

Table S1. Bond lengths [$\text{\AA}$] and angles [$^\circ$] for compound **1** according to X-Ray diffraction data.

S(1)-C(1)	1.8142(4)	C(1)-S(1)-C(2)	90.413(15)
S(1)-C(2)	1.8605(4)	N(1A)-N(1)-C(2A)	104.81(3)
N(1)-N(1A)	1.4411(5)	N(1A)-N(1)-C(1)	108.538(19)
N(1)-C(2A)	1.4435(4)	C(2A)-N(1)-C(1)	113.59(2)
N(1)-C(1)	1.4891(4)	N(1)-C(1)-S(1)	105.061(19)
C(1)-H(1B)	0.988(8)	N(1)-C(1)-H(1B)	107.2(5)
C(1)-H(1A)	0.974(8)	S(1)-C(1)-H(1B)	109.3(5)
C(2)-N(1A)	1.4435(4)	N(1)-C(1)-H(1A)	112.6(5)
C(2)-H(2B)	0.968(9)	S(1)-C(1)-H(1A)	108.7(5)
C(2)-H(2A)	0.963(8)	H(1B)-C(1)-H(1A)	113.6(7)
		N(1A)-C(2)-S(1)	107.16(2)
		N(1A)-C(2)-H(2B)	110.3(5)
		S(1)-C(2)-H(2B)	110.6(5)
		N(1A)-C(2)-H(2A)	111.7(5)
		S(1)-C(2)-H(2A)	109.6(5)
		H(2B)-C(2)-H(2A)	107.4(7)

Table S2. Torsion and pseudo-torsion angles [$^\circ$] for compound **1** according to X-Ray diffraction data.

C(1)-N(1)-N(1A)-C(2)	53.32
N(1)-C(1)-S(1)-C(2)	13.19
N(1A)-C(2)-S(1)-C(1)	15.5
C(1)-N(1)-N(1A)-C(1A)	-68.36
S(1)-C(1)-N(1)-N(1A)	-40.21
S(1)-C(2)-N(1A)-N(1)	-40.6
C(2A)-N(1)-N(1A)-C(2)	175.01
S(1)-C(1)-N(1)-C(2A)	-156.33
S(1)-C(2)-N(1A)-C(1A)	77.69
C(1)-N(1)-N(1A)-lp	171
lp-N(1)-N(1A)-lp	51
S(1)-C(1)-N(1)-lp	80
H(1A)-C(1)-N(1)-C(2A)	-38
H(1B)-C(1)-N(1)-C(2A)	88
S(1)-C(2)-N(1A)-lp	-159
H(2A)-C(2)-N(1A)-lp	-40
H(2B)-C(2)-N(1A)-lp	80
lp-N(1)-N(1A)-C(2)	-67
H(1A)-C(1)-S(1)-C(2)	-107
H(1A)-C(1)-N(1)-lp	-162
H(1B)-C(1)-N(1)-lp	-35
H(2A)-C(2)-S(1)-C(1)	-104
H(2A)-C(2)-N(1A)-N(1)	78
H(2B)-C(2)-N(1A)-N(1)	-162
lp-N(1)-N(1A)-C(1A)	171
H(1B)-C(1)-S(1)-C(2)	128
H(1A)-C(1)-N(1)-N(1A)	78
H(1B)-C(1)-N(1)-N(1A)	-155
H(2B)-C(2)-S(1)-C(1)	138

H(2A)-C(2)-N(1A)-C(1A)	-164
H(2B)-C(2)-N(1A)-C(1A)	-44

Table S3. Summary of symmetrically independent critical points (3,-1) for intermolecular contacts in the crystal of **1**.

	symmetry operation	$\rho(r)$, eÅ ⁻³	$\nabla^2\rho(r)$, eÅ ⁻³	$v(r)$, a.u.	E, kcal/mol
S(1) - H(2A)	x, y-1, z	0.04	0.67	-0.00344	1.1
S(1) - H(1B)	1/2-x, -1/2+y, 1/2-z	0.05	0.56	-0.00343	1.1
S(1) - H(2B)	1/2-x, 1/2-y, -z	0.04	0.47	-0.00287	0.9
S(1) - S(1)	1/2-x, 1/2-y, -z	0.04	0.37	-0.00238	0.7
S(1) - S(1)	1/2-x, -1/2+y, 1/2-z	0.04	0.45	-0.0029	0.9
H(2B) - H(2B)	1-x, 1-y, -z	0.04	0.56	-0.00328	1.0
H(1A) - N(1)	x, y-1, z	0.03	0.56	-0.00258	0.8
H(2B) - H(1A)	x, -y, -1/2+z	0.04	0.56	-0.00288	0.9

Table S4. Summary of symmetrically independent $n \rightarrow \sigma^*$ NBO interactions in conformer **1** with energy > 1 kcal/mol according B3LYP/6-311G*

Atoms	NBO energy (kcal/mol)
<i>lp</i> -S(1)-C(1)-H(1A)	4.31
<i>lp</i> -S(1)-C(1)-H(1E)	3.82
<i>lp</i> -S(1)-C(2)-H(2E)	1.75
<i>lp</i> -S(1)-C(2)-H(2A)	4.18
<i>lp</i> -N(1)-C(1)-H(1A)	5.22
<i>lp</i> -N(1)-C(1)-H(1E)	2.95
<i>lp</i> -N(1A)-C(2)-S(1)	12.14
<i>lp</i> -N(1A)-N(1)-C(1)	7.43
<i>lp</i> -N(1A)-C(4)-H(2A)	2.96

Table S5. Summary of symmetrically independent $n \rightarrow \sigma^*$ NBO interactions in conformer **1B** with energy > 1 kcal/mol according B3LYP/6-311G*

Atoms	NBO energy (kcal/mol)
<i>lp</i> -S(1)-C(1)-N(1)	2.59
<i>lp</i> -S(1)-C(1)-H(1A)	4.74
<i>lp</i> -N(1)-C(1)-H(1E)	2.10
<i>lp</i> -N(1)-C(1)-H(1A)	5.89

Table S6. Summary of symmetrically independent $n \rightarrow \sigma^*$ NBO interactions in conformer **1C** with energy > 1 kcal/mol according B3LYP/6-311G*

Atoms	NBO energy (kcal/mol)
<i>lp</i> -S(1)-C(1)-N(1)	3.20
<i>lp</i> -S(1)-C(1)-H(1A)	5.07
<i>lp</i> -N(1)-C(1)-H(1A)	5.61
<i>lp</i> -N(1)-C(1)-H(1E)	3.52
<i>lp</i> -N(1)-N(1A)-C(1A)	1.83
<i>lp</i> -N(1)-C(2)-S(2)	10.41
<i>lp</i> -N(1)-N(1A)-C(2A)	1.74
<i>lp</i> -N(1)-C(2)-H(2A)	3.91

Table S7. Summary of symmetrically independent $n \rightarrow \sigma^*$ NBO interactions in conformer **1A** with energy > 1 kcal/mol according B3LYP/6-311G*

Atoms	NBO energy (kcal/mol)
<i>lp</i> -S(1)-C(1)-N(1)	2.95
<i>lp</i> -S(1)-C(1)-H(1A)	5.02
<i>lp</i> -N(1)-C(1)-H(1A)	6.52
<i>lp</i> -N(1)-C(1)-H(1E)	3.62
<i>lp</i> -N(1)-N(1A)-C(1A)	1.73

Table S8. Summary of symmetrically independent $n \rightarrow \sigma^*$ NBO interactions in conformer **1D** with energy > 1 kcal/mol according B3LYP/6-311G*

Atoms	NBO energy (kcal/mol)
<i>lp</i> -S(1)-C(1)-N(1)	3.07
<i>lp</i> -S(1)-C(1)-H(1A)	4.67
<i>lp</i> -N(1)-C(1)-S(1)	9.28
<i>lp</i> -N(1)-N(1A)-C(1A)	1.65
<i>lp</i> -N(1)-C(1)-H(1A)	4.16

Multipole refinement

Table S9. Monopole Populations, Radial Parameters and Net Atomic Charges.

Atom	Pval	Kappa	P00	Kappa'	Net charge
S (1)	6.007 (19)	0.998	0.000	0.999	-0.006 (19)
N (1)	5.079 (13)	0.984	0.000	0.975	-0.079 (13)
C (1)	4.171 (21)	0.978	0.000	0.981	-0.171 (21)
C (2)	4.155 (20)	0.979	0.000	1.000	-0.155 (20)
H (1B)	0.910 (15)	1.200	0.000	1.200	+0.089 (15)
H (2B)	0.876 (14)	1.200	0.000	1.200	+0.123 (14)
H (2A)	0.898 (14)	1.200	0.000	1.200	+0.102 (14)
H (1A)	0.903 (13)	1.200	0.000	1.200	+0.096 (13)

Table S10. Dipole Population Parameters.

Atom	D11+	D11-	D10	Kappa'
S (1)	0.012 (11)	-0.030 (12)	-0.038 (10)	0.999
N (1)	-0.054 (7)	0.107 (6)	-0.025 (7)	0.975
C (1)	0.017 (9)	0.038 (9)	0.030 (9)	0.981
C (2)	-0.017 (8)	0.053 (10)	0.012 (9)	1.000
H (1B)	0.000	0.000	0.105 (10)	1.200
H (2B)	0.000	0.000	0.078 (11)	1.200
H (2A)	0.000	0.000	0.130 (11)	1.200
H (1A)	0.000	0.000	0.127 (10)	1.200

Table S11. Quadrupole Population Parameters.

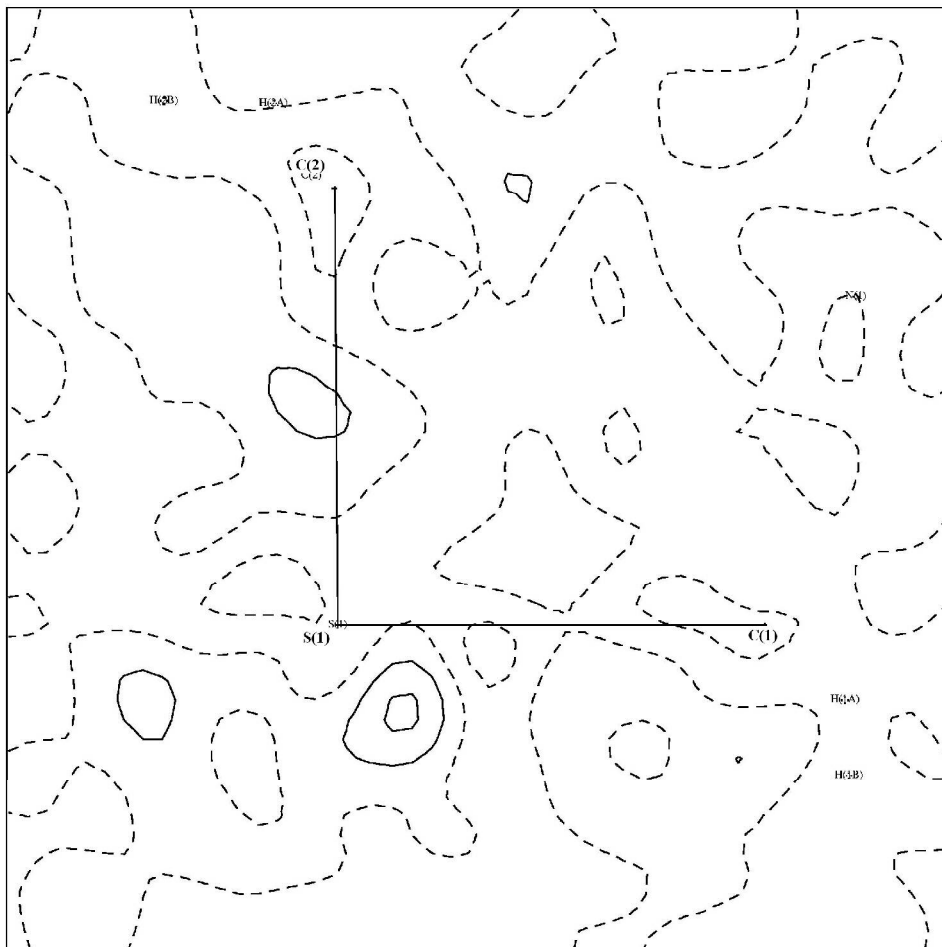
Atom	Q20	Q21+	Q21-	Q22+	Q22-	Kappa'
S (1)	-0.174 (13)	-0.060 (10)	0.054 (11)	0.126 (9)	0.020 (12)	0.999
N (1)	-0.018 (7)	0.016 (7)	-0.033 (7)	-0.001 (7)	-0.088 (7)	0.975
C (1)	0.051 (9)	0.011 (9)	-0.010 (8)	-0.064 (9)	-0.002 (9)	0.981
C (2)	0.055 (9)	0.009 (8)	-0.032 (8)	-0.076 (8)	0.000 (9)	1.000
H (1B)	0.000	0.000	0.000	0.000	0.000	1.200
H (2B)	0.000	0.000	0.000	0.000	0.000	1.200
H (2A)	0.000	0.000	0.000	0.000	0.000	1.200
H (1A)	0.000	0.000	0.000	0.000	0.000	1.200

Table S12. Octupole Population Parameters.

Atom	O30	O31+	O31-	O32+	O32-	O33+	O33-	Kappa'
S (1)	0.127 (12)	0.017 (11)	-0.051 (11)	0.032 (11)	-0.019 (10)	0.007 (10)	-0.055 (11)	0.999
N (1)	0.150 (10)	-0.006 (10)	0.007 (10)	-0.023 (9)	-0.058 (9)	0.114 (9)	0.037 (9)	0.975
C (1)	0.234 (12)	0.029 (11)	0.033 (11)	0.008 (11)	-0.022 (12)	0.015 (12)	-0.182 (11)	0.981
C (2)	0.222 (12)	0.007 (11)	-0.042 (11)	-0.001 (11)	0.005 (12)	0.007 (11)	-0.204 (12)	1.000

Differences of Mean-Squares Displacement Amplitudes (DMSDA)
(1.E4 Å**2) along interatomic vectors (*bonds)

ATOM-->	ATOM	/	DIST	DMSDA	ATOM	/	DIST	DMSDA	ATOM	/	DIST	DMSDA
S(1)	N(1)		2.6289	9	C(1)		1.8145	12	C(2)		1.8602	14
N(1)	C(1)		1.4882	3	C(2)		2.2852	4				
C(1)	C(2)		2.6077	2								



The residual electron density map in the S(1)C(1)C(2) plane. Contours are drawn with 0.1 eÅ⁻³ step, the negative and zero contours are dashed.