) .. ?
C9 C4 C3 119.0(2) .. ?
C6 C5 C4 121.0(2) .. ?

C6 C5 H5 119.5 . . ?
C4 C5 H5 119.5 . . ?
C5 C6 C7 119.4(2) . . ?
C5 C6 H6 120.3 . . ?
C7 C6 H6 120.3 . . ?
C8 C7 C6 121.0(2) . . ?
C8 C7 H7 119.5 . . ?
C6 C7 H7 119.5 . . ?
C7 C8 C9 121.5(2) . . ?
C7 C8 H8 119.3 . . ?
C9 C8 H8 119.3 . . ?
C8 C9 C4 117.4(2) . . ?
C8 C9 C10 123.51(19) . . ?
C4 C9 C10 119.1(2) . . ?
C11 C10 C1 118.8(2) . . ?
C11 C10 C9 121.6(2) . . ?
C1 C10 C9 119.55(19) . . ?
N1 C11 C10 122.8(2) . . ?
N1 C11 H11 118.6 . . ?
C10 C11 H11 118.6 . . ?
C17 C12 C13 119.3(2) . . ?
C17 C12 N1 116.8(2) . . ?
C13 C12 N1 123.8(2) . . ?
C14 C13 C12 119.8(2) . . ?
C14 C13 H13 120.1 . . ?
C12 C13 H13 120.1 . . ?
C15 C14 C13 120.6(2) . . ?
C15 C14 H14 119.7 . . ?
C13 C14 H14 119.7 . . ?
C14 C15 C16 119.9(2) . . ?
C14 C15 S1 119.37(17) . . ?
C16 C15 S1 120.70(18) . . ?
C15 C16 C17 119.3(2) . . ?

C15 C16 H16 120.3 . . ?
C17 C16 H16 120.3 . . ?
C12 C17 C16 121.0(2) . . ?
C12 C17 H17 119.5 . . ?
C16 C17 H17 119.5 . . ?
N2 C18 N3 119.9(2) . . ?
N2 C18 S2 130.88(18) . . ?
N3 C18 S2 109.21(18) . . ?
C20 C19 S2 112.0(2) . . ?
C20 C19 H19 124.0 . . ?
S2 C19 H19 124.0 . . ?
C19 C20 N3 112.5(2) . . ?
C19 C20 H20 123.7 . . ?
N3 C20 H20 123.7 . . ?

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_geom_torsion_site_symmetry_1
_geom_torsion_site_symmetry_2
_geom_torsion_site_symmetry_3
_geom_torsion_site_symmetry_4
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O2 S1 N2 C18 -165.13(17) ?
O3 S1 N2 C18 -35.7(2) ?
C15 S1 N2 C18 80.88(19) ?
O1 C1 C2 C3 179.8(2) ?
C10 C1 C2 C3 -0.4(4) ?
C1 C2 C3 C4 0.5(4) ?
C2 C3 C4 C5 178.1(3) ?

C2 C3 C4 C9 0.5(4) ?
C9 C4 C5 C6 0.1(4) ?
C3 C4 C5 C6 -177.5(3) ?
C4 C5 C6 C7 0.5(4) ?
C5 C6 C7 C8 -0.6(4) ?
C6 C7 C8 C9 0.1(4) ?
C7 C8 C9 C4 0.5(4) ?
C7 C8 C9 C10 178.8(2) ?
C5 C4 C9 C8 -0.6(3) ?
C3 C4 C9 C8 177.1(2) ?
C5 C4 C9 C10 -179.0(2) ?
C3 C4 C9 C10 -1.4(3) ?
O1 C1 C10 C11 -0.9(3) ?
C2 C1 C10 C11 179.3(2) ?
O1 C1 C10 C9 179.3(2) ?
C2 C1 C10 C9 -0.5(3) ?
C8 C9 C10 C11 3.3(3) ?
C4 C9 C10 C11 -178.4(2) ?
C8 C9 C10 C1 -176.9(2) ?
C4 C9 C10 C1 1.4(3) ?
C12 N1 C11 C10 -179.7(2) ?
C1 C10 C11 N1 2.2(3) ?
C9 C10 C11 N1 -178.0(2) ?
C11 N1 C12 C17 -179.6(2) ?
C11 N1 C12 C13 0.6(4) ?
C17 C12 C13 C14 0.5(4) ?
N1 C12 C13 C14 -179.7(2) ?
C12 C13 C14 C15 -0.7(4) ?
C13 C14 C15 C16 1.0(4) ?
C13 C14 C15 S1 179.64(19) ?
O2 S1 C15 C14 -40.8(2) ?
O3 S1 C15 C14 -169.98(18) ?
N2 S1 C15 C14 70.5(2) ?

O2 S1 C15 C16 137.92(19) ?
 O3 S1 C15 C16 8.7(2) ?
 N2 S1 C15 C16 -110.84(19) ?
 C14 C15 C16 C17 -0.9(3) ?
 S1 C15 C16 C17 -179.56(18) ?
 C13 C12 C17 C16 -0.4(3) ?
 N1 C12 C17 C16 179.7(2) ?
 C15 C16 C17 C12 0.6(3) ?
 S1 N2 C18 N3 178.04(16) ?
 S1 N2 C18 S2 -0.2(3) ?
 C20 N3 C18 N2 180.0(2) ?
 C20 N3 C18 S2 -1.4(3) ?
 C19 S2 C18 N2 179.9(2) ?
 C19 S2 C18 N3 1.55(18) ?
 C18 S2 C19 C20 -1.4(2) ?
 S2 C19 C20 N3 0.9(3) ?
 C18 N3 C20 C19 0.3(3) ?

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N1 H1 O1 0.98(2) 1.72(2) 2.530(3) 138(2) . yes
 N3 H3A S1 0.89(2) 2.89(3) 3.746(2) 163(2) 3_666 yes
 N3 H3A O2 0.89(2) 2.50(2) 3.185(3) 134(2) 3_666 yes
 N3 H3A N2 0.89(2) 2.07(2) 2.922(3) 161(2) 3_666 yes
 C11 H11 O3 0.93 2.64 3.521(3) 157.4 2_655 yes

C13 H13 O3 0.93 2.58 3.463(3) 158.7 2_655 yes

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TITL NAPSULY_0m in P2(1)/c RED PRISM

shelx.res

created by SHELXL-2019/2 at 12:42:26 on 18-Feb-2023

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ZERR 4.00 0.0008 0.0016 0.0021 0.000 0.004 0.000

LATT 1

SYMM -x, y+1/2, -z+1/2

SFAC C H N O S

UNIT 80 60 12 12 8

OMIT -3 54

L.S. 40

ACTA

BOND \$H

FMAP 2

PLAN 10

CONF

HTAB

TEMP 23.000

SIZE 0.32 0.28 0.23

HTAB N1 O1

EQIV \$1 -x+1, -y+1, -z+1

HTAB N3 S1_\$1

HTAB N3 O2_\$1

HTAB N3 N2_\$1

EQIV \$2 -x+1, y+1/2, -z+1/2

HTAB C11 O3_\$2

HTAB C13 O3_\$2

OMIT 0 1 1

MPLA 6 N2 C18 C19 C20 S2 N3

MPLA 12 C1 C2 C3 C4 C5 C6 C7 C8 C9 C10 C11 O1

MPLA 7 N1 C12 C13 C14 C15 C16 C17

MPLA 3 S1 O2 O3

MPLA 6 N2 C18 C19 C20 S2 N3

MPLA 7 N1 C12 C13 C14 C15 C16 C17

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MPLA 5 C18 C19 C20 S2 N3

WGHT 0.042900 0.313200

FVAR 0.09598

S1 5 0.628057 0.373793 0.359833 11.00000 0.06087 0.04754 =
0.03397 0.00184 0.00113 0.00970

S2 5 0.270187 0.276639 0.412422 11.00000 0.07069 0.05016 =
0.06133 -0.01180 0.00315 -0.01084

O1 4 0.121683 0.415423 -0.065008 11.00000 0.08385 0.04670 =
0.07113 -0.00012 -0.00997 0.00069

O2 4 0.788859 0.422998 0.369546 11.00000 0.05088 0.08175 =
0.04944 0.00204 0.00020 0.00012

O3 4 0.625926 0.270120 0.364266 11.00000 0.09287 0.04482 =
0.04200 0.00183 0.00191 0.02066

N1 3 0.300087 0.481441 0.051420 11.00000 0.05439 0.05107 =
0.03935 0.00374 -0.00274 0.00327

H1 2 0.238428 0.429667 0.022870 11.00000 -1.20000

N2 3 0.515331 0.422184 0.423883 11.00000 0.06085 0.03769 =
0.03636 -0.00150 0.00791 -0.00083

N3 3 0.283740 0.422420 0.499368 11.00000 0.06113 0.04441 =
0.04787 -0.00396 0.01028 -0.00303

H3A 2 0.321545 0.474981 0.524945 11.00000 -1.20000

C1 1 0.123148 0.499229 -0.095202 11.00000 0.05042 0.05084 =

0.05121 -0.00314 0.00113 0.00818

C2 1 0.036279 0.514085 -0.170260 11.00000 0.05739 0.06149 =
0.05134 -0.01130 -0.00670 -0.00432

AFIX 43

H2 2 -0.019101 0.462165 -0.195502 11.00000 -1.20000

AFIX 0

C3 1 0.032738 0.600287 -0.204843 11.00000 0.05419 0.07498 =
0.03926 -0.00359 -0.00714 -0.00005

AFIX 43

H3 2 -0.025917 0.606851 -0.253469 11.00000 -1.20000

AFIX 0

C4 1 0.116162 0.683350 -0.169763 11.00000 0.04839 0.05775 =
0.03686 0.00069 -0.00107 0.00162

C5 1 0.114817 0.772307 -0.208268 11.00000 0.07536 0.07401 =
0.04361 0.01547 -0.01344 -0.00161

AFIX 43

H5 2 0.055085 0.778511 -0.256678 11.00000 -1.20000

AFIX 0

C6 1 0.199669 0.850179 -0.176016 11.00000 0.09102 0.06040 =
0.05615 0.02085 -0.01190 -0.00611

AFIX 43

H6 2 0.197231 0.909269 -0.201978 11.00000 -1.20000

AFIX 0

C7 1 0.289205 0.840560 -0.104515 11.00000 0.08374 0.05122 =
0.05233 0.00275 -0.00733 -0.00587

AFIX 43

H7 2 0.348124 0.893463 -0.082612 11.00000 -1.20000

AFIX 0

C8 1 0.292632 0.754694 -0.065476 11.00000 0.06816 0.05254 =
0.03611 -0.00046 -0.00834 0.00051

AFIX 43

H8 2 0.353845 0.750163 -0.017294 11.00000 -1.20000

AFIX 0

C9 1 0.206431 0.673458 -0.096136 11.00000 0.04371 0.04969 =
0.03400 -0.00011 0.00425 0.00660

C10 1 0.208811 0.579934 -0.057408 11.00000 0.04164 0.05011 =
0.03465 -0.00191 0.00207 0.00546

C11 1 0.291866 0.566535 0.015789 11.00000 0.04949 0.04829 =
0.04010 -0.00015 0.00315 0.00327

AFIX 43

H11 2 0.343968 0.619554 0.040893 11.00000 -1.20000

AFIX 0

C12 1 0.379533 0.460351 0.125034 11.00000 0.04356 0.04903 =
0.03774 0.00568 0.00510 0.00730

C13 1 0.462201 0.528157 0.172658 11.00000 0.07304 0.04085 =
0.04574 0.00979 -0.00401 -0.00278

AFIX 43

H13 2 0.467179 0.592349 0.156691 11.00000 -1.20000

AFIX 0

C14 1 0.537379 0.500798 0.243863 11.00000 0.07131 0.04609 =
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AFIX 43

H14 2 0.594324 0.546593 0.275698 11.00000 -1.20000

AFIX 0

C15 1 0.529174 0.406454 0.268431 11.00000 0.04883 0.04457 =
0.03357 0.00105 0.00473 0.00795

C16 1 0.447293 0.338205 0.221093 11.00000 0.06591 0.03984 =
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AFIX 43

H16 2 0.442650 0.274010 0.237083 11.00000 -1.20000

AFIX 0

C17 1 0.372273 0.365881 0.149795 11.00000 0.05983 0.04701 =
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AFIX 43

H17 2 0.315809 0.319999 0.117875 11.00000 -1.20000

AFIX 0

C18 1 0.371849 0.381591 0.444147 11.00000 0.05738 0.03751 =
0.03402 0.00235 -0.00002 0.00302

C19 1 0.110644 0.296188 0.474215 11.00000 0.05894 0.06214 =
0.08122 -0.00098 0.00802 -0.01077

AFIX 43

H19 2 0.016556 0.255785 0.477583 11.00000 -1.20000

AFIX 0

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0.06277 0.00052 0.01600 0.00118

AFIX 43

H20 2 0.064756 0.396229 0.553157 11.00000 -1.20000

AFIX 0

HKLF 4 1 1 0 0 0 1 0 0 0 1

REM NAPSULY_0m in P2(1)/c RED PRISM

REM wR2 = 0.1087, GooF = S = 1.018, Restrained GooF = 1.018 for all data

REM R1 = 0.0437 for 2548 Fo > 4sig(Fo) and 0.0852 for all 4040 data

REM 259 parameters refined using 0 restraints

END

CIF of ACPIN without hkl data

data_global

#=====

PROCESSING SUMMARY (IUCr Office Use only)

_publ_contact_author

;

Muhammad Nawaz Tahir
University of Sargodha
Department of Physics
Sargodha
Pakistan

;

_publ_contact_author_phone '0092 48 92 30 914'
_publ_contact_author_fax '0092 48 32 22 121'
_publ_contact_author_email 'dmntahir_uos@yahoo.com'
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'Muhammad Ashfaq'
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Av. Santa Maria 6400 Vitacura,
Postal code: 7660251,
Chile

;

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TEXT

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(E)-1-(((4-amino-5-(4-chlorophenyl)-6-ethylpyrimidin-2-yl)iminio)methyl)

naphthalen-2-olate

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_publ_section_references

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Bruker (2007). <i>SADABS</i>. Bruker AXS Inc., Madison, Wisconsin, USA.

Bruker (2007). <i>APEX2</i> and <i>SAINT</i>. Bruker AXS Inc.,
Madison, Wisconsin, USA.

Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.

Farrugia, L. J. (2012). *J. Appl. Cryst.* **45**, 849--854.

Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112--122.

Sheldrick, G. M. (2015). *Acta Cryst.* **C71**, 3--8.

Spek, A. L. (2009). *Acta Cryst.* **D65**, 148--155.

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naphthalen-2-olate

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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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  '-x, -y, -z'

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  _cell_length_c          17.673(2)
  _cell_angle_alpha       88.308(8)
  _cell_angle_beta        89.195(8)
  _cell_angle_gamma       82.611(9)
  _cell_volume            996.9(2)
  _cell_formula_units_Z   2
  _cell_measurement_reflns_used    2397
  _cell_measurement_theta_min      1.640
  _cell_measurement_theta_max      27.448
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  _exptl_crystal_size_min      0.20
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  _exptl_crystal_density_meas     ?
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  _exptl_crystal_F_000          420
  _exptl_absorpt_coefficient_mu    0.214
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  _exptl_absorpt_process_details   '(SADABS; Bruker, 2007)'
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Refinement of $\langle F \rangle^2$ against ALL reflections. The weighted $\langle R \rangle$ -factor $\langle wR \rangle$ and goodness of fit $\langle S \rangle$ are based on $\langle F \rangle^2$, conventional $\langle R \rangle$ -factors $\langle R \rangle$ are based on $\langle F \rangle$, with $\langle F \rangle$ set to zero for negative $\langle F \rangle^2$. The threshold expression of $\langle F \rangle^2 > \sqrt{\langle F \rangle^2}$ is used only for calculating $\langle R \rangle$ -factors(gt) etc. and is not relevant to the choice of reflections for refinement.

$\langle R \rangle$ -factors based on $\langle F \rangle^2$ are statistically about twice as large as those based on $\langle F \rangle$, and $\langle R \rangle$ -factors based on ALL data will be even larger.

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'w=1/[\(\langle F \rangle^2 + (0.0757P)^2 + 0.7915P\)] where $P = (F_o^2 + 2F_c^2)/3$ '


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O1 O 1.5052(8) 0.6311(2) 0.22359(17) 0.0647(9) Uani 1 1 d
N1 N 1.1691(8) 0.7121(2) 0.11607(17) 0.0506(9) Uani 1 1 d
H1 H 1.316681 0.667310 0.131723 0.061 Uiso 1 1 calc R U . . .
N2 N 1.2336(8) 0.6206(3) 0.00662(17) 0.0515(9) Uani 1 1 d
N3 N 1.3058(9) 0.5222(3) -0.09936(18) 0.0620(10) Uani 1 1 d
H3A H 1.422638 0.476107 -0.073272 0.074 Uiso 1 1 calc R U . . .
H3B H 1.275403 0.511139 -0.146252 0.074 Uiso 1 1 calc R U . . .
N4 N 0.8944(8) 0.7821(2) 0.01133(18) 0.0490(8) Uani 1 1 d
C1 C 1.3595(10) 0.6975(3) 0.2669(2) 0.0519(10) Uani 1 1 d
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C3 C 1.2799(11) 0.7618(4) 0.3950(2) 0.0606(12) Uani 1 1 d
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C4 C 1.0470(10) 0.8437(3) 0.3713(2) 0.0530(11) Uani 1 1 d
C5 C 0.8982(12) 0.9153(4) 0.4231(2) 0.0629(13) Uani 1 1 d
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C6 C 0.6814(12) 0.9948(4) 0.4005(3) 0.0683(13) Uani 1 1 d
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C7 C 0.6019(11) 1.0036(4) 0.3247(2) 0.0625(12) Uani 1 1 d
H7 H 0.451165 1.056642 0.308993 0.075 Uiso 1 1 calc R U . . .
C8 C 0.7432(10) 0.9353(3) 0.2731(2) 0.0554(11) Uani 1 1 d
H8 H 0.687906 0.943491 0.222608 0.067 Uiso 1 1 calc R U . . .
C9 C 0.9688(10) 0.8531(3) 0.2940(2) 0.0455(9) Uani 1 1 d
C10 C 1.1248(9) 0.7779(3) 0.2408(2) 0.0441(9) Uani 1 1 d
C11 C 1.0389(10) 0.7792(3) 0.1662(2) 0.0479(10) Uani 1 1 d
H11 H 0.881995 0.829698 0.150473 0.057 Uiso 1 1 calc R U . . .
C12 C 1.0917(10) 0.7062(3) 0.0402(2) 0.0459(9) Uani 1 1 d
C13 C 1.1704(10) 0.6115(3) -0.0668(2) 0.0488(10) Uani 1 1 d
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C15 C 0.8397(9) 0.7750(3) -0.0636(2) 0.0448(9) Uani 1 1 d
C16 C 0.6198(10) 0.8645(3) -0.0940(2) 0.0551(11) Uani 1 1 d

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H16B H 0.571511 0.848972 -0.145513 0.066 Uiso 1 1 calc R U . . .
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C18 C 0.9410(9) 0.6853(3) -0.1905(2) 0.0461(9) Uani 1 1 d
C19 C 1.0302(11) 0.7619(3) -0.2399(2) 0.0570(11) Uani 1 1 d
H19 H 1.112091 0.819797 -0.220837 0.068 Uiso 1 1 calc R U . . .
C20 C 1.0012(12) 0.7551(4) -0.3177(2) 0.0683(13) Uani 1 1 d
H20 H 1.061097 0.807895 -0.350439 0.082 Uiso 1 1 calc R U . . .
C21 C 0.8825(12) 0.6690(4) -0.3453(2) 0.0665(13) Uani 1 1 d
C22 C 0.7958(13) 0.5910(4) -0.2986(3) 0.0749(15) Uani 1 1 d
H22 H 0.716106 0.532999 -0.318188 0.090 Uiso 1 1 calc R U . . .
C23 C 0.8277(11) 0.5989(4) -0.2210(2) 0.0635(13) Uani 1 1 d
H23 H 0.771632 0.544828 -0.188875 0.076 Uiso 1 1 calc R U . . .

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O1 0.076(2) 0.0589(18) 0.0561(19) -0.0029(14) -0.0072(16) 0.0057(16)
N1 0.066(2) 0.0479(18) 0.0349(17) -0.0006(14) -0.0026(15) 0.0031(16)
N2 0.070(2) 0.0467(19) 0.0353(17) 0.0002(14) 0.0037(16) 0.0028(17)
N3 0.086(3) 0.058(2) 0.0346(18) -0.0019(15) 0.0001(17) 0.0190(19)
N4 0.058(2) 0.0459(18) 0.0418(18) -0.0051(14) -0.0021(15) 0.0005(16)
C1 0.069(3) 0.049(2) 0.039(2) 0.0005(18) -0.005(2) -0.012(2)
C2 0.072(3) 0.062(3) 0.047(2) 0.007(2) -0.012(2) -0.004(2)

C3 0.079(3) 0.069(3) 0.036(2) 0.004(2) -0.018(2) -0.020(3)
C4 0.067(3) 0.058(2) 0.038(2) -0.0038(18) 0.0008(19) -0.024(2)
C5 0.086(4) 0.074(3) 0.034(2) -0.009(2) 0.001(2) -0.026(3)
C6 0.082(4) 0.071(3) 0.053(3) -0.019(2) 0.015(2) -0.014(3)
C7 0.069(3) 0.065(3) 0.051(3) -0.008(2) 0.005(2) -0.002(2)
C8 0.065(3) 0.061(3) 0.041(2) -0.0016(19) 0.000(2) -0.008(2)
C9 0.056(3) 0.049(2) 0.0337(19) -0.0009(16) 0.0001(17) -0.0170(19)
C10 0.055(3) 0.043(2) 0.0352(19) 0.0019(15) -0.0002(17) -0.0104(18)
C11 0.063(3) 0.040(2) 0.040(2) 0.0009(16) -0.0051(19) -0.0041(18)
C12 0.061(3) 0.045(2) 0.0326(19) 0.0024(16) -0.0010(17) -0.0079(19)
C13 0.063(3) 0.046(2) 0.036(2) -0.0004(16) 0.0069(18) -0.0004(19)
C14 0.051(3) 0.044(2) 0.0357(19) 0.0006(16) -0.0001(17) -0.0093(18)
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C16 0.056(3) 0.060(3) 0.046(2) -0.0112(19) -0.0108(19) 0.006(2)
C17 0.088(4) 0.052(3) 0.085(4) 0.002(2) -0.032(3) 0.007(2)
C18 0.053(3) 0.045(2) 0.039(2) 0.0001(16) -0.0007(17) -0.0042(18)
C19 0.073(3) 0.057(3) 0.044(2) 0.0058(19) -0.009(2) -0.016(2)
C20 0.092(4) 0.069(3) 0.046(2) 0.015(2) -0.004(2) -0.021(3)
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C22 0.107(5) 0.079(3) 0.045(3) -0.009(2) -0.006(3) -0.035(3)
C23 0.087(4) 0.063(3) 0.045(2) 0.002(2) 0.004(2) -0.028(2)

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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.

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O1 C1 1.263(5) . ?
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N1 C12 1.397(5) . ?
N1 H1 0.8600 . ?
N2 C12 1.330(5) . ?
N2 C13 1.344(5) . ?
N3 C13 1.346(5) . ?
N3 H3A 0.8600 . ?
N3 H3B 0.8600 . ?
N4 C12 1.316(5) . ?
N4 C15 1.358(5) . ?
C1 C10 1.440(6) . ?
C1 C2 1.458(5) . ?
C2 C3 1.332(6) . ?
C2 H2 0.9300 . ?
C3 C4 1.433(6) . ?
C3 H3 0.9300 . ?
C4 C5 1.404(6) . ?
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C5 C6 1.362(7) . ?
C5 H5 0.9300 . ?
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C6 H6 0.9300 . ?
C7 C8 1.366(6) . ?
C7 H7 0.9300 . ?
C8 C9 1.401(6) . ?
C8 H8 0.9300 . ?

C9 C10 1.461(5) . ?
C10 C11 1.379(5) . ?
C11 H11 0.9300 . ?
C13 C14 1.419(5) . ?
C14 C15 1.387(5) . ?
C14 C18 1.501(5) . ?
C15 C16 1.495(5) . ?
C16 C17 1.506(6) . ?
C16 H16A 0.9700 . ?
C16 H16B 0.9700 . ?
C17 H17A 0.9600 . ?
C17 H17B 0.9600 . ?
C17 H17C 0.9600 . ?
C18 C19 1.375(5) . ?
C18 C23 1.383(6) . ?
C19 C20 1.389(6) . ?
C19 H19 0.9300 . ?
C20 C21 1.371(6) . ?
C20 H20 0.9300 . ?
C21 C22 1.354(6) . ?
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C22 H22 0.9300 . ?
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C11 N1 H1 116.7 .. ?
C12 N1 H1 116.7 .. ?
C12 N2 C13 115.6(3) .. ?
C13 N3 H3A 120.0 .. ?
C13 N3 H3B 120.0 .. ?
H3A N3 H3B 120.0 .. ?
C12 N4 C15 115.8(3) .. ?
O1 C1 C10 123.1(3) .. ?
O1 C1 C2 119.3(4) .. ?
C10 C1 C2 117.6(4) .. ?
C3 C2 C1 121.3(4) .. ?
C3 C2 H2 119.3 .. ?
C1 C2 H2 119.3 .. ?
C2 C3 C4 123.0(4) .. ?
C2 C3 H3 118.5 .. ?
C4 C3 H3 118.5 .. ?
C5 C4 C9 119.5(4) .. ?
C5 C4 C3 121.4(4) .. ?
C9 C4 C3 119.1(4) .. ?
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C6 C5 H5 119.2 .. ?
C4 C5 H5 119.2 .. ?
C5 C6 C7 119.1(4) .. ?
C5 C6 H6 120.5 .. ?
C7 C6 H6 120.5 .. ?
C8 C7 C6 120.6(5) .. ?
C8 C7 H7 119.7 .. ?
C6 C7 H7 119.7 .. ?
C7 C8 C9 122.0(4) .. ?
C7 C8 H8 119.0 .. ?
C9 C8 H8 119.0 .. ?
C8 C9 C4 117.2(4) .. ?
C8 C9 C10 123.9(3) .. ?

C4 C9 C10 119.0(4) . . ?
C11 C10 C1 119.3(4) . . ?
C11 C10 C9 120.5(4) . . ?
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N1 C11 C10 123.0(4) . . ?
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C10 C11 H11 118.5 . . ?
N4 C12 N2 128.5(3) . . ?
N4 C12 N1 118.6(3) . . ?
N2 C12 N1 113.0(3) . . ?
N2 C13 N3 115.0(4) . . ?
N2 C13 C14 121.8(3) . . ?
N3 C13 C14 123.2(3) . . ?
C15 C14 C13 116.1(3) . . ?
C15 C14 C18 123.7(3) . . ?
C13 C14 C18 120.2(3) . . ?
N4 C15 C14 122.1(4) . . ?
N4 C15 C16 113.6(3) . . ?
C14 C15 C16 124.3(3) . . ?
C15 C16 C17 113.0(4) . . ?
C15 C16 H16A 109.0 . . ?
C17 C16 H16A 109.0 . . ?
C15 C16 H16B 109.0 . . ?
C17 C16 H16B 109.0 . . ?
H16A C16 H16B 107.8 . . ?
C16 C17 H17A 109.5 . . ?
C16 C17 H17B 109.5 . . ?
H17A C17 H17B 109.5 . . ?
C16 C17 H17C 109.5 . . ?
H17A C17 H17C 109.5 . . ?
H17B C17 H17C 109.5 . . ?
C19 C18 C23 117.5(4) . . ?
C19 C18 C14 121.6(4) . . ?

C23 C18 C14 120.8(3) . . ?
 C18 C19 C20 121.7(4) . . ?
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 C20 C19 H19 119.2 . . ?
 C21 C20 C19 118.7(4) . . ?
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 C22 C21 C20 121.5(4) . . ?
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 C20 C21 Cl1 118.8(4) . . ?
 C21 C22 C23 119.0(4) . . ?
 C21 C22 H22 120.5 . . ?
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 N3 H3A N2 0.86 2.19 3.037(5) 170.8 2_865 yes
 N3 H3B O1 0.86 2.38 3.003(5) 129.2 2_865 yes

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TITL PYMTVAN_0m in P-1 YELLOW NEEDLE

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created by SHELXL-2019/2 at 14:15:37 on 18-Feb-2023

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ZERR 2.00 0.0006 0.0016 0.0022 0.008 0.008 0.009

LATT 1

SFAC C H N O Cl

UNIT 46 38 8 2 2

L.S. 20

ACTA

LIST 4

BOND \$H

HTAB

HTAB N1 O1

EQIV \$1 -x+3, -y+1, -z

HTAB N3 N2_\$1

HTAB N3 O1_\$1

FMAP 2

PLAN 10

SIZE 0.40 0.24 0.20

TEMP 23.000

MPLA 9 C12 N2 C13 C14 C15 N4 C16 C17 N3

MPLA 7 C18 C19 C20 C21 C22 C23 Cl1

MPLA 11 C1 C2 C3 C4 C5 C6 C7 C8 C9 C10 C11 C12 O1

MPLA 9 C12 N2 C13 C14 C15 N4 C16 C17 N3

WGHT 0.075700 0.791500

BASF 0.18748

FVAR 10.25644

CL1 5 0.843720 0.660723 -0.442590 11.00000 0.18910 0.15579 =
0.03694 -0.00168 -0.01765 -0.05358

O1 4 1.505207 0.631141 0.223588 11.00000 0.07565 0.05889 =
0.05611 -0.00290 -0.00718 0.00569

N1 3 1.169096 0.712051 0.116073 11.00000 0.06622 0.04789 =
0.03492 -0.00057 -0.00264 0.00309

AFIX 43

H1 2 1.316681 0.667310 0.131723 11.00000 -1.20000

AFIX 0

N2 3 1.233611 0.620577 0.006622 11.00000 0.06953 0.04674 =
0.03526 0.00020 0.00371 0.00276

N3 3 1.305780 0.522156 -0.099363 11.00000 0.08581 0.05811 =
0.03457 -0.00187 0.00007 0.01899

AFIX 93

H3A 2 1.422638 0.476107 -0.073272 11.00000 -1.20000

H3B 2 1.275403 0.511139 -0.146252 11.00000 -1.20000

AFIX 0

N4 3 0.894449 0.782108 0.011335 11.00000 0.05776 0.04586 =
0.04184 -0.00506 -0.00208 0.00046

C1 1 1.359490 0.697474 0.266882 11.00000 0.06921 0.04862 =
0.03885 0.00054 -0.00460 -0.01164

C2 1 1.427473 0.693255 0.347355 11.00000 0.07188 0.06156 =
0.04700 0.00682 -0.01196 -0.00411

AFIX 43

H2 2 1.576803 0.641685 0.365800 11.00000 -1.20000

AFIX 0

C3 1 1.279931 0.761810 0.395009 11.00000 0.07914 0.06919 =
0.03637 0.00406 -0.01830 -0.02048

AFIX 43

H3 2 1.330356 0.756311 0.445923 11.00000 -1.20000

AFIX 0

C4 1 1.047010 0.843650 0.371347 11.00000 0.06659 0.05842 =
0.03825 -0.00378 0.00076 -0.02433

C5 1 0.898223 0.915265 0.423055 11.00000 0.08559 0.07381 =
0.03386 -0.00880 0.00065 -0.02634

AFIX 43

H5 2 0.948896 0.908272 0.473926 11.00000 -1.20000

AFIX 0

C6 1 0.681392 0.994754 0.400479 11.00000 0.08208 0.07121 =
0.05332 -0.01888 0.01469 -0.01382

AFIX 43

H6 2 0.587866 1.042410 0.435310 11.00000 -1.20000

AFIX 0

C7 1 0.601943 1.003635 0.324724 11.00000 0.06935 0.06545 =
0.05101 -0.00796 0.00462 -0.00178

AFIX 43

H7 2 0.451165 1.056642 0.308993 11.00000 -1.20000

AFIX 0

C8 1 0.743179 0.935250 0.273123 11.00000 0.06461 0.06087 =
0.04087 -0.00159 0.00019 -0.00830

AFIX 43

H8 2 0.687906 0.943491 0.222608 11.00000 -1.20000

AFIX 0

C9 1 0.968758 0.853075 0.293964 11.00000 0.05619 0.04942 =
0.03367 -0.00095 0.00007 -0.01702

C10 1 1.124755 0.777884 0.240784 11.00000 0.05516 0.04292 =
0.03515 0.00189 -0.00020 -0.01039

C11 1 1.038895 0.779186 0.166198 11.00000 0.06320 0.04007 =
0.03971 0.00086 -0.00515 -0.00412

AFIX 43

H11 2 0.881995 0.829698 0.150473 11.00000 -1.20000

AFIX 0

C12 1 1.091721 0.706242 0.040238 11.00000 0.06054 0.04476 =
0.03257 0.00237 -0.00097 -0.00791

C13 1 1.170420 0.611475 -0.066810 11.00000 0.06281 0.04575 =
0.03590 -0.00042 0.00692 -0.00035

C14 1 0.978035 0.691656 -0.106490 11.00000 0.05104 0.04426 =
0.03574 0.00055 -0.00008 -0.00927

C15 1 0.839686 0.774960 -0.063588 11.00000 0.05277 0.04665 =
0.03496 -0.00239 -0.00522 -0.00602

C16 1 0.619837 0.864460 -0.094022 11.00000 0.05594 0.06020 =
0.04648 -0.01122 -0.01085 0.00573

AFIX 23

H16A 2 0.438920 0.867660 -0.063915 11.00000 -1.20000

H16B 2 0.571511 0.848972 -0.145513 11.00000 -1.20000

AFIX 0

C17 1 0.731879 0.972490 -0.093568 11.00000 0.08795 0.05179 =
0.08462 0.00233 -0.03228 0.00698

AFIX 137

H17A 2 0.808426 0.983545 -0.044450 11.00000 -1.50000

H17B 2 0.571904 1.028182 -0.105070 11.00000 -1.50000

H17C 2 0.886726 0.974607 -0.130896 11.00000 -1.50000

AFIX 0

C18 1 0.940957 0.685305 -0.190453 11.00000 0.05336 0.04536 =
0.03893 0.00012 -0.00072 -0.00422

C19 1 1.030220 0.761935 -0.239937 11.00000 0.07253 0.05666 =
0.04353 0.00583 -0.00932 -0.01588

AFIX 43

H19 2 1.112091 0.819797 -0.220837 11.00000 -1.20000

AFIX 0

C20 1 1.001206 0.755110 -0.317715 11.00000 0.09196 0.06916 =
0.04561 0.01483 -0.00366 -0.02119

AFIX 43

H20 2 1.061097 0.807895 -0.350439 11.00000 -1.20000

AFIX 0

C21 1 0.882531 0.669017 -0.345280 11.00000 0.08426 0.08161 =
0.03437 -0.00273 -0.00423 -0.01290

C22 1 0.795820 0.590999 -0.298596 11.00000 0.10743 0.07921 =
0.04463 -0.00893 -0.00591 -0.03502

AFIX 43

H22 2 0.716106 0.532999 -0.318188 11.00000 -1.20000

AFIX 0

C23 1 0.827735 0.598919 -0.221049 11.00000 0.08686 0.06283 =
0.04518 0.00211 0.00356 -0.02785

AFIX 43

H23 2 0.771632 0.544828 -0.188875 11.00000 -1.20000

AFIX 0

HKLF 5

REM PYMTVAN_0m in P-1 YELLOW NEEDLE

REM wR2 = 0.2358, GooF = S = 1.139, Restrained GooF = 1.139 for all data

REM R1 = 0.0763 for 2397 Fo > 4sig(Fo) and 0.1311 for all 4351 data

REM 264 parameters refined using 0 restraints

END

Table S5: Selected structural parameters for the compounds **DSPIN** and **ACPIN**, calculated using the B3LYP/6-311+G(d,p) approach with the implicit solvent effects from ethanol.

Selected distances (Å) in DSPIN		Selected distances (Å) in ACPIN	
H11—H13	2.070	H11—N4	2.394
H11—N1	2.055	H1—N2	2.468
H13—N1	2.744	H3—N2	2.456
N2—H3	2.534	-	-
Selected dihedral angles (°) in DSPIN		Selected dihedral angles (°) in ACPIN	
N1—C11—C10—C1	-0.566	N1—C11—C10—C1	-0.042
C11—C10—C1—O1	0.658	C11—C10—C1—O1	0.257
H1—N1—C1—O1	-1.358	H1—N1—C1—O1	0.102
H13—C13—C11—H1	-13.103	H11—C11—C12—N4	-0.139
O3—S1—C18—S2	-2.349	C15—C14—C18—C19	75.940
C10—C11—N1—C12	179.224	C10—C11—N1—C12	179.570

Table S6. Second-order perturbation theory analysis results for the compounds **DSPIN** and **ACPIN**, calculated using the B3LYP/6-311+G(d,p) approach with the implicit solvent effects from ethanol.

Compound	Donor	Type	Acceptor	Type	E(2), ^a kcal/mol	E(j)-E(i), A.U.	F(i,j), A.U.
DSPIN	S1-C15	σ	C13-C14	π^*	11.88	0.05	0.024
	S2-C18	σ	C13-C14	π^*	73.93	0.03	0.048
	S2-C19	σ	C13-C14	π^*	488.92	0.01	0.078
	O1-C1	π	C10	LP*	16.04	0.09	0.052
	N1-C11	σ	C13-C14	π^*	45.75	0.23	0.100
	N1-C11	σ	C16-H16	σ^*	721.29	0.03	0.137
	N1-C11	σ	C16-C17	σ^*	10.00	0.84	0.082
	N1-C11	σ	C10	LP*	67.16	0.11	0.113
	N1-C11	π	O1-C1	π^*	16.96	0.22	0.059
	N1-C11	π	N1-C11	π^*	11.58	0.16	0.043
	N1-C11	π	N3-C18	σ^*	11.91	0.53	0.072
	N1-C11	π	C3-C4	σ^*	14.40	0.52	0.079
	N1-C11	π	C8-H8	σ^*	19.88	0.42	0.084
	N1-C11	π	C12-C17	π^*	28.92	0.36	0.098
	N1-C11	π	C14-H14	σ^*	11.58	0.54	0.072
	N1-C11	π	C14-C15	σ^*	14.94	0.66	0.090
	N1-C11	π	C16-C17	σ^*	39.19	0.34	0.106
	N1-C11	π	C19-C20	σ^*	14.83	1.65	0.143
	N1-C12	σ	O1-C1	σ^*	12.76	1.16	0.109
	N1-C12	σ	O1-C1	π^*	11.77	0.72	0.091
	N1-C12	σ	N1-C11	π^*	12.79	0.66	0.094
	N1-C12	σ	N3-H3	σ^*	14.54	1.03	0.110
	N1-C12	σ	N3-C18	σ^*	18.32	1.03	0.124
	N1-C12	σ	N3-C20	σ^*	11.80	1.09	0.101
	N1-C12	σ	C3-C4	σ^*	22.97	1.02	0.137
	N1-C12	σ	C5-H5	σ^*	11.25	1.13	0.101
	N1-C12	σ	C8-H8	σ^*	29.43	0.92	0.147

	N1-C12	σ	C13-C14	σ^*	232.83	0.23	0.225
	N1-C12	σ	C14-H14	σ^*	18.62	1.04	0.124
	N1-C12	σ	C14-C15	σ^*	27.12	1.15	0.158
	N1-C12	σ	C16-H16	σ^*	3398.88	0.03	0.297
	N1-C12	σ	C16-C17	σ^*	72.22	0.84	0.220
	N1-C12	σ	C19-C20	σ^*	33.24	2.15	0.239
	N3-H3	σ	O1-C1	π^*	11.45	0.54	0.078
	N3-H3	σ	N3-C18	σ^*	15.56	0.85	0.103
	N3-H3	σ	C7-H7	σ^*	20.49	1.03	0.130
	N3-H3	σ	C13-C14	π^*	411.36	0.06	0.145
	N3-C18	σ	C14-C15	σ^*	13.79	1.15	0.113
	N3-C18	σ	C16-H16	σ^*	1845.82	0.03	0.208
	N3-C18	σ	C16-C17	σ^*	11.91	0.84	0.089
	N3-C18	σ	C19-C20	σ^*	14.53	2.15	0.158
	N3-C20	σ	C10	LP*	58.17	0.73	0.279
	N3-C20	σ	S1-N2	σ^*	12.45	1.09	0.109
	N3-C20	σ	O1-C1	σ^*	64.05	1.28	0.256
	N3-C20	σ	O1-C1	π^*	87.73	0.84	0.269
	N3-C20	σ	N1-C10	π^*	51.63	0.78	0.205
	N3-C20	σ	N1-C12	σ^*	17.15	1.35	0.136
	N3-C20	σ	N2-C18	σ^*	50.01	1.40	0.236
	N3-C20	σ	N3-H3	σ^*	83.44	1.15	0.277
	N3-C20	σ	N3-C18	σ^*	151.39	1.15	0.375
	N3-C20	σ	N3-C20	σ^*	36.47	1.21	0.187
	N3-C20	σ	C1-C10	σ^*	12.80	1.40	0.121
	N3-C20	σ	C2-H2	σ^*	18.56	1.36	0.142
	N3-C20	σ	C3-C4	σ^*	117.54	1.14	0.327
	N3-C20	σ	C4-C9	σ^*	19.73	1.43	0.151
	N3-C20	σ	C5-H5	σ^*	45.14	1.25	0.212
	N3-C20	σ	C6-H6	σ^*	11.61	1.38	0.113
	N3-C20	σ	C6-C7	σ^*	30.48	1.46	0.188
	N3-C20	σ	C7-H7	σ^*	108.06	1.32	0.338

	N3-C20	σ	C8-H8	σ^*	131.77	1.04	0.330
	N3-C20	σ	C9-C10	σ^*	22.32	1.42	0.160
	N3-C20	σ	C11-H11	σ^*	24.91	1.28	0.159
	N3-C20	σ	C13-C14	π^*	969.38	0.35	0.563
	N3-C20	σ	C14-H14	σ^*	83.01	1.16	0.277
	N3-C20	σ	C14-C15	σ^*	78.10	1.27	0.282
	N3-C20	σ	C15-C16	π^*	12.92	0.96	0.110
	N3-C20	σ	C16-H16	σ^*	2881.41	0.15	0.590
	N3-C20	σ	C16-C17	σ^*	196.69	0.96	0.388
	N3-C20	σ	C19-C20	σ^*	207.91	2.27	0.613
	N3-C20	σ	C19-C20	π^*	13.27	2.57	0.174
	C1-C2	σ	O1-C1	σ^*	24.75	1.01	0.141
	C1-C2	σ	O1-C1	π^*	13.65	0.56	0.087
	C1-C2	σ	N1-C10	π^*	16.22	0.50	0.092
	C1-C2	σ	N3-H3	σ^*	16.27	0.88	0.107
	C1-C2	σ	N3-C18	σ^*	19.40	0.87	0.117
	C1-C2	σ	N3-C20	σ^*	12.29	0.93	0.096
	C1-C2	σ	C1-C10	σ^*	10.15	1.13	0.096
	C1-C2	σ	C3-C4	σ^*	24.50	0.86	0.130
	C1-C2	σ	C5-H5	σ^*	12.33	0.97	0.098
	C1-C2	σ	C8-H8	σ^*	31.78	0.76	0.139
	C1-C2	σ	C13-C14	π^*	652.58	0.08	0.214
	C1-C2	σ	C14-H14	σ^*	20.55	0.88	0.120
	C1-C2	σ	C14-C15	σ^*	27.87	1.00	0.149
	C1-C2	σ	C16-C17	σ^*	55.81	0.68	0.175
	C1-C2	σ	C19-C20	σ^*	32.91	1.99	0.229
	C2-C3	π	C10	LP*	27.40	0.16	0.078
	C2-C3	π	O1-C1	π^*	86.00	0.11	0.093
	C3-C4	σ	C13-C14	π^*	96.51	0.03	0.050
	C5-C6	π	C10	LP*	41.86	0.15	0.088
	C5-C6	π	C7-C8	π^*	20.55	0.28	0.068
	C7-C8	π	C9	LP*	19.68	0.35	0.091

	C7-C8	π	C10	LP*	130.61	0.19	0.185
	C7-C8	π	S1-N2	σ^*	38.33	0.55	0.133
	C7-C8	π	O1-C1	σ^*	70.65	0.74	0.221
	C7-C8	π	O1-C1	π^*	131.51	0.30	0.182
	C7-C8	π	N1-C11	π^*	168.05	0.24	0.188
	C7-C8	π	N1-C12	σ^*	14.71	0.81	0.105
	C7-C8	π	N2-C18	σ^*	35.24	0.86	0.167
	C7-C8	π	N3-H3	σ^*	76.43	0.61	0.208
	C7-C8	π	N3-C18	σ^*	113.45	0.61	0.252
	C7-C8	π	N3-C20	σ^*	64.25	0.67	0.200
	C7-C8	π	C1-C10	σ^*	18.61	0.87	0.121
	C7-C8	π	C2-H2	σ^*	25.20	0.82	0.139
	C7-C8	π	C3-C4	σ^*	130.11	0.60	0.268
	C7-C8	π	C4-C9	σ^*	23.14	0.89	0.138
	C7-C8	π	C5-H5	σ^*	59.27	0.71	0.198
	C7-C8	π	C5-C6	σ^*	12.10	0.49	0.069
	C7-C8	π	C6-H6	σ^*	14.18	0.84	0.105
	C7-C8	π	C6-C7	σ^*	25.16	0.92	0.146
	C7-C8	π	C7-H7	σ^*	39.74	0.78	0.170
	C7-C8	π	C8-H8	σ^*	170.45	0.50	0.281
	C7-C8	π	C9-C10	σ^*	17.31	0.88	0.119
	C7-C8	π	C11-H11	σ^*	36.39	0.74	0.158
	C7-C8	π	C12-C13	σ^*	10.35	0.96	0.096
	C7-C8	π	C12-C17	π^*	16.40	0.44	0.077
	C7-C8	π	C14-H14	σ^*	109.14	0.62	0.250
	C7-C8	π	C14-C15	σ^*	145.20	0.73	0.314
	C7-C8	π	C15-C16	π^*	35.76	0.42	0.112
	C7-C8	π	C16-C17	σ^*	311.99	0.42	0.349
	C7-C8	π	C19-C20	σ^*	168.27	1.73	0.520
	C7-C8	π	C19-C20	π^*	41.34	2.03	0.264
	C8-H8	σ	C4-C9	σ^*	11.67	0.97	0.095
	C8-H8	σ	C6-C7	σ^*	12.28	0.99	0.099

	C8-H8	σ	C8-H8	σ^*	16.39	0.58	0.087
	C8-H8	σ	C16-C17	σ^*	24.59	0.50	0.099
	C8-C9	σ	C13-C14	π^*	231.30	0.06	0.113
	C8-C9	σ	C16-C17	σ^*	16.68	0.67	0.094
	C8-C9	σ	C19-C20	σ^*	12.61	1.98	0.141
	C12-C13	σ	C13-C14	π^*	19.77	0.08	0.038
	C12-C17	σ	O1-C1	σ^*	11.17	1.04	0.097
	C12-C17	σ	O1-C1	π^*	10.20	0.60	0.077
	C12-C17	σ	N1-C11	σ^*	12.24	0.54	0.083
	C12-C17	σ	N3-H3	σ^*	11.74	0.91	0.092
	C12-C17	σ	N3-C18	σ^*	15.15	0.91	0.105
	C12-C17	σ	C3-C4	σ^*	18.64	0.90	0.115
	C12-C17	σ	C8-H8	σ^*	23.39	0.80	0.122
	C12-C17	σ	C13-C14	π^*	333.15	0.11	0.184
	C12-C17	σ	C14-H14	σ^*	15.18	0.92	0.106
	C12-C17	σ	C14-C15	σ^*	20.97	1.03	0.131
	C12-C17	σ	C16-C17	σ^*	72.89	0.72	0.205
	C12-C17	σ	C19-C20	σ^*	32.78	2.03	0.231
	C12-C17	π	C10	LP*	82.01	0.01	0.038
	C12-C17	π	O1-C1	π^*	16.42	0.12	0.040
	C12-C17	π	C8-H8	σ^*	12.14	0.32	0.062
	C12-C17	π	C15-C16	π^*	12.95	0.24	0.050
	C12-C17	π	C16-C17	σ^*	24.11	0.24	0.076
	C13-C14	π	C12-C17	π^*	24.51	0.25	0.071
	C13-C14	π	C15-C16	π^*	21.09	0.23	0.063
	C14-C15	σ	C8-H8	σ^*	11.40	0.80	0.085
	C14-C15	σ	C13-C14	π^*	158.26	0.11	0.126
	C14-C15	σ	C14-H14	σ^*	16.31	0.91	0.109
	C14-C15	σ	C14-C15	σ^*	10.33	1.03	0.092
	C14-C15	σ	C16-C17	σ^*	20.20	0.72	0.108
	C14-C15	σ	C19-C20	σ^*	16.24	2.03	0.162
	C15-C16	π	C12-C17	π^*	16.49	0.26	0.059

	C19-H19	σ	C10	LP*	17.35	0.33	0.102
	C19-H19	σ	O1-C1	π^*	21.29	0.44	0.096
	C19-H19	σ	N3-C18	σ^*	35.50	0.75	0.147
	C19-H19	σ	C3-C4	σ^*	18.19	0.74	0.104
	C19-H19	σ	C7-H7	σ^*	24.00	0.93	0.133
	C19-H19	σ	C8-H8	σ^*	20.22	0.64	0.102
	C19-H19	σ	C14-H14	σ^*	10.37	0.76	0.079
	C19-H19	σ	C16-C17	σ^*	29.30	0.56	0.115
	C19-H19	σ	C10	LP*	32.14	0.41	0.156
	C19-C20	σ	O1-C1	σ^*	14.12	0.97	0.105
	C19-C20	σ	O1-C1	π^*	0.97	0.52	0.146
	C19-C20	σ	N1-C11	π^*	12.50	0.46	0.078
	C19-C20	σ	N3-H3	σ^*	37.62	0.84	0.159
	C19-C20	σ	N3-C18	σ^*	83.30	0.84	0.237
	C19-C20	σ	C3-C4	σ^*	32.01	0.82	0.145
	C19-C20	σ	C7-H7	σ^*	53.34	1.01	0.207
	C19-C20	σ	C8-H8	σ^*	32.23	0.73	0.137
	C19-C20	σ	C13-C14	π^*	1909.55	0.04	0.262
	C19-C20	σ	C14-H14	σ^*	17.35	0.84	0.108
	C19-C20	σ	C14-C15	σ^*	12.04	0.96	0.096
	C19-C20	σ	C16-C17	σ^*	42.21	0.65	0.147
	C19-C20	σ	C19-C20	σ^*	11.45	1.96	0.134
	C19-C20	σ	C19-C20	π^*	12.62	2.25	0.159
	C19-C20	σ	C10	LP*	10.99	0.04	0.029
	C19-C20	π	N1-C11	π^*	10.25	0.09	0.03
	C19-C20	π	C16-C17	σ^*	15.59	0.28	0.059
	C19-C20	π	C19-C20	σ^*	18.87	1.59	0.156
	C20-H20	σ	S1-N2	σ^*	108.32	0.30	0.169
	C20-H20	σ	S1-C15	σ^*	11.29	0.36	0.060
	C20-H20	σ	S2-C18	σ^*	15.63	0.34	0.066
	C20-H20	σ	S2-C19	σ^*	12.63	0.40	0.063
	C20-H20	σ	O1-C1	σ^*	153.06	0.50	0.247

	C20-H20	σ	O1-C1	π^*	514.97	0.05	0.160
	C20-H20	σ	N1-C12	σ^*	25.31	0.56	0.107
	C20-H20	σ	N2-C18	σ^*	58.84	0.61	0.169
	C20-H20	σ	N3-H3	σ^*	127.53	0.36	0.193
	C20-H20	σ	N3-C18	σ^*	123.77	0.36	0.190
	C20-H20	σ	N3-C20	σ^*	178.70	0.42	0.245
	C20-H20	σ	C1-C10	σ^*	35.48	0.62	0.133
	C20-H20	σ	C2-H2	σ^*	47.34	0.57	0.147
	C20-H20	σ	C3-C4	σ^*	280.99	0.35	0.281
	C20-H20	σ	C4-C5	σ^*	17.90	0.71	0.101
	C20-H20	σ	C4-C9	σ^*	43.58	0.64	0.150
	C20-H20	σ	C5-H5	σ^*	128.94	0.46	0.219
	C20-H20	σ	C6-H6	σ^*	32.84	0.60	0.125
	C20-H20	σ	C6-C7	σ^*	44.44	0.67	0.15
	C20-H20	σ	C7-H7	σ^*	26.15	0.54	0.106
	C20-H20	σ	C8-H8	σ^*	483.59	0.25	0.313
	C20-H20	σ	C9-C10	σ^*	35.16	0.64	0.134
	C20-H20	σ	C11-H11	σ^*	94.60	0.49	0.193
	C20-H20	σ	C12-C13	σ^*	43.74	0.71	0.157
	C20-H20	σ	C12-C17	σ^*	13.27	0.73	0.088
	C20-H20	σ	C12-C17	π^*	40.87	0.19	0.086
	C20-H20	σ	C14-H14	σ^*	232.13	0.37	0.263
	C20-H20	σ	C14-C15	σ^*	297.77	0.49	0.340
	C20-H20	σ	C15-C16	σ^*	16.07	0.81	0.102
	C20-H20	σ	C15-C16	π^*	85.43	0.17	0.119
	C20-H20	σ	C16-C17	σ^*	1183.19	0.17	0.404
	C20-H20	σ	C19-C20	σ^*	254.36	1.48	0.550
	C20-H20	σ	C19-C20	π^*	41.82	1.78	0.256
	S2	LP	N2-C18	π^*	35.28	0.21	0.082
	O1	LP	N1-H1	σ^*	16.20	0.67	0.094
	O1	LP	C1-C2	σ^*	15.17	0.79	0.099
	O1	LP	C1-C10	σ^*	12.60	0.70	0.085

	O2	LP	C10	LP*	27.79	0.05	0.046
	O2	LP	S1-N2	σ^*	33.71	0.41	0.106
	O2	LP	S1-C15	σ^*	16.06	0.47	0.078
	O2	LP	O1-C1	π^*	15.95	0.16	0.048
	O2	LP	N1-C11	π^*	26.86	0.10	0.050
	O2	LP	C3-C4	σ^*	12.82	0.46	0.071
	O2	LP	C8-H8	σ^*	18.50	0.36	0.076
	O2	LP	C14-C15	σ^*	14.20	0.59	0.085
	O2	LP	C16-C17	σ^*	38.46	0.28	0.096
	O2	LP	C19-C20	σ^*	11.85	1.59	0.128
	O2	LP	S1-O3	σ^*	21.00	0.56	0.097
	O2	LP	S1-N2	σ^*	10.24	0.39	0.057
	O3	LP	S1-C15	σ^*	15.65	0.46	0.076
	O3	LP	S1-O2	σ^*	16.80	0.57	0.088
	O3	LP	S1-N2	σ^*	19.54	0.40	0.079
	N2	LP	S2-C18	σ^*	19.06	0.54	0.092
	N2	LP	N2-C18	π^*	68.12	0.24	0.117
	C16	LP	C2-C3	π^*	48.99	0.15	0.099
	C16	LP	C5-C6	π^*	63.96	0.14	0.104
	C10	LP	C9	LP*	110.94	0.16	0.123
	C10	LP	S1-N2	σ^*	33.64	0.36	0.119
	C10	LP	O1-C1	σ^*	48.54	0.55	0.193
	C10	LP	O1-C1	π^*	14.12	0.11	0.040
	C10	LP	N1-C11	π^*	23.53	0.05	0.034
	C10	LP	N2-C18	σ^*	24.53	0.67	0.150
	C10	LP	N3-H3	σ^*	61.89	0.42	0.189
	C10	LP	N3-C18	σ^*	73.59	0.42	0.204
	C10	LP	N3-C20	σ^*	42.25	0.48	0.167
	C10	LP	C1-C10	σ^*	12.17	0.68	0.105
	C10	LP	C2-H2	σ^*	16.26	0.63	0.119
	C10	LP	C3-C4	σ^*	94.51	0.41	0.230
	C10	LP	C4-C9	σ^*	14.95	0.70	0.119

	C10	LP	C5-H5	σ^*	40.85	0.52	0.171
	C10	LP	C6-C7	σ^*	15.75	0.73	0.126
	C10	LP	C7-H7	σ^*	22.29	0.59	0.135
	C10	LP	C8-H8	σ^*	138.98	0.31	0.244
	C10	LP	C9-C10	σ^*	11.39	0.69	0.104
	C10	LP	C11-H11	σ^*	25.91	0.55	0.140
	C10	LP	C12-C17	π^*	16.34	0.25	0.065
	C10	LP	C14-H14	σ^*	77.78	0.43	0.214
	C10	LP	C14-C15	σ^*	98.43	0.54	0.270
	C10	LP	C15-C16	π^*	39.87	0.23	0.097
	C10	LP	C16-C17	σ^*	292.73	0.23	0.305
	C10	LP	C19-C20	σ^*	89.46	1.54	0.435
	C10	LP	C19-C20	π^*	23.71	1.84	0.226
ACPIN	C11-C21	σ	C22-C23	σ^*	10.92	1.07	0.097
	N1-C11	σ	C1	LP*	15.24	0.75	0.123
	N1-C11	σ	C11-C21	σ^*	22.06	0.98	0.132
	N1-C11	σ	O1-C1	σ^*	13.43	1.41	0.123
	N1-C11	σ	C1-C2	σ^*	54.37	1.09	0.219
	N1-C11	σ	C1-C10	σ^*	23.67	1.28	0.157
	N1-C11	σ	C4-C5	σ^*	94.15	0.96	0.269
	N1-C11	σ	C4-C5	π^*	68.66	0.66	0.213
	N1-C11	σ	C4-C9	σ^*	44.00	1.18	0.204
	N1-C11	σ	C6-C7	σ^*	18.50	1.45	0.146
	N1-C11	σ	C6-C7	π^*	17.24	0.92	0.125
	N1-C11	σ	C7-C8	σ^*	135.10	0.84	0.301
	N1-C11	σ	C8-H8	σ^*	175.12	0.68	0.308
	N1-C11	σ	C8-C9	σ^*	25.94	1.31	0.165
	N1-C11	σ	C9-C10	σ^*	30.07	1.33	0.179
	N1-C11	σ	C10-C11	π^*	13.04	0.91	0.105
	N1-C11	σ	C13-C18	σ^*	10.20	1.41	0.108
	N1-C11	σ	C15-C16	σ^*	37.42	1.22	0.191
	N1-C11	σ	C16-H16A	σ^*	213.17	0.59	0.317

	N1-C11	σ	C17-H17A	σ^*	13.95	1.33	0.122
	N1-C11	σ	C18-C19	π^*	125.52	0.64	0.278
	N1-C11	σ	C18-C23	σ^*	11.38	3.35	0.175
	N1-C11	σ	C19-C20	σ^*	111.63	1.04	0.304
	N1-C11	σ	C20-H20	σ^*	19.91	1.28	0.143
	N1-C11	σ	C20-C21	σ^*	163.51	0.87	0.339
	N1-C11	σ	C20-C21	π^*	4137.13	0.04	0.388
	N1-C11	σ	C22-C23	σ^*	318.92	1.34	0.584
	N1-C11	σ	C22-C23	π^*	210.62	1.15	0.476
	N1-C12	σ	C1	LP*	36.66	0.97	0.217
	N1-C12	σ	C11-C21	σ^*	55.54	1.20	0.231
	N1-C12	σ	O1-C1	σ^*	31.19	1.63	0.202
	N1-C12	σ	N1-C11	σ^*	11.22	1.69	0.123
	N1-C12	σ	N3-H3A	σ^*	16.50	1.57	0.144
	N1-C12	σ	N3-C13	σ^*	23.81	1.59	0.174
	N1-C12	σ	N4-C12	σ^*	28.51	1.67	0.196
	N1-C12	σ	N4-C15	σ^*	11.63	1.70	0.126
	N1-C12	σ	C1-C2	σ^*	142.93	1.31	0.389
	N1-C12	σ	C1-C10	σ^*	58.66	1.50	0.267
	N1-C12	σ	C2-C3	σ^*	21.08	1.75	0.172
	N1-C12	σ	C2-C3	π^*	22.85	1.18	0.15
	N1-C12	σ	C3-H3	σ^*	25.36	1.56	0.17
	N1-C12	σ	C4-C5	σ^*	233.44	1.17	0.468
	N1-C12	σ	C4-C5	π^*	159.87	0.88	0.374
	N1-C12	σ	C4-C9	σ^*	111.39	1.39	0.353
	N1-C12	σ	C6-C7	σ^*	50.19	1.66	0.258
	N1-C12	σ	C6-C7	π^*	43.05	1.14	0.219
	N1-C12	σ	C7-C8	σ^*	342.15	1.05	0.536
	N1-C12	σ	C8-H8	σ^*	404.57	0.89	0.537
	N1-C12	σ	C8-C9	σ^*	76.17	1.52	0.305
	N1-C12	σ	C9-C10	σ^*	44.21	1.55	0.234
	N1-C12	σ	C10-C11	π^*	32.57	1.13	0.185

	N1-C12	σ	C13-C14	σ^*	25.55	1.62	0.183
	N1-C12	σ	C15-C16	σ^*	102.89	1.43	0.344
	N1-C12	σ	C16-H16A	σ^*	486.04	0.81	0.560
	N1-C12	σ	C16-H16B	σ^*	3645.75	0.19	0.739
	N1-C12	σ	C17-H17A	σ^*	36.77	1.55	0.213
	N1-C12	σ	C18-C19	π^*	292.45	0.86	0.490
	N1-C12	σ	C18-C23	σ^*	32.68	3.57	0.30
	N1-C12	σ	C19-C20	σ^*	284.68	1.26	0.534
	N1-C12	σ	C20-H20	σ^*	52.51	1.50	0.251
	N1-C12	σ	C20-C21	σ^*	404.70	1.09	0.595
	N1-C12	σ	C20-C21	π^*	1853.10	0.25	0.678
	N1-C12	σ	C22-C23	σ^*	854.05	1.56	1.029
	N1-C12	σ	C22-C23	π^*	549.35	1.36	0.838
	N2-C12	σ	C8-H8	σ^*	11.48	0.55	0.071
	N2-C12	σ	C16-H16B	σ^*	15.19	0.47	0.075
	N2-C12	σ	C22-C23	σ^*	19.62	1.21	0.138
	N2-C12	σ	C22-C23	π^*	11.89	1.02	0.107
	N2-C13	π	N4-C12	π^*	40.61	0.31	0.103
	N2-C13	σ	C1-C2	σ^*	22.74	1.02	0.137
	N2-C13	σ	C1-C10	σ^*	14.43	1.21	0.119
	N2-C13	σ	C4-C5	σ^*	54.14	0.89	0.196
	N2-C13	σ	C4-C5	π^*	36.89	0.59	0.148
	N2-C13	σ	C4-C9	σ^*	23.23	1.11	0.144
	N2-C13	σ	C6-C7	σ^*	10.39	1.38	0.107
	N2-C13	σ	C6-C7	π^*	10.59	0.85	0.094
	N2-C13	σ	C7-C8	σ^*	71.83	0.77	0.21
	N2-C13	σ	C8-H8	σ^*	87.74	0.61	0.206
	N2-C13	σ	C8-C9	σ^*	16.17	1.24	0.127
	N2-C13	σ	C13-C14	σ^*	13.28	1.34	0.120
	N2-C13	σ	C15-C16	σ^*	14.75	1.15	0.117
	N2-C13	σ	C16-H16B	σ^*	128.65	0.52	0.232
	N2-C13	σ	C18-C19	π^*	44.47	0.57	0.156

	N2-C13	σ	C19-C20	σ^*	76.74	0.97	0.244
	N2-C13	σ	C20-C21	σ^*	111.52	0.80	0.269
	N2-C13	σ	C22-C23	σ^*	198.68	1.27	0.449
	N2-C13	σ	C22-C23	π^*	130.96	1.08	0.364
	N4-C12	π	C8-H8	σ^*	22.95	0.01	0.016
	C2-C3	π	C1	LP*	169.24	0.04	0.085
	C4-C5	π	C1	LP*	13.36	0.02	0.014
	C4-C5	π	C2-C3	π^*	32.58	0.23	0.082
	C4-C5	π	C6-C7	π^*	49.83	0.18	0.086
	C4-C5	π	C7-C8	σ^*	18.42	0.10	0.043
	C4-C5	π	C8-C9	π^*	22.75	0.28	0.071
	C4-C5	π	C20-C21	σ^*	21.16	0.14	0.053
	C4-C5	π	C22-C23	σ^*	12.57	0.60	0.086
	C4-C5	π	C22-C23	π^*	14.04	0.41	0.069
	C4-C9	σ	C4-C5	σ^*	27.48	0.65	0.120
	C4-C9	σ	C8-H8	σ^*	42.41	0.37	0.113
	C4-C9	σ	C16-H16B	σ^*	19.54	0.29	0.067
	C4-C9	σ	C20-C21	σ^*	11.46	0.57	0.072
	C4-C9	σ	C22-C23	σ^*	12.72	1.04	0.103
	C4-C9	σ	C22-C23	π^*	14.59	0.84	0.107
	C6-H6	σ	C4-C5	σ^*	44.10	0.50	0.132
	C6-H6	σ	C4-C5	π^*	19.45	0.20	0.063
	C6-H6	σ	C8-H8	σ^*	52.16	0.22	0.095
	C6-H6	σ	C16-H16B	σ^*	80.08	0.13	0.091
	C6-H6	σ	C18-C19	π^*	34.71	0.18	0.077
	C6-H6	σ	C19-C20	σ^*	17.42	0.58	0.090
	C6-H6	σ	C20-C21	σ^*	30.13	0.41	0.100
	C6-H6	σ	C22-C23	σ^*	37.78	0.88	0.163
	C6-H6	σ	C22-C23	π^*	32.30	0.69	0.144
	C6-C7	σ	C8-H8	σ^*	10.88	0.38	0.058
	C6-C7	σ	C8-C9	π^*	18.83	0.28	0.065
	C7-H7	σ	C8-H8	σ^*	14.63	0.21	0.049

	C8-C9	π	C1	LP*	22.37	0.02	0.019
	C8-C9	π	C6-C7	π^*	25.79	0.19	0.062
	C8-C9	π	C10-C11	π^*	10.53	0.18	0.040
	C10-C11	π	C1	LP*	233.38	0.03	0.093
	C10-C11	π	C8-C9	π^*	18.92	0.30	0.068
	C13-C14	σ	C14-C15	σ^*	13.60	0.93	0.101
	C13-C14	σ	C18-C19	σ^*	33.62	0.36	0.106
	C14-C15	π	N2-C13	π^*	33.02	0.25	0.085
	C14-C15	π	N4-C12	π^*	10.48	0.26	0.048
	C14-C15	π	C7-C8	σ^*	11.60	0.12	0.036
	C14-C18	σ	C11-C21	σ^*	10.20	0.63	0.072
	C14-C18	σ	C1-C2	σ^*	29.89	0.75	0.134
	C14-C18	σ	C4-C5	π^*	20.60	0.32	0.080
	C14-C18	σ	C7-C8	σ^*	34.08	0.49	0.116
	C14-C18	σ	C8-H8	σ^*	68.48	0.33	0.135
	C14-C18	σ	C15-C16	σ^*	19.75	0.87	0.117
	C14-C18	σ	C16-H16B	σ^*	44.62	0.25	0.09
	C14-C18	σ	C18-C19	π^*	140.64	0.29	0.198
	C17-H17A	σ	C16-H16B	σ^*	15.28	0.10	0.035
	C18-C19	σ	C18-C19	π^*	19.84	0.36	0.082
	C18-C19	π	C22-C23	π^*	21.82	0.42	0.086
	C18-C23	σ	C1	LP*	47.34	0.50	0.177
	C18-C23	σ	C11-C21	σ^*	71.39	0.73	0.204
	C18-C23	σ	O1-C1	σ^*	32.25	1.17	0.174
	C18-C23	σ	N3-H3B	σ^*	13.36	1.10	0.109
	C18-C23	σ	N3-C13	σ^*	23.52	1.13	0.146
	C18-C23	σ	N4-C12	σ^*	16.67	1.21	0.127
	C18-C23	σ	C1-C2	σ^*	186.71	0.85	0.356
	C18-C23	σ	C1-C10	σ^*	36.44	1.03	0.174
	C18-C23	σ	C2-C3	σ^*	13.89	1.29	0.120
	C18-C23	σ	C2-C3	π^*	23.00	0.72	0.119
	C18-C23	σ	C3-H3	σ^*	20.72	1.10	0.135

	C18-C23	σ	C4-C5	σ^*	192.97	0.71	0.331
	C18-C23	σ	C4-C5	π^*	207.61	0.42	0.292
	C18-C23	σ	C4-C9	σ^*	100.53	0.93	0.273
	C18-C23	σ	C6-C7	σ^*	32.05	1.20	0.176
	C18-C23	σ	C6-C7	π^*	27.87	0.67	0.135
	C18-C23	σ	C7-C8	σ^*	385.71	0.59	0.428
	C18-C23	σ	C8-H8	σ^*	581.82	0.43	0.448
	C18-C23	σ	C8-C9	σ^*	52.23	1.06	0.210
	C18-C23	σ	C9-C10	σ^*	35.78	1.08	0.176
	C18-C23	σ	C10-C11	π^*	21.98	0.67	0.116
	C18-C23	σ	C13-C14	σ^*	25.86	1.16	0.155
	C18-C23	σ	C15-C16	σ^*	166.52	0.97	0.359
	C18-C23	σ	C16-H16B	σ^*	604.48	0.34	0.409
	C18-C23	σ	C17-H17A	σ^*	63.59	1.08	0.236
	C18-C23	σ	C18-C19	π^*	758.97	0.39	0.532
	C18-C23	σ	C18-C23	σ^*	17.47	3.11	0.208
	C18-C23	σ	C19-C20	σ^*	126.43	0.79	0.284
	C18-C23	σ	C20-H20	σ^*	41.35	1.04	0.186
	C18-C23	σ	C20-C21	σ^*	214.08	0.63	0.327
	C18-C23	σ	C22-C23	σ^*	389.83	1.09	0.585
	C18-C23	σ	C22-C23	π^*	384.68	0.90	0.567
	C19-C20	σ	C20-C21	σ^*	11.52	0.59	0.074
	C20-C21	π	C22-C23	π^*	14.87	0.44	0.072
	C21-C22	σ	C18-C19	π^*	26.13	0.39	0.099
	C21-C22	σ	C20-C21	σ^*	19.11	0.62	0.098
	C22-H22	σ	C1-C2	σ^*	36.67	0.65	0.139
	C22-H22	σ	C4-C5	σ^*	18.85	0.52	0.088
	C22-H22	σ	C4-C5	π^*	39.21	0.22	0.093
	C22-H22	σ	C4-C9	σ^*	11.18	0.74	0.081
	C22-H22	σ	C7-C8	σ^*	57.89	0.40	0.136
	C22-H22	σ	C8-H8	σ^*	120.41	0.24	0.151
	C22-H22	σ	C15-C16	σ^*	24.66	0.78	0.124

	C22-H22	σ	C16-H16B	σ^*	107.54	0.15	0.115
	C22-H22	σ	C17-H17A	σ^*	11.80	0.89	0.092
	C22-H22	σ	C18-C19	π^*	225.24	0.20	0.208
	C22-H22	σ	C22-C23	σ^*	60.17	0.90	0.208
	C22-H22	σ	C22-C23	π^*	22.57	0.71	0.122
	C22-C23	σ	C1	LP*	48.73	0.48	0.174
	C22-C23	σ	C11-C21	σ^*	121.29	0.70	0.261
	C22-C23	σ	O1-C1	σ^*	32.26	1.14	0.172
	C22-C23	σ	N3-H3B	σ^*	12.91	1.08	0.106
	C22-C23	σ	N3-C13	σ^*	25.70	1.10	0.151
	C22-C23	σ	N4-C12	σ^*	16.29	1.18	0.124
	C22-C23	σ	C1-C2	σ^*	198.02	0.82	0.360
	C22-C23	σ	C1-C10	σ^*	45.20	1.01	0.192
	C22-C23	σ	C2-C3	σ^*	13.83	1.26	0.118
	C22-C23	σ	C2-C3	π^*	22.92	0.69	0.116
	C22-C23	σ	C3-H3	σ^*	20.17	1.07	0.132
	C22-C23	σ	C4-C5	σ^*	197.10	0.68	0.328
	C22-C23	σ	C4-C5	π^*	226.23	0.39	0.294
	C22-C23	σ	C4-C9	σ^*	103.13	0.90	0.272
	C22-C23	σ	C6-C7	σ^*	38.19	1.17	0.190
	C22-C23	σ	C6-C7	π^*	34.95	0.65	0.148
	C22-C23	σ	C7-C8	σ^*	402.60	0.56	0.427
	C22-C23	σ	C8-H8	σ^*	624.78	0.40	0.449
	C22-C23	σ	C8-C9	σ^*	60.54	1.03	0.224
	C22-C23	σ	C9-C10	σ^*	35.87	1.05	0.174
	C22-C23	σ	C10-C11	π^*	23.06	0.64	0.116
	C22-C23	σ	C13-C14	σ^*	22.97	1.13	0.144
	C22-C23	σ	C15-C16	σ^*	139.41	0.94	0.324
	C22-C23	σ	C16-H16B	σ^*	642.46	0.32	0.404
	C22-C23	σ	C17-H17A	σ^*	63.66	1.06	0.233
	C22-C23	σ	C18-C19	π^*	827.31	0.36	0.535
	C22-C23	σ	C19-C20	σ^*	138.68	0.76	0.292

	C22-C23	σ	C20-H20	σ^*	54.36	1.01	0.210
	C22-C23	σ	C20-C21	σ^*	266.14	0.60	0.356
	C22-C23	σ	C22-C23	σ^*	832.74	1.07	0.844
	C22-C23	σ	C22-C23	π^*	393.95	0.87	0.564
	C23-H23	σ	C11-C21	σ^*	77.20	0.08	0.072
	C23-H23	σ	O2-C1	σ^*	21.79	0.52	0.095
	C23-H23	σ	N3-H3B	σ^*	20.65	0.46	0.087
	C23-H23	σ	N3-C13	σ^*	18.35	0.48	0.084
	C23-H23	σ	N4-C12	σ^*	16.21	0.56	0.085
	C23-H23	σ	C1-C2	σ^*	135.13	0.20	0.147
	C23-H23	σ	C1-C10	σ^*	75.75	0.39	0.154
	C23-H23	σ	C2-C3	σ^*	17.67	0.64	0.095
	C23-H23	σ	C2-C3	π^*	83.12	0.07	0.072
	C23-H23	σ	C3-H3	σ^*	22.83	0.45	0.091
	C23-H23	σ	C4-C5	σ^*	1349.18	0.06	0.260
	C23-H23	σ	C4-C9	σ^*	147.28	0.28	0.183
	C23-H23	σ	C6-C7	σ^*	39.33	0.55	0.132
	C23-H23	σ	C6-C7	π^*	492.58	0.03	0.113
	C23-H23	σ	C8-C9	σ^*	79.59	0.41	0.162
	C23-H23	σ	C9-C10	σ^*	37.99	0.43	0.115
	C23-H23	σ	C10-C11	π^*	654.23	0.02	0.107
	C23-H23	σ	C13-C14	σ^*	19.16	0.51	0.089
	C23-H23	σ	C15-C16	σ^*	92.33	0.32	0.154
	C23-H23	σ	C17-H17A	σ^*	22.22	0.44	0.088
	C23-H23	σ	C18-C23	σ^*	12.83	2.46	0.159
	C23-H23	σ	C19-C20	σ^*	806.57	0.15	0.306
	C23-H23	σ	C20-H20	σ^*	38.13	0.39	0.109
	C23-H23	σ	C22-C23	σ^*	701.89	0.45	0.500
	C23-H23	σ	C22-C23	π^*	709.38	0.25	0.409
	C11	LP	C20-C21	σ^*	11.15	0.83	0.086
	C11	LP	C8-H8	σ^*	11.24	0.02	0.013
	C11	LP	C20-C21	σ^*	27.33	0.21	0.068

	C11	LP	C7-C8	σ^*	10.18	0.18	0.039
	C11	LP	C8-H8	σ^*	94.00	0.02	0.039
	C11	LP	C20-C21	σ^*	11.49	0.22	0.045
	C11	LP	C22-C23	σ^*	12.04	0.68	0.082
	C11	LP	C22-C23	π^*	10.13	0.49	0.067
	O1	LP	N1-H1	σ^*	11.43	0.67	0.079
	O1	LP	C1-C2	σ^*	32.45	0.39	0.102
	O1	LP	C1-C10	σ^*	15.72	0.58	0.086
	O1	LP	C1	LP*	1242.28	0.02	0.155
	N1	LP	N4-C12	π^*	50.69	0.26	0.103
	N1	LP	C10-C11	π^*	66.53	0.20	0.107
	N2	LP	N4-C12	σ^*	13.02	0.83	0.094
	N2	LP	C13-C14	σ^*	10.38	0.79	0.082
	N3	LP	N2-C13	σ^*	53.91	0.26	0.112
	N4	LP	C1	LP*	46.37	0.23	0.115
	N4	LP	C11-C21	σ^*	40.17	0.46	0.123
	N4	LP	O1-C1	σ^*	15.88	0.89	0.109
	N4	LP	N1-C12	σ^*	5.40	0.84	0.103
	N4	LP	N3-C13	σ^*	14.40	0.85	0.101
	N4	LP	N4-C12	σ^*	11.13	0.93	0.092
	N4	LP	C1-C2	σ^*	90.14	0.57	0.205
	N4	LP	C1-C10	σ^*	36.56	0.76	0.150
	N4	LP	C2-C3	π^*	17.37	0.44	0.080
	N4	LP	C3-H3	σ^*	12.81	0.82	0.093
	N4	LP	C4-C5	σ^*	172.63	0.43	0.249
	N4	LP	C4-C5	π^*	292.13	0.14	0.197
	N4	LP	C4-C9	σ^*	66.76	0.65	0.190
	N4	LP	C6-C7	σ^*	25.18	0.92	0.139
	N4	LP	C6-C7	π^*	36.66	0.40	0.116
	N4	LP	C7-C8	σ^*	305.10	0.32	0.283
	N4	LP	C8-H8	σ^*	631.04	0.15	0.284
	N4	LP	C8-C9	σ^*	40.88	0.78	0.163

	N4	LP	C9-C10	σ^*	22.42	0.81	0.122
	N4	LP	C10-C11	π^*	27.77	0.39	0.098
	N4	LP	C13-C14	σ^*	5.93	0.88	0.107
	N4	LP	C15-C16	σ^*	38.06	0.69	0.148
	N4	LP	C16-H16B	σ^*	1709.60	0.07	0.312
	N4	LP	C17-H17A	σ^*	17.64	0.81	0.109
	N4	LP	C18-C19	π^*	659.02	0.12	0.265
	N4	LP	C19-C20	σ^*	185.98	0.52	0.282
	N4	LP	C20-H20	σ^*	29.51	0.76	0.137
	N4	LP	C20-C21	σ^*	342.64	0.35	0.315
	N4	LP	C22-C23	σ^*	460.09	0.82	0.559
	N4	LP	C22-C23	π^*	336.03	0.62	0.433

^aThe interaction energy threshold is 10 kcal/mol.