

Supplementary information for

## Sigmatropic rearrangements in 5-allyloxytetrazoles

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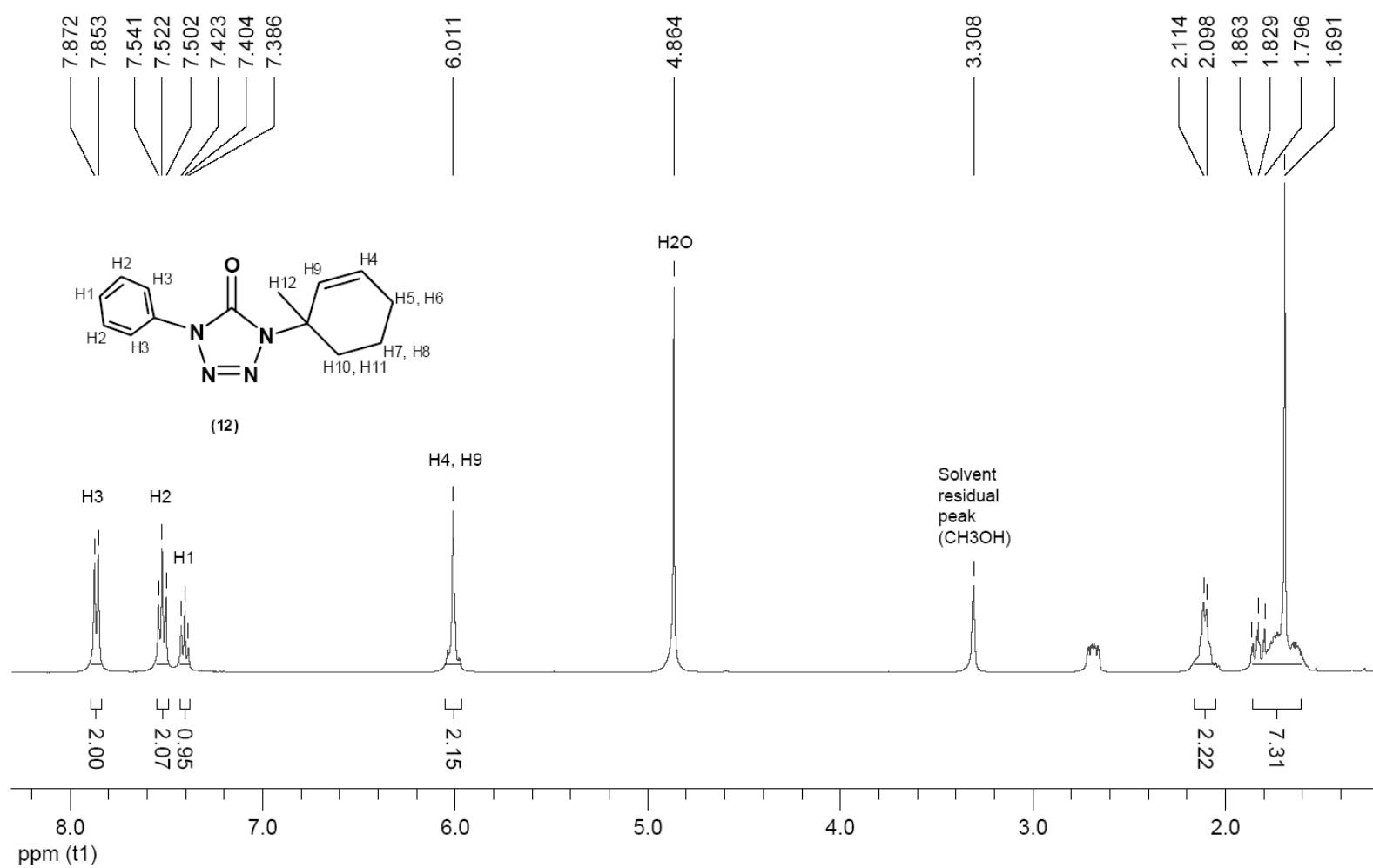
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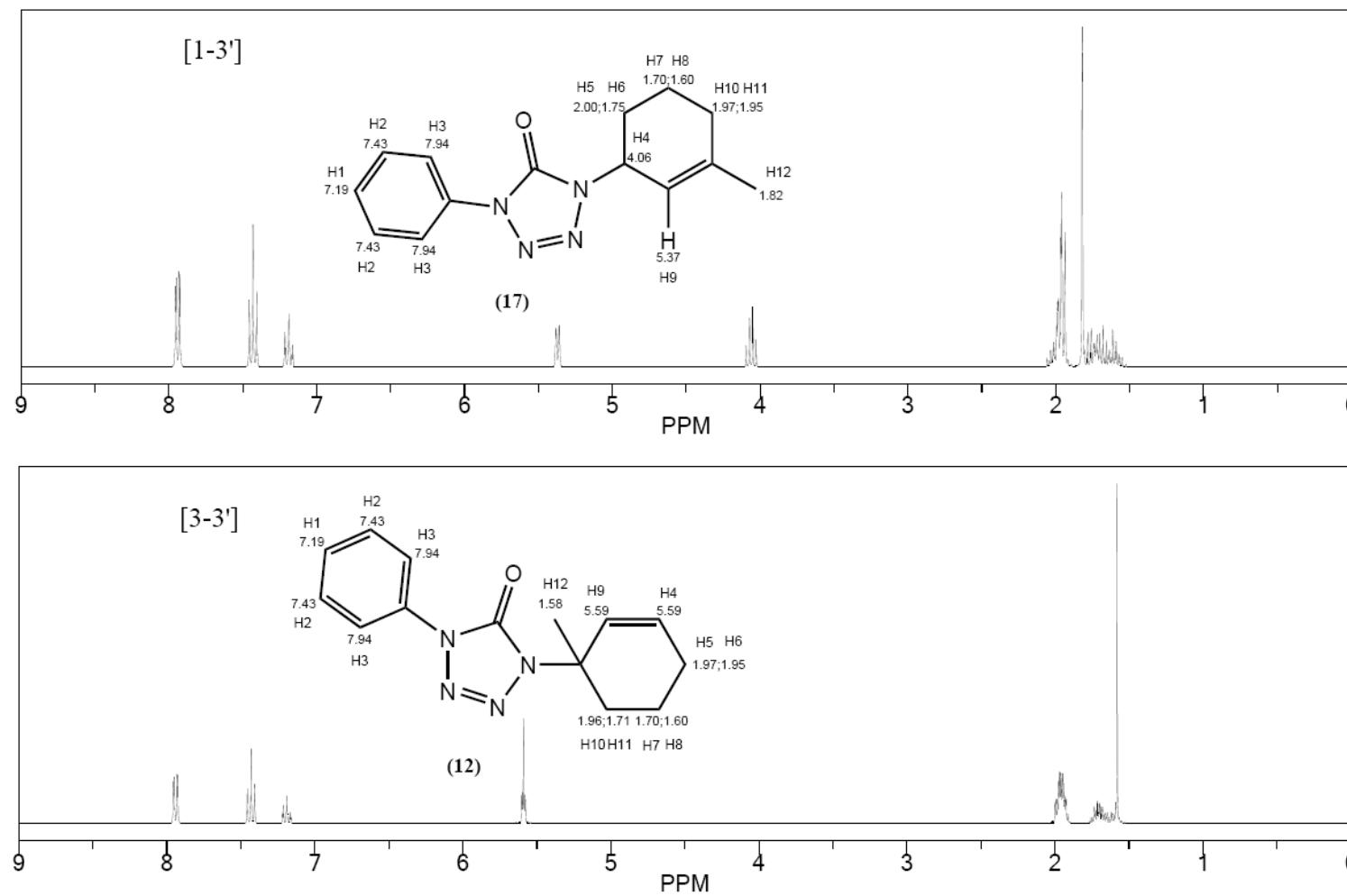
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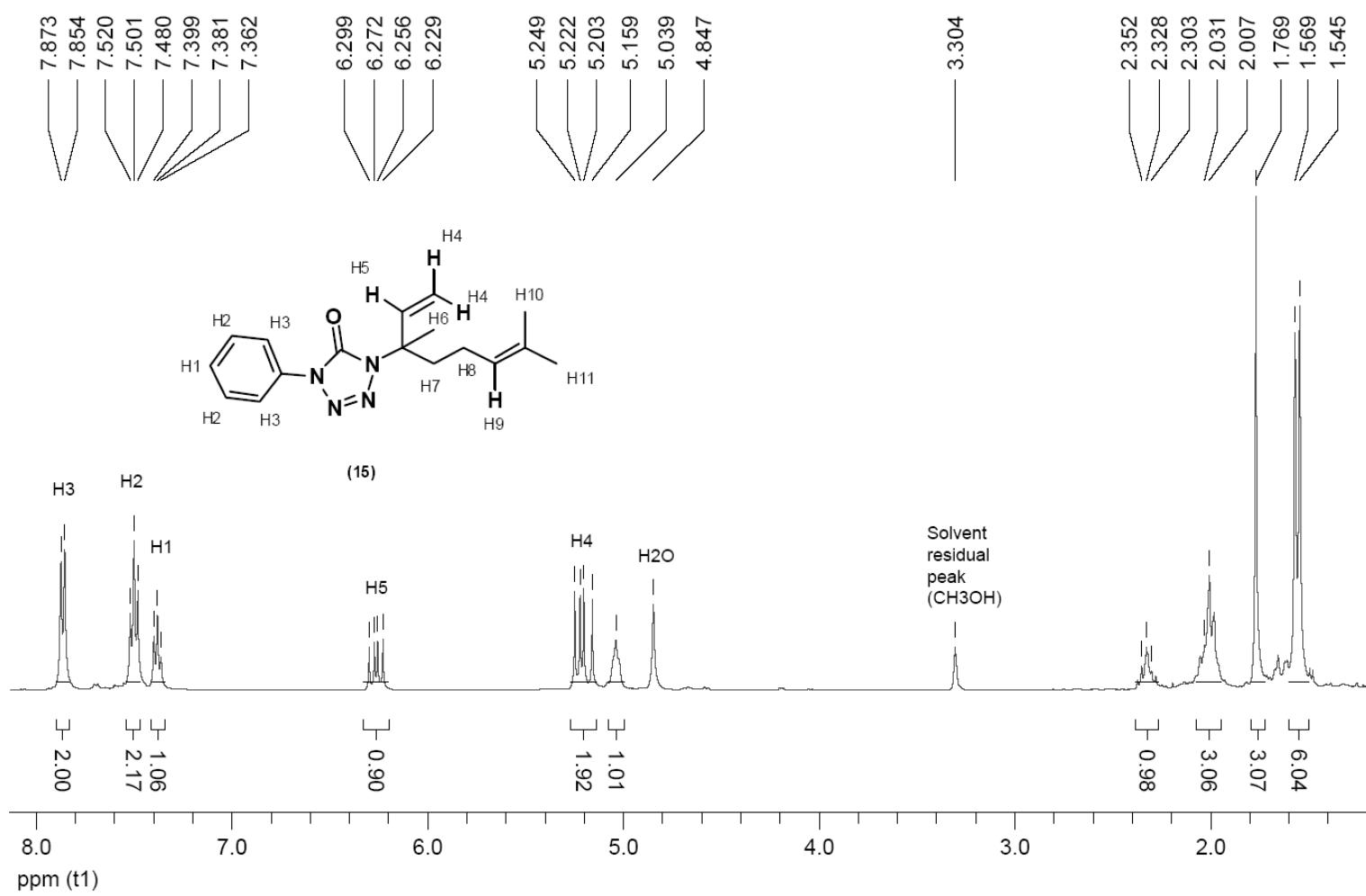
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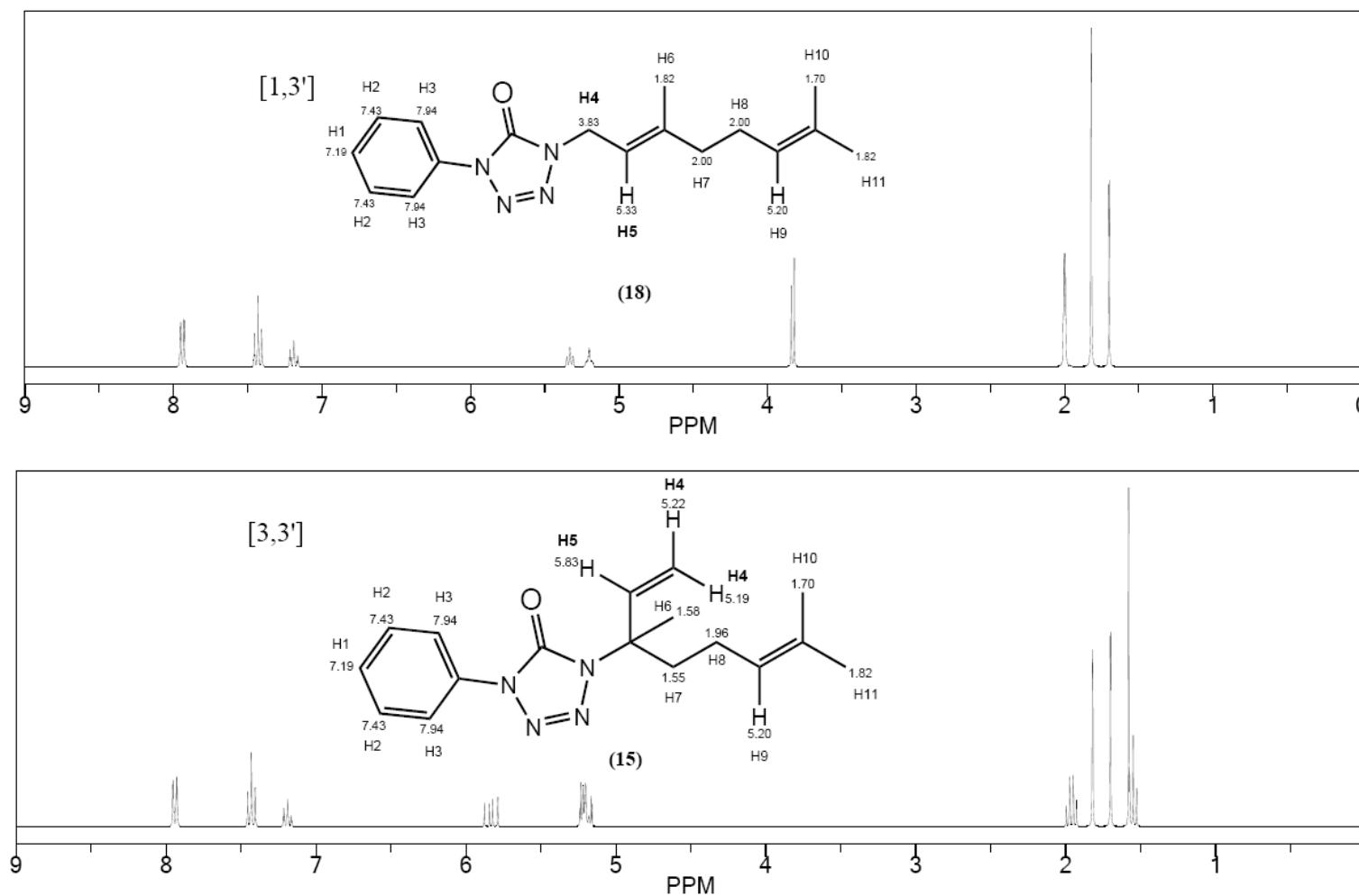
**Figure 1S.** <sup>1</sup>H-NMR spectrum of 4-(3-methylcyclohex-2-enyl)-1-phenyl-1*H*-tetrazol-5(4*H*)-one (**12**) (400 MHz, CD<sub>3</sub>OD).



**Figure 2S.** Predicted <sup>1</sup>H-NMR spectra of tetrazolones **17** and **12** ([1,3]- and [3,3]- migration products). (<sup>1</sup>H-NMR shifts were acquired using the ChemDraw (Ultra 9.0) software)



**Figure 3S.** <sup>1</sup>H-NMR spectrum of 1-(3,7-dimethylocta-1,6-dien-3-yl)-4-phenyl-1*H*-tetrazol-5(4*H*)-one (**15**) (400 MHz,  $\text{CD}_3\text{OD}$ ).



**Figure 4S.** Predicted <sup>1</sup>H-NMR spectra of tetrazolones **18** and **15** ([1,3]- and [3,3]- migration products).  
(<sup>1</sup>H-NMR shifts were acquired using the ChemDraw (Ultra 9.0) software)

**Table 1S.** Conformationally relevant geometric parameters and relative energies of conformers of 5-(Cyclohex-2-enyloxy)-1-phenyl-1*H*-tetrazole **9** [in brief: Ether **9** (R)] and corresponding isomeric 1-(cyclohex-2-enyl)-4-phenyltetrazol-5-one **16** [in brief: Tetrazolone **16** (R)] calculated at the DFT(B3LYP)/6-31G(d,p) level of theory.

Compound / Conformer		Dihedral angle / degrees				Relative energy / kJ mol <sup>-1</sup>		
Nº	Ether <b>9</b> (R)	A	B	C	D	ΔE <sub>el</sub>	ΔE <sub>ZPE</sub>	ΔE <sub>G</sub>
1	Ph <sup>+</sup> TG <sup>-</sup> G <sup>-</sup>	22.3	179.1	-83.6	-59.6	0.00	0.00	0.00
2	Ph <sup>+</sup> TTG <sup>-</sup>	28.3	177.6	-158.0	-58.9	0.14	0.04	0.53
3	Ph <sup>-</sup> TTG <sup>-</sup>	-27.0	-176.9	-163.3	-59.6	0.21	0.37	0.83
4	Ph <sup>-</sup> TG <sup>-</sup> G <sup>+</sup>	-20.5	-175.8	-82.6	-59.4	0.34	0.47	0.99
5	Ph <sup>-</sup> TG <sup>-</sup> G <sup>+</sup>	-24.9	-176.3	-81.2	60.7	0.82	0.63	1.05
6	Ph <sup>+</sup> TG <sup>-</sup> G <sup>+</sup>	21.9	177.8	-80.7	60.7	1.39	1.23	0.73
7	Ph <sup>+</sup> TTG <sup>+</sup>	27.6	176.8	-155.6	60.9	2.01	1.81	1.65
8	Ph <sup>-</sup> TTG <sup>+</sup>	-26.4	-178.8	-158.6	60.9	2.42	2.27	2.28
9	Ph <sup>+</sup> TG <sup>+</sup> G <sup>+</sup>	28.7	170.2	68.0	60.8	11.63	11.55	12.57
10	Ph <sup>-</sup> TG <sup>+</sup> G <sup>+</sup>	-28.2	179.6	65.6	61.0	12.22	11.86	12.27
11	Ph <sup>+</sup> TG <sup>+</sup> G <sup>-</sup>	30.5	160.5	77.4	-52.9	17.00	16.65	17.25
12	Ph <sup>-</sup> TG <sup>+</sup> G <sup>-</sup>	-28.3	172.7	71.9	-52.3	18.11	17.58	16.85
Nº	Tetrazolone <b>16</b> (R)	A		C	D	ΔE <sub>el</sub>	ΔE <sub>ZPE</sub>	ΔE <sub>G</sub>
1	CG <sup>+</sup>	-0.7		5.0	60.8	-69.31	-68.22	-68.35
2	CG <sup>-</sup>	0.1		9.9	-59.2	-68.08	-66.70	-66.66
3	TG <sup>+</sup>	1.7		178.7	60.9	-65.43	-64.54	-64.84
4	TG <sup>-</sup>	3.6		-171.5	-57.5	-62.46	-61.06	-61.25
Distance / pm								
Nº	TS type	OC(1)	OC(3)	NC(1)	NC(3)	ΔE <sub>el</sub>	ΔE <sub>ZPE</sub>	ΔE <sub>G</sub>
1	1,3'-shift	273.7	338.0	270.5	394.1	161.89	150.32	149.67
2	1,3'-shift	266.1	400.0	267.8	332.8	160.49	149.06	148.31
3	3,3'-shift	230.5	331.6	339.5	244.8	115.41	107.50	110.76
4	3,3'-shift	224.0	328.3	333.6	239.1	107.37	100.06	102.77

Definition of the geometric parameters (see also Figure 1 for the graphical representation). Compound “Ether” (R-enantiomer): A is defined as CCNC dihedral angle; B is defined as N–COC dihedral angle; C is defined as COC(1)C(6); D is defined as C(1)C(6)C(5)C(4) dihedral angle. Compound “Tetrazolone” resulting from 1,3'-shift (R-enantiomer): A is defined as CCNC dihedral angle; dihedral angle “B” (analogous of ether) is not applicable; C is defined as C–NC(1)H dihedral angle; D is defined as C(1)C(6)C(5)C(4). By the nature of compound, the structures resulting from the 3,3'-shift represent S-enantiomeric variety and otherwise are symmetrically identical to R-structures resulting from the 1,3'-shift. “TS” stands for transition state. Numbering of the carbon atoms corresponds to that of the starting compound. Nitrogen atom is N(5) of the ether. All relative energies are calculated with respect to Ph<sup>+</sup>TG<sup>-</sup>G<sup>-</sup> conformer (Nº1) of Ether **9**. ΔE<sub>el</sub>, ΔE<sub>ZPE</sub>, ΔE<sub>G</sub> state for the relative electronic, zero-point-corrected and Gibbs free energy (at 298.15 K), respectively. The absolute values calculated for the most stable conformer Ph<sup>+</sup>TG<sup>-</sup>G<sup>-</sup> of Ether **9** at the DFT(B3LYP)/6-31G(d,p) level are: E<sub>el</sub>=-797.995940; E<sub>ZPE</sub>=-797.736523, E<sub>G</sub>=-797.780470 hartree.

**Table 2S.** Conformationally relevant geometric parameters and relative energies of conformers of 5-(Cyclohex-2-enyloxy)-1-phenyl-1*H*-tetrazole **9** [in brief: Ether **9** (R)] and corresponding isomeric 1-(cyclohex-2-enyl)-4-phenyltetrazol-5-one **16** [in brief: Tetrazolone **16** (R)] calculated at the MP2/6-31G(d,p) level of theory.

Compound / Conformer		Dihedral angle / degrees				Relative Energy
Nº	Ether <b>9</b> (R)	A	B	C	D	$\Delta E_{el}$
1	Ph <sup>+</sup> TTG <sup>-</sup>	36.6	-180.0	-164.9	-62.2	0.00
2	Ph <sup>-</sup> TTG <sup>-</sup>	-35.2	-173.7	-168.4	-62.6	0.06
3	Ph <sup>+</sup> TG <sup>-</sup> G <sup>-</sup>	33.7	174.4	-78.9	-62.8	0.96
4	Ph <sup>+</sup> TG <sup>-</sup> G <sup>-</sup>	-34.1	-177.1	-79.9	-62.6	1.32
5	Ph <sup>+</sup> TG <sup>-</sup> G <sup>+</sup>	-34.6	-177.0	-78.2	63.6	3.90
6	Ph <sup>+</sup> TG <sup>-</sup> G <sup>+</sup>	34.8	175.9	-76.7	63.6	4.31
7	Ph <sup>+</sup> TTG <sup>+</sup>	35.6	177.4	-161.4	63.9	4.32
8	Ph <sup>-</sup> TTG <sup>+</sup>	-35.1	-177.7	-163.1	63.8	4.64
9	Ph <sup>+</sup> TG <sup>+</sup> G <sup>+</sup>	36.8	167.0	71.1	63.4	11.16
10	Ph <sup>-</sup> TG <sup>+</sup> G <sup>+</sup>	-37.1	177.6	67.4	63.5	11.88
11	Ph <sup>+</sup> A <sup>+</sup> G <sup>+</sup> G <sup>-</sup>	38.6	91.6	104.6	-60.4	13.03
12	Ph <sup>-</sup> A <sup>+</sup> G <sup>+</sup> G <sup>-</sup>	-39.4	157.3	86.4	-56.4	19.18
Nº	Tetrazolone <b>16</b> (R)	A	B	C	D	$\Delta E_{el}$
1	Ph <sup>-</sup> CG <sup>+</sup>	-22.2	-25.7	91.4	62.7	-67.54
2	Ph <sup>+</sup> CG <sup>+</sup>	22.3	-26.7	90.3	62.8	-67.52
3	Ph <sup>-</sup> CG <sup>-</sup>	-22.9	-9.6	108.1	-63.6	-65.74
4	Ph <sup>+</sup> CG <sup>-</sup>	23.1	-10.5	107.2	-63.6	-65.74
5	Ph <sup>+</sup> TG <sup>+</sup>	23.8	179.8	61.7	63.7	-63.52
6	Ph <sup>-</sup> TG <sup>+</sup>	-23.5	-179.9	62.0	63.7	-63.38
7	Ph <sup>+</sup> TG <sup>-</sup>	25.6	-164.5	78.5	-60.8	-62.16
8	Ph <sup>-</sup> TG <sup>-</sup>	-25.3	-164.2	78.8	-60.8	-61.97
Distance / pm						
Nº	TS type	OC(1)	OC(3)	NC(1)	NC(3)	$\Delta E_{el}$
1	1,3'-shift	271.4	312.5	264.7	376.6	213.87
2	1,3'-shift	261.3	377.6	267.5	311.4	209.60
3	3,3'-shift	204.6	310.8	319.8	213.9	128.27
4	3,3'-shift	215.7	317.0	328.2	220.8	121.62

For the definition of the geometric parameters see caption of Table 1S and also Figure 1 for the graphical representation. All relative electronic energies ( $\Delta E_{el}$ , in  $\text{kJ mol}^{-1}$ ) are calculated with respect to Ph<sup>+</sup>TTG<sup>-</sup> conformer (Nº1) of Ether **9**. The absolute value calculated for Ph<sup>+</sup>TTG<sup>-</sup> conformer at the MP2/6-31G(d,p) level is  $E_{el} = -795.620869$  hartree.

**Table 3S.** Conformationally relevant geometric parameters and relative energies of conformers of (E)-5-(3,7-dimethylocta-2,6-dienyloxy)-1-phenyl-1*H*-tetrazole (**14**) [in brief: Ether **14**] and corresponding isomeric 1-(3,7-dimethylocta-1,6-dien-3-yl)-4-phenyl-1*H*-tetrazol-5(4*H*)-one (**15**) [in brief: Tetrazolone **15** (R)] and isomeric (E)-1-(3,7-dimethylocta-2,6-dienyl)-4-phenyl-1*H*-tetrazol-5(4*H*)-one (**18**) [in brief: Tetrazolone **18**] calculated at the DFT(B3LYP)/6-31G(d,p) level of theory for selected structures.

Compound / Form		Dihedral angle / degrees								Relative energy / kJ mol <sup>-1</sup>		
No.	Ether <b>14</b>	A	B	C	O12=3	12=34	2=345	3456	456=7	ΔE <sub>el</sub>	ΔE <sub>ZPE</sub>	ΔE <sub>G</sub>
1	Ph <sup>+</sup> TG <sup>-</sup> A <sup>+</sup> CA <sup>-</sup> TA <sup>-</sup>	27.4	171.7	-82.9	131.8	-0.7	-104.5	-173.6	-111.7	0.00	0.00	0.00
2	Ph <sup>+</sup> TG <sup>+</sup> A <sup>-</sup> CA <sup>-</sup> TA <sup>+</sup>	-26.8	-171.5	84.2	-120.5	0.3	-89.2	-178.5	119.8	1.45	1.17	-0.98
3	Ph <sup>+</sup> TG <sup>-</sup> A <sup>+</sup> TA <sup>-</sup> TA <sup>+</sup>	26.1	171.3	-85.1	119.6	177.2	-115.0	176.2	115.5	1.50	1.20	-0.95
4	Ph <sup>-</sup> TG <sup>-</sup> A <sup>+</sup> TA <sup>-</sup> TA <sup>+</sup>	-25.8	175.9	-87.2	118.9	177.1	-114.5	176.4	116.8	1.64	1.53	-1.53
5	Ph <sup>+</sup> TG <sup>-</sup> A <sup>+</sup> TA <sup>+</sup> TA <sup>+</sup>	25.4	171.1	-85.1	118.1	178.7	103.0	175.0	115.2	1.91	1.56	-1.50
6	Ph <sup>-</sup> TG <sup>-</sup> A <sup>+</sup> TA <sup>+</sup> TA <sup>+</sup>	-25.0	175.9	-86.5	118.9	178.2	105.6	175.8	118.9	2.11	1.58	-1.89
7	Ph <sup>+</sup> TG <sup>+</sup> A <sup>-</sup> CA <sup>-</sup> TA <sup>+</sup>	26.0	-176.2	86.8	-120.0	0.3	-90.8	-178.8	119.0	1.49	1.69	0.87
8	Ph <sup>-</sup> TG <sup>-</sup> A <sup>+</sup> CA <sup>-</sup> TA <sup>+</sup>	-25.3	176.4	-85.6	117.9	-0.5	-119.4	173.2	112.0	2.53	2.45	0.17
9	Ph <sup>+</sup> T T A <sup>+</sup> CA <sup>-</sup> TA <sup>-</sup>	21.8	177.6	175.1	90.0	-1.8	-95.5	-175.9	-116.8	3.05	2.46	-0.47
10	Ph <sup>+</sup> TG <sup>-</sup> A <sup>+</sup> T C TA <sup>+</sup>	26.3	171.4	-84.9	119.7	177.4	-2.3	177.6	116.0	3.20	2.54	-1.07
11	Ph <sup>+</sup> TG <sup>-</sup> A <sup>+</sup> CA <sup>-</sup> TA <sup>+</sup>	26.5	171.7	-83.9	117.7	-0.7	-120.6	172.9	111.1	2.31	2.60	1.51
12	Ph <sup>-</sup> TG <sup>-</sup> A <sup>+</sup> T C TA <sup>+</sup>	-26.2	176.3	-87.5	119.3	177.1	-2.3	177.9	117.0	3.32	2.65	0.30
13	Ph <sup>+</sup> T T A <sup>+</sup> CA <sup>-</sup> TA <sup>+</sup>	22.2	178.7	176.0	92.0	-1.9	-92.0	179.3	116.7	3.97	3.08	-2.53
14	Ph <sup>-</sup> T T A <sup>+</sup> CA <sup>-</sup> TA <sup>+</sup>	-23.3	-176.6	176.9	109.2	-1.9	-93.0	-180.0	116.4	4.36	3.56	-3.02
15	Ph <sup>-</sup> TG <sup>+</sup> A <sup>+</sup> CA <sup>-</sup> TA <sup>+</sup>	-26.9	-176.0	74.1	89.4	-1.5	-91.0	-179.5	117.3	3.85	3.91	1.56
16	Ph <sup>-</sup> TG <sup>-</sup> A <sup>-</sup> CA <sup>-</sup> TA <sup>+</sup>	-26.2	-178.5	-80.6	-112.9	-0.3	-97.4	179.9	116.7	4.33	4.10	0.01
17	Ph <sup>+</sup> TG <sup>+</sup> A <sup>+</sup> CA <sup>-</sup> TA <sup>+</sup>	26.1	177.1	78.3	101.3	-1.6	-93.8	-179.8	117.7	4.65	4.38	-0.02
18	Ph <sup>+</sup> TG <sup>-</sup> A <sup>-</sup> CA <sup>-</sup> TA <sup>+</sup>	25.2	177.6	-77.5	-108.9	0.0	-98.1	179.6	117.4	4.83	4.50	-0.86
19	Ph <sup>+</sup> TG <sup>+</sup> A <sup>+</sup> CA <sup>-</sup> GA <sup>+</sup>	24.4	176.0	74.5	95.1	-1.6	-124.3	59.4	129.6	7.28	7.75	7.05
No.	Tetrazolone <b>15</b> (R)	A	CN39		1=234	2345	3456	456=7	ΔE <sub>el</sub>	ΔE <sub>ZPE</sub>	ΔE <sub>G</sub>	
1	T A <sup>+</sup> T T A <sup>+</sup>	-2.8			-170.1			132.6	179.6	173.2	116.6	-43.50
2	T C T T A <sup>+</sup>	1.4			-178.1			-4.1	-179.2	172.0	117.3	-43.55
3	G <sup>+</sup> A <sup>+</sup> T T A <sup>+</sup>	0.7			49.4			-120.5	178.9	173.8	114.8	-42.85

4	G <sup>-</sup> A <sup>+</sup> G <sup>+</sup> T A <sup>+</sup>	-2.5	-70.2	93.8	50.8	175.0	112.5	-41.80	-43.56	-42.66	
5	T A <sup>+</sup> T G <sup>+</sup> A <sup>-</sup>	1.2	-172.6	134.2	174.9	71.6	-128.9	-40.00	-41.21	-36.91	
6	T A <sup>+</sup> T G <sup>+</sup> A <sup>+</sup>	-0.3	-172.0	132.5	171.2	62.1	85.4	-37.93	-38.52	-30.80	
7	C A <sup>+</sup> T G <sup>+</sup> A <sup>+</sup>	0.7	14.9	119.6	167.9	59.9	91.0	-31.57	-32.45	-29.90	
8	A <sup>-</sup> A <sup>+</sup> G <sup>+</sup> G <sup>+</sup> A <sup>+</sup>	1.1	-131.9	110.2	47.7	61.9	93.5	-32.18	-32.39	-29.20	
No.	Tetrazolone <b>18</b>	A	CN12	N12=3	12=34	2=345	3456	456=7	$\Delta E_{el}$	$\Delta E_{ZPE}$	$\Delta E_G$
1	A <sup>-</sup> A <sup>+</sup> C A <sup>-</sup> T A <sup>-</sup>	-0.1	-88.8	122.4	-0.5	-96.5	-172.5	-111.2	-78.90	-77.48	-77.40
2	A <sup>-</sup> A <sup>+</sup> C A <sup>-</sup> T A <sup>+</sup>	0.5	-90.1	118.5	-0.6	-99.1	-179.2	113.0	-75.19	-74.53	-78.14
3	A <sup>-</sup> A <sup>+</sup> C A <sup>+</sup> T A <sup>-</sup>	1.5	-89.9	108.3	0.0	92.2	178.9	-117.5	-74.40	-73.54	-77.07
4	A <sup>-</sup> A <sup>+</sup> C A <sup>+</sup> T A <sup>+</sup>	1.7	-90.2	108.3	-0.2	94.4	173.3	117.0	-74.16	-73.34	-78.31
5	A <sup>+</sup> A <sup>-</sup> C A <sup>-</sup> G <sup>+</sup> A <sup>-</sup>	-2.7	89.3	-108.0	0.6	-98.1	74.7	-113.7	-73.59	-72.72	-75.58
6	A <sup>+</sup> A <sup>+</sup> C A <sup>-</sup> T A <sup>+</sup>	1.2	107.6	112.1	-1.4	-96.8	179.6	116.2	-72.86	-72.13	-76.27
7	A <sup>+</sup> A <sup>+</sup> T A <sup>+</sup> T A <sup>+</sup>	1.0	107.4	117.5	179.7	109.0	178.4	116.5	-71.58	-71.58	-79.25
8	A <sup>+</sup> A <sup>+</sup> C A <sup>+</sup> T A <sup>+</sup>	-0.8	107.5	116.0	-0.3	97.8	174.6	117.3	-71.98	-71.51	-76.69
9	A <sup>+</sup> A <sup>+</sup> C A <sup>+</sup> T A <sup>-</sup>	0.7	104.5	116.2	0.0	95.6	179.8	-117.3	-71.68	-70.98	-75.99
10	A <sup>-</sup> A <sup>+</sup> C A <sup>+</sup> G <sup>+</sup> A <sup>+</sup>	0.2	-88.2	109.2	0.4	82.9	56.8	108.7	-71.16	-69.72	-71.25

Definition of the geometric parameters (see also Figure 4 for the graphical representation). Compound “Ether **14**”: dihedral angles A, B, C, are defined similar to the ethers derived from the carbocyclic allylic alcohols **9** and **10** (compare with Figure 1). The numbers from 1 to 7 in the names of the remaining dihedral angles correspond to the numbers of the carbon atoms of the side chain (as shown in Figure 4), e.g. “2=345” is defined as C(2)=C(3)C(4)C(5) dihedral angle. Compound “Tetrazolone **15** (R)” resulting from 3,3'-shift (R-enantiomer): A is defined as CCNC dihedral angle describing the mutual orientation of the phenyl and tetrazolone rings; dihedral angles “B” and “C” (analogous of ether) are not applicable; CN39 is defined as C(tetrazolone)=NC(3)C(9) dihedral angle. Compound “Tetrazolone **18**” resulting from 1,3'-shift: A is defined as CCNC dihedral angle in Tetrazolone **15**; dihedral angles “B” and “C” (analogous of ether) are not applicable; CN12 is defined as C(tetrazolone)=NC(1)C(2) dihedral angle. All relative energies are calculated with respect to Ph<sup>+</sup>TG<sup>-</sup>A<sup>+</sup>CA<sup>-</sup>TA<sup>-</sup> conformer (N°1) of Ether **14**.  $\Delta E_{el}$ ,  $\Delta E_{ZPE}$ ,  $\Delta E_G$  state for the relative electronic, zero-point-corrected and Gibbs free energy (at 298.15 K), respectively. The absolute values calculated for the most stable conformer of Ether **14** at the DFT(B3LYP)/6-31G(d,p) level are:  $E_{el}=-955.237317$ ;  $E_{ZPE}=-954.868515$ ,  $E_G=-954.923266$  hartree.

**Table 4S.** Conformationally relevant geometric parameters and relative energies of conformers of (E)-5-(3,7-dimethylocta-2,6-dienyloxy)-1-phenyl-1*H*-tetrazole (**14**) [in brief: Ether **14**] and corresponding isomeric 1-(3,7-dimethylocta-1,6-dien-3-yl)-4-phenyl-1*H*-tetrazol-5(4*H*)-one (**15**) [in brief: Tetrazolone **15** (R)] and isomeric (E)-1-(3,7-dimethylocta-2,6-dienyl)-4-phenyl-1*H*-tetrazol-5(4*H*)-one (**18**) [in brief: Tetrazolone **18**] calculated at the MP2/6-31G(d,p) level of theory for selected structures.

Compound / Form		Dihedral angle / degrees								Energy
No.	Ether <b>14</b>	A	B	C	O12=3	12=34	2=345	3456	456=7	$\Delta E_{el}$
1	Ph <sup>+</sup> TG <sup>+</sup> G <sup>+</sup> CA <sup>-</sup> G <sup>+</sup> A <sup>+</sup>	32.7	172.8	67.2	60.9	-3.4	-118.6	53.8	138.7	0.0
2	Ph <sup>+</sup> T T G <sup>+</sup> CA <sup>-</sup> TA <sup>-</sup>	31.1	176.1	171.4	73.9	-1.7	-97.7	-175.7	-85.9	10.7
3	Ph <sup>+</sup> T G <sup>-</sup> A <sup>+</sup> CA <sup>-</sup> TA <sup>-</sup>	35.5	162.3	-72.0	118.6	2.3	-123.6	-173.3	-107.7	11.6
4	Ph <sup>+</sup> T G <sup>-</sup> A <sup>+</sup> CA <sup>-</sup> TA <sup>+</sup>	35.7	160.7	-78.4	111.6	-1.2	-118.9	166.3	95.1	13.8
5	Ph <sup>-</sup> T G <sup>-</sup> A <sup>+</sup> CA <sup>-</sup> TA <sup>+</sup>	-36.4	165.8	-81.0	111.2	-1.1	-118.2	166.4	95.0	14.3
6	Ph <sup>+</sup> A <sup>+</sup> G <sup>-</sup> A <sup>+</sup> TA <sup>+</sup> TA <sup>+</sup>	31.4	125.1	-64.2	101.7	-168.0	90.9	179.9	89.8	14.4
7	Ph <sup>-</sup> A <sup>+</sup> G <sup>-</sup> A <sup>+</sup> TA <sup>+</sup> TA <sup>+</sup>	-16.8	122.3	-64.1	98.8	-167.1	100.0	-178.1	110.6	17.2
8	Ph <sup>-</sup> T G <sup>+</sup> A <sup>-</sup> CA <sup>-</sup> TA <sup>+</sup>	-35.7	-164.3	76.5	-120.5	-1.7	-86.1	-178.5	106.0	18.8
9	Ph <sup>+</sup> T G <sup>+</sup> A <sup>-</sup> CA <sup>-</sup> TA <sup>-</sup>	36.3	-170.4	79.2	-120.7	-1.6	-86.0	-178.5	106.0	19.4
10	Ph <sup>+</sup> T G <sup>-</sup> A <sup>+</sup> TA <sup>+</sup> TA <sup>+</sup>	35.7	164.5	-76.1	120.5	177.3	-113.9	177.6	96.3	19.7
11	Ph <sup>-</sup> T G <sup>-</sup> A <sup>+</sup> TA <sup>+</sup> TA <sup>+</sup>	-36.3	170.5	-78.8	120.7	177.2	-113.9	177.6	96.1	20.2
12	Ph <sup>-</sup> T G <sup>-</sup> A <sup>+</sup> TA <sup>+</sup> TA <sup>+</sup>	-36.4	168.0	-75.9	117.3	-177.1	99.5	176.7	101.6	20.7
13	Ph <sup>+</sup> T G <sup>-</sup> A <sup>+</sup> TC TA <sup>+</sup>	35.6	163.5	-75.7	119.9	178.4	-1.2	177.8	101.8	22.7
14	Ph <sup>-</sup> T G <sup>-</sup> A <sup>+</sup> TC TA <sup>+</sup>	-36.4	169.3	-78.3	120.1	178.4	-1.1	177.7	101.7	23.2
No.	Tetrazolone <b>15</b> (R)	A	CN39		1=234	2345	3456	456=7	$\Delta E_{el}$	
1	Ph <sup>+</sup> T A <sup>+</sup> T G <sup>+</sup> A <sup>-</sup>	20.1	-174.9		135.0	-178.4	70.7	-124.3	-70.1	
2	Ph <sup>-</sup> T A <sup>+</sup> T G <sup>+</sup> G <sup>+</sup>	-20.5	-174.6		134.3	177.3	61.8	80.0	-67.1	
3	Ph <sup>+</sup> T A <sup>+</sup> T G <sup>+</sup> G <sup>+</sup>	10.6	-175.2		134.4	175.6	59.4	81.0	-66.4	
4	Ph <sup>+</sup> A <sup>-</sup> A <sup>+</sup> G <sup>+</sup> G <sup>+</sup> A <sup>-</sup>	26.6	-125.2		111.3	50.6	59.0	-124.4	-64.8	
5	Ph <sup>-</sup> A <sup>-</sup> A <sup>+</sup> G <sup>+</sup> G <sup>+</sup> G <sup>+</sup>	-22.1	-129.5		113.2	53.1	56.6	77.2	-60.7	
6	Ph <sup>-</sup> C A <sup>+</sup> T G <sup>+</sup> G <sup>+</sup>	-22.9	-3.9		125.6	173.4	54.6	79.8	-60.2	
7	Ph <sup>-</sup> T A <sup>+</sup> T T A <sup>+</sup>	-25.5	-165.4		128.3	-175.1	173.1	105.3	-50.8	
8	Ph <sup>+</sup> T A <sup>+</sup> T T A <sup>+</sup>	26.0	-164.6		127.1	-175.4	172.2	106.2	-50.7	

9	Ph <sup>-</sup> T C T A <sup>+</sup>	-24.6	-177.2	-1.0	-176.4	171.2	105.0	-49.1	
10	Ph <sup>+</sup> T C T A <sup>+</sup>	24.4	-177.1	-1.2	-176.2	171.4	105.0	-48.9	
11	Ph <sup>-</sup> G <sup>-</sup> G <sup>+</sup> G <sup>+</sup> G <sup>+</sup>	-24.3	-70.7	88.1	51.9	-179.3	89.6	-48.7	
12	Ph <sup>+</sup> G <sup>-</sup> G <sup>+</sup> G <sup>+</sup> G <sup>+</sup>	22.6	-71.1	87.6	52.1	-179.1	89.6	-48.6	
13	Ph <sup>+</sup> G <sup>+</sup> A <sup>-</sup> T A <sup>+</sup>	24.2	52.8	-123.6	178.8	173.8	169.4	-47.9	
14	Ph <sup>-</sup> G <sup>+</sup> A <sup>-</sup> T A <sup>+</sup>	-25.6	52.7	-123.4	178.9	169.5	100.8	-47.8	
No.	Tetrazolone <b>18</b>	A	CN12	N12=3	12=34	2=345	3456	456=7	$\Delta E_{el}$
1	Ph <sup>-</sup> A <sup>+</sup> A <sup>-</sup> C A <sup>-</sup> G <sup>+</sup> A <sup>-</sup>	-23.0	90.0	-105.9	-2.1	-93.1	70.0	-97.4	-63.2
2	Ph <sup>+</sup> A <sup>-</sup> G <sup>+</sup> C A <sup>-</sup> T A <sup>+</sup>	39.0	-110.9	73.9	-1.8	-99.3	173.5	133.5	-62.0
3	Ph <sup>+</sup> A <sup>-</sup> A <sup>+</sup> C A <sup>-</sup> T A <sup>-</sup>	24.7	-92.2	123.0	-0.7	-93.2	-172.1	-99.4	-61.5
4	Ph <sup>+</sup> A <sup>-</sup> A <sup>+</sup> C A <sup>+</sup> T A <sup>+</sup>	24.7	-101.1	99.5	0.5	89.9	180.0	83.9	-57.6
5	Ph <sup>+</sup> A <sup>-</sup> A <sup>+</sup> C A <sup>+</sup> G <sup>+</sup> A <sup>+</sup>	23.3	-87.7	108.0	1.8	81.4	54.6	95.7	-55.1
6	Ph <sup>+</sup> A <sup>-</sup> A <sup>+</sup> C A <sup>+</sup> T A <sup>-</sup>	23.4	-92.2	104.9	1.4	88.3	178.7	-105.9	-53.8
7	Ph <sup>+</sup> G <sup>+</sup> A <sup>+</sup> C A <sup>-</sup> T A <sup>+</sup>	23.0	75.1	106.7	-1.6	-90.5	-178.9	93.4	-51.9
8	Ph <sup>-</sup> G <sup>+</sup> A <sup>+</sup> C A <sup>+</sup> T A <sup>+</sup>	-22.5	68.4	85.8	0.9	92.3	178.9	83.6	-50.8
9	Ph <sup>+</sup> G <sup>+</sup> A <sup>+</sup> C A <sup>+</sup> T A <sup>-</sup>	22.9	80.2	96.9	1.8	91.2	178.9	-101.2	-49.2
10	Ph <sup>+</sup> A <sup>+</sup> A <sup>+</sup> T A <sup>+</sup> T A <sup>+</sup>	23.4	107.7	90.9	-176.2	112.4	-179.4	100.6	-48.1

For the definition of the geometric parameters see caption of Table 3S and also Figure 4 for the graphical representation. All relative electronic energies ( $\Delta E_{el}$ , in kJ mol<sup>-1</sup>) are calculated with respect to Ph<sup>+</sup>TG<sup>+</sup>G<sup>+</sup>CA<sup>-</sup>G<sup>+</sup>A<sup>+</sup> conformer (N°1) of Ether **14**. The absolute value calculated for Ph<sup>+</sup>TG<sup>+</sup>G<sup>+</sup>CA<sup>-</sup>G<sup>+</sup>A<sup>+</sup> conformer at the MP2/6-31G(d,p) level is  $E_{el}=-952.323974$  hartree.

This Section of Supplementary information contains Cartesian coordinates for optimized structures of the transition states for the sigmatropic isomerisation of 5-allyloxytetrazoles to the corresponding tetrazolones. For brevity, before each structure, the theoretical methods are specified briefly as DFT or MP2, which correspond to DFT(B3LYP)/6-31G(d,p) and MP2/6-31G(d,p), respectively. The theory level is followed by a short name of the structure: TS1a, TS1a', TS1b, TS1b', TS2a, TS2a', TS2b, TS2b', TS3a, TS3a', TS3a'', TS3b and TS3b'.

In these names “TS” designates “Transition State”. Numbers 1, 2 and 3 correspond to the tetrazoles derived from the carbocyclic allylic alcohols (1) cyclohex-2-enol; (2) 3-methylcyclohex-2-enol; (3) natural terpene alcohol nerol.

Some names finish with one or two apostrophes, which designate transition states occurring on alternative “parallel” pathways. These structures have relative energies slightly higher than the base structure (without apostrophe). The calculated electronic energies (in hartree) and total dipole moments (debye) are also shown.

DFT, TS1a, -797.93481127 hartree, 7.7 debye

N	0.80257800	1.92563300	-0.04402300
N	-0.47814500	1.97239800	-0.13656600
N	-1.02000200	0.74216900	-0.22650200
C	-0.00998300	-0.16931700	-0.19803100
N	1.14675600	0.58658900	-0.07568000
C	2.49971700	0.17943400	-0.01058800
C	3.50476300	1.15147300	0.09605800
C	4.83796600	0.75503200	0.16159600
C	5.18330200	-0.59746600	0.12187200
C	4.17532400	-1.55591600	0.01532500
C	2.83382300	-1.18111900	-0.05156500
H	3.22862300	2.19782100	0.12528600
H	5.61186700	1.51292400	0.24373400
H	6.22512200	-0.89940800	0.17264900
H	4.42994800	-2.61158900	-0.01749100
H	2.04383600	-1.91639800	-0.13633000
O	-0.11580300	-1.41770400	-0.26978400
C	-2.75530200	-1.26885200	-0.57084800
C	-2.92029900	-1.45559100	0.81195400
C	-3.56082800	-0.48570400	1.53670500
C	-4.02201300	0.76650000	0.89063700
C	-4.57671800	0.49813200	-0.51866200
C	-3.59523000	-0.33330100	-1.36552000
H	-2.07931000	-1.92875000	-1.10034200
H	-2.42181900	-2.28807900	1.29581400
H	-4.73831100	1.30683200	1.51534800
H	-3.10726400	1.38666600	0.80297200
H	-5.52645400	-0.04145400	-0.42457800
H	-4.79482400	1.44494500	-1.01927000
H	-2.91411800	0.29703100	-1.94465700
H	-4.13211500	-0.94475000	-2.10791400
H	-3.62419700	-0.58152100	2.61884100

DFT, TS1a', -797.93427933 hartree, 8.6 debye

N	-1.41384700	2.31525400	0.00450300
N	-0.23227900	2.82370500	-0.05032500
N	0.71852200	1.87674600	-0.11240100
C	0.10658600	0.66522000	-0.12705600
N	-1.24900300	0.94411000	-0.02522200
C	-2.36486000	0.07595900	0.00268700
C	-3.65182900	0.61717900	0.13601500
C	-4.75385600	-0.23315100	0.16883000
C	-4.59172800	-1.61675200	0.07051100
C	-3.30743400	-2.14474400	-0.06211500
C	-2.19013300	-1.31111200	-0.09806500
H	-3.76895200	1.69075100	0.21108200
H	-5.74772500	0.19291300	0.27231400
H	-5.45593700	-2.27370000	0.09590600
H	-3.16681200	-3.21921600	-0.14115200
H	-1.18949500	-1.70959100	-0.20334900
O	0.65688600	-0.46351200	-0.22016600
C	3.09952100	0.70177600	-0.62980200
C	3.42013900	0.81564800	0.73442300
C	3.57535700	-0.32199800	1.47921900
C	3.34373100	-1.66405000	0.88447700
C	3.83209200	-1.71734000	-0.57370400
C	3.25686400	-0.55514400	-1.40018900
H	2.82189100	1.60203800	-1.16444300
H	3.42725300	1.79770100	1.19358700
H	3.79035000	-2.45572800	1.49244900
H	2.24581800	-1.79666100	0.90203700
H	4.92769500	-1.66442100	-0.58050700
H	3.55535300	-2.67321300	-1.02629100
H	2.25381000	-0.79276200	-1.78024100
H	3.87371400	-0.33817200	-2.28445400
H	3.76750000	-0.23950900	2.54713200

DFT, TS1b, -797.95473353 hartree, 5.2 debye

N	0.89407200	2.17726300	-0.17393700
N	-0.34542100	2.48115000	-0.35645600
N	-1.08874900	1.38186200	-0.63944900
C	-0.25378000	0.30815300	-0.59786300
N	1.00440800	0.82254400	-0.29588100
C	2.25128400	0.16937800	-0.11846800
C	3.33197500	0.89030100	0.40676200
C	4.55883400	0.25496100	0.58019600
C	4.71775200	-1.09074000	0.24282800
C	3.63428000	-1.79870000	-0.27729500
C	2.39880100	-1.17965500	-0.46528500
H	3.19940600	1.93332000	0.66551200
H	5.39453000	0.81811500	0.98503500
H	5.67640100	-1.58078500	0.38292000
H	3.74680300	-2.84470600	-0.54734800
H	1.55453700	-1.72238400	-0.86904000
O	-0.56316300	-0.90604100	-0.73144600
C	-2.14080300	-1.45801100	0.76055200
C	-2.17034500	-0.24856100	1.44943000
C	-2.84820000	0.81662900	0.87830300
C	-3.92809700	0.60921400	-0.14335800
C	-4.38507900	-0.85500200	-0.20031000
C	-3.18724800	-1.81375500	-0.25406500
H	-1.47584100	-2.24447300	1.10060300
H	-1.45632100	-0.06657100	2.24670300
H	-4.77029200	1.27233700	0.08948600
H	-3.53202900	0.94276200	-1.10889800
H	-4.97968800	-1.07811100	0.69411900
H	-5.03679400	-1.01504100	-1.06452500
H	-2.70501800	-1.79303200	-1.23849600
H	-3.50964100	-2.84919100	-0.08850000
H	-2.72717100	1.81107600	1.29550400

DFT, TS1b', -797.95167216 hartree, 5.8 debye

N	-0.25351300	2.14655400	0.32602800
N	0.93772000	2.48635600	-0.05218200
N	1.45061900	1.60842300	-0.94460800
C	0.51599300	0.62755900	-1.12657900
N	-0.56433100	0.98493000	-0.30931300
C	-1.82175200	0.35560800	-0.11298500
C	-2.65460200	0.79824800	0.92371700
C	-3.89026600	0.18629100	1.11765700
C	-4.30164200	-0.86466300	0.29513900
C	-3.46429800	-1.29740700	-0.73300600
C	-2.22510400	-0.69456400	-0.94873200
H	-2.32970100	1.61615400	1.55457100
H	-4.53413100	0.53524500	1.91956900
H	-5.26554800	-1.33853600	0.45321400
H	-3.77561000	-2.10963600	-1.38335200
H	-1.57226700	-1.02261800	-1.74670200
O	0.62318200	-0.42219700	-1.80462400
C	2.14491100	-1.70088000	-0.63722400
C	3.12521200	-0.72657000	-0.75722900
C	3.18015100	0.29226700	0.18158200
C	2.58264500	0.15111300	1.54899600
C	2.16397800	-1.29675700	1.84618600
C	1.42368700	-1.93110400	0.65866000
H	2.00330600	-2.41607300	-1.43931400
H	3.64809300	-0.61210800	-1.70026500
H	3.31699700	0.49900700	2.28677400
H	1.73923200	0.85050700	1.64635800
H	3.06486600	-1.88704500	2.05286800
H	1.54170100	-1.33706900	2.74495300
H	0.40825100	-1.52430800	0.56864100
H	1.29317800	-3.00891000	0.81225800
H	3.80923600	1.15426600	-0.00914100

MP2, TS1a, -795.54103565 hartree, 9.0 debye

N	0.75323400	1.90374800	-0.38207600
N	-0.55928200	1.94111000	-0.54112700
N	-1.08146100	0.69039000	-0.54089600
C	-0.06642500	-0.20146900	-0.37716700
N	1.08099200	0.58463100	-0.27567000
C	2.42521300	0.19010100	-0.08280900
C	3.37656600	1.15612700	0.26937800
C	4.70227400	0.76902500	0.45105600
C	5.08099800	-0.56725100	0.30009900
C	4.11992000	-1.51874700	-0.04324100
C	2.78842400	-1.15249900	-0.24229900
H	3.06890400	2.18519300	0.38352000
H	5.43971300	1.51542500	0.71866700
H	6.11218500	-0.86147600	0.44794400
H	4.40457300	-2.55652800	-0.16478000
H	2.03069300	-1.87556300	-0.50347500
O	-0.14999600	-1.46167600	-0.30684400
C	-2.76140200	-1.38312000	-0.36294800
C	-2.62735300	-1.30360400	1.02642400
C	-3.11203500	-0.18080600	1.65405300
C	-3.78130600	0.90205000	0.90464200
C	-4.58877000	0.33428900	-0.26045000
C	-3.71400700	-0.55811100	-1.13670700
H	-2.18435000	-2.13313700	-0.88518100
H	-2.00140400	-2.01664700	1.54055400
H	-4.38997900	1.51845000	1.56517800
H	-2.95320400	1.51996000	0.51596200
H	-5.41890600	-0.25261500	0.14038300
H	-5.01666700	1.14111700	-0.85367100
H	-3.09682400	0.02643200	-1.82311000
H	-4.30996400	-1.23733100	-1.75586000
H	-2.89177800	-0.03493000	2.70681500

MP2, TS1a', -795.53941160 hartree, 10.2 debye

N	-1.35155200	2.32906000	-0.02859100
N	-0.14476900	2.86016800	-0.12657800
N	0.79159300	1.89822100	-0.31216900
C	0.16771300	0.69307900	-0.33933700
N	-1.18152700	0.98229900	-0.15571500
C	-2.28516500	0.10423300	-0.07011400
C	-3.48615400	0.56901300	0.48057200
C	-4.57792400	-0.29304600	0.55486700
C	-4.47601300	-1.61124100	0.10331300
C	-3.27008100	-2.06196000	-0.43519700
C	-2.16825000	-1.21178000	-0.53301400
H	-3.54829900	1.59054200	0.82641900
H	-5.50889500	0.06620300	0.97525800
H	-5.32680500	-2.27722400	0.16970100
H	-3.18393400	-3.08094900	-0.79187200
H	-1.22664600	-1.54226400	-0.94414800
O	0.69322400	-0.45318600	-0.49062200
C	3.14852800	0.70205000	-0.44670900
C	3.08203400	0.73321100	0.95100300
C	2.99956900	-0.46337500	1.61860600
C	3.00830600	-1.76177500	0.90583100
C	3.88495700	-1.68721400	-0.34212600
C	3.46075900	-0.51463600	-1.22000200
H	3.00332800	1.63252300	-0.97794800
H	2.93712500	1.67701700	1.45413100
H	3.31173900	-2.56833400	1.57250100
H	1.95945600	-1.92104700	0.61469500
H	4.92557600	-1.55304600	-0.03583000
H	3.82800300	-2.61914700	-0.90303400
H	2.52763200	-0.73113100	-1.75258800
H	4.20973600	-0.26201900	-1.97612600
H	2.82028500	-0.45322700	2.68904000

MP2, TS1b, -795.57454659 hartree, 6.0 debye

N	-0.06026300	2.13455100	0.34940800
N	1.19081100	2.42634000	-0.00707000
N	1.65810100	1.51187700	-0.89788100
C	0.64906700	0.60519200	-1.13289600
N	-0.40552900	1.01217900	-0.30451500
C	-1.68773700	0.43060800	-0.13005200
C	-2.38552000	0.67885500	1.05708400
C	-3.64713900	0.11299700	1.22710600
C	-4.19152700	-0.71476200	0.24183400
C	-3.47418300	-0.96263900	-0.92959700
C	-2.21681700	-0.39232400	-1.12983000
H	-1.94884700	1.32249500	1.80761100
H	-4.19842900	0.30866500	2.13779700
H	-5.16875000	-1.15738300	0.38413100
H	-3.89376600	-1.59899200	-1.69811800
H	-1.65148400	-0.57444000	-2.03130700
O	0.69460500	-0.43975500	-1.83475300
C	1.92670000	-1.74879200	-0.64326000
C	2.99585600	-0.86790300	-0.64021200
C	2.98135700	0.18554900	0.26990400
C	2.25654300	0.06039800	1.57989400
C	1.77912400	-1.37052500	1.81564700
C	1.06862100	-1.91371800	0.57835500
H	1.85834000	-2.49593600	-1.42337000
H	3.64489600	-0.81635100	-1.50339400
H	2.93511500	0.37576700	2.37624000
H	1.42075600	0.76725100	1.62396600
H	2.64380100	-2.00266800	2.03076000
H	1.11796900	-1.41224200	2.68221900
H	0.11670700	-1.39578200	0.43417800
H	0.82157400	-2.97002500	0.70314500
H	3.73737000	0.95545200	0.19069500

MP2, TS1b', -795.57201317 hartree, 4.9 debye

N	0.80281800	2.19570400	-0.21368500
N	-0.47474700	2.49579900	-0.40753100
N	-1.18169600	1.36851800	-0.70163200
C	-0.31792900	0.31358600	-0.67290100
N	0.91742900	0.85842100	-0.34235100
C	2.15918800	0.20363000	-0.14281900
C	3.12674700	0.81866100	0.65788900
C	4.34950600	0.17836600	0.84700300
C	4.59388300	-1.06863000	0.26627400
C	3.61111800	-1.67221700	-0.51997800
C	2.38720900	-1.03999700	-0.73925400
H	2.91725500	1.78115500	1.10243400
H	5.10719600	0.65060200	1.45887300
H	5.54288500	-1.56417600	0.42436200
H	3.79771200	-2.63589800	-0.97614600
H	1.61913700	-1.49506200	-1.34629700
O	-0.60822200	-0.92076200	-0.75015800
C	-1.86733700	-1.37191700	0.79779000
C	-1.97646900	-0.12072700	1.40563300
C	-2.69323200	0.85332200	0.72162200
C	-3.80245200	0.45460900	-0.21828900
C	-4.21205800	-0.99566000	0.02100800
C	-2.99423700	-1.91270700	-0.04085100
H	-1.17120500	-2.09078800	1.21342900
H	-1.25625700	0.17738400	2.15718800
H	-4.65005700	1.12688400	-0.07013100
H	-3.46420700	0.59315600	-1.24544900
H	-4.67102600	-1.07768600	1.00884200
H	-4.95780300	-1.30499800	-0.71235300
H	-2.64352500	-2.02487500	-1.06815100
H	-3.24977300	-2.91213400	0.32018300
H	-2.68480200	1.87151200	1.09211800

DFT, TS2a, -837.26441743 hartree, 8.8 debye

N	1.06105700	1.92976500	0.26993900
N	-0.20804900	2.03400400	0.09476300
N	-0.75515300	0.87751500	-0.33187600
C	0.24092400	-0.04475700	-0.44683400
N	1.39373100	0.62917300	-0.05896100
C	2.73106900	0.17845500	0.01617200
C	3.73268000	1.06430200	0.43993800
C	5.05110900	0.62333000	0.51845800
C	5.38667300	-0.68932700	0.18001200
C	4.38294400	-1.56195200	-0.24105000
C	3.05613000	-1.14148600	-0.32690400
H	3.46468100	2.08046100	0.69995300
H	5.82157300	1.31511500	0.84728600
H	6.41713600	-1.02608900	0.24303600
H	4.62955700	-2.58566800	-0.50892300
H	2.26907000	-1.80899700	-0.65359600
O	0.13552300	-1.23790800	-0.81737500
C	-2.54122300	-0.96579900	-1.17258900
C	-2.75144200	-1.43092200	0.12639300
C	-3.41120700	-0.63505900	1.04123200
C	-3.85230700	0.72633300	0.62176200
C	-4.37163100	0.73873500	-0.82378700
C	-3.33541200	0.13323600	-1.78337100
H	-1.83067800	-1.49849000	-1.79382700
H	-2.25386100	-2.34097300	0.44507300
H	-4.58244500	1.14410600	1.32136800
H	-2.93380600	1.34234900	0.67533400
H	-5.30324500	0.16206600	-0.87338900
H	-4.61221000	1.76196700	-1.12410700
H	-2.61780400	0.88186400	-2.13384000
H	-3.81336000	-0.27043000	-2.68867200
C	-3.55088600	-1.03179900	2.47001500
H	-3.03387900	-0.30027300	3.10549900
H	-4.60608400	-1.00250700	2.77039500
H	-3.14446300	-2.02416800	2.67367000

DFT, TS2a', -837.26395925 hartree, 9.8 debye

N	-1.68982700	2.32227200	0.06953300
N	-0.52727300	2.85938500	-0.06213000
N	0.43300400	1.94018000	-0.26905600
C	-0.15643800	0.71799600	-0.29465000
N	-1.50435900	0.95963000	-0.05844600
C	-2.59606100	0.06639500	0.02295400
C	-3.87320800	0.56712800	0.31656500
C	-4.95200300	-0.30895400	0.40177900
C	-4.77731900	-1.67965700	0.19884900
C	-3.50360100	-2.16765800	-0.09338800
C	-2.40973100	-1.30762900	-0.18478600
H	-4.00048900	1.63093500	0.47169700
H	-5.93785300	0.08666500	0.62931800
H	-5.62347800	-2.35690100	0.26629200
H	-3.35333800	-3.23143600	-0.25638600
H	-1.41772700	-1.67397300	-0.41447200
O	0.39774500	-0.39609900	-0.49332000
C	2.83683200	0.80800400	-0.97727000
C	3.22409900	0.82615700	0.36410300
C	3.41957200	-0.35844100	1.04368600
C	3.15152800	-1.64700500	0.33759600
C	3.58457300	-1.59129800	-1.13602600
C	2.94628800	-0.38881900	-1.84619600
H	2.51209400	1.74115900	-1.42261800
H	3.23927900	1.77341800	0.89287600
H	3.60689100	-2.49111300	0.86449400
H	2.05319800	-1.76285800	0.38131000
H	4.67787900	-1.51340400	-1.18608300
H	3.30683000	-2.52021000	-1.64139100
H	1.91835500	-0.61351500	-2.16220200
H	3.49575600	-0.10883100	-2.75608600
C	3.74964200	-0.38592700	2.49564000
H	2.96209400	-0.92088600	3.04313200
H	4.67352000	-0.95502700	2.66216200
H	3.85917100	0.61276700	2.92207800

DFT, TS2b, -837.28137234 hartree, 5.5 debye

N	0.96944900	2.02639500	-0.44243900
N	-0.28357600	2.21201200	-0.68312400
N	-0.93979500	1.03755000	-0.83980400
C	-0.03451200	0.03356800	-0.65100000
N	1.17965000	0.67788100	-0.39959700
C	2.46760500	0.14713500	-0.13562700
C	3.51789300	1.02066700	0.17889800
C	4.78450600	0.50390000	0.43831900
C	5.01630700	-0.87237800	0.39163500
C	3.96364400	-1.73205600	0.07813700
C	2.68802200	-1.23558500	-0.18881700
H	3.33048600	2.08643700	0.21224900
H	5.59503800	1.18547200	0.67950900
H	6.00636300	-1.26860100	0.59578800
H	4.13133800	-2.80443800	0.03529600
H	1.86640000	-1.89486100	-0.43488900
O	-0.25171000	-1.20018000	-0.65308400
C	-2.04969500	-1.61548200	0.87816600
C	-2.15043800	-0.32249500	1.34608400
C	-2.83253000	0.64738600	0.59857400
C	-3.77767500	0.21829300	-0.49684000
C	-4.17622600	-1.25984200	-0.39468300
C	-2.94568400	-2.15237800	-0.19281300
H	-1.38541800	-2.30858100	1.38389400
H	-1.52555600	0.00048500	2.17395500
H	-4.66413300	0.86385600	-0.47343800
H	-3.26914700	0.42283100	-1.44495600
H	-4.85869400	-1.39264900	0.45420900
H	-4.72338800	-1.56174600	-1.29303500
H	-2.34301600	-2.21104100	-1.10819100
H	-3.23916400	-3.17941700	0.05559600
C	-2.89967200	2.06804400	1.05432800
H	-2.90087300	2.75182300	0.20284200
H	-3.84394900	2.20963300	1.60023800
H	-2.07196500	2.33272500	1.71418400

DFT, TS2b', -837.27860054 hartree, 5.7 debye

N	0.38022600	-2.04148500	0.32097400
N	-0.83304300	-2.27637800	-0.06344300
N	-1.26516600	-1.35838200	-0.95627400
C	-0.24960300	-0.45630700	-1.13954800
N	0.79488200	-0.91192600	-0.31712600
C	2.09963700	-0.39395200	-0.11273500
C	2.93261900	-0.98882600	0.84542600
C	4.21323000	-0.48168000	1.04925500
C	4.67242000	0.61262100	0.31298200
C	3.83572600	1.19575100	-0.63838300
C	2.55049300	0.70180400	-0.86176300
H	2.57100600	-1.83906500	1.40981000
H	4.85553400	-0.94821400	1.79046300
H	5.67185800	1.00331900	0.47802700
H	4.18290800	2.04359300	-1.22183500
H	1.89716400	1.14393300	-1.60220800
O	-0.26523500	0.58569500	-1.82406300
C	-1.89253400	1.95112900	-0.58543400
C	-2.85653000	0.98657000	-0.75880700
C	-2.99006300	-0.06647700	0.15943000
C	-2.38048700	0.05156800	1.53305600
C	-1.93340100	1.48012200	1.87392400
C	-1.16472300	2.12450300	0.71274300
H	-1.71435300	2.66968100	-1.37780700
H	-3.36823700	0.91057200	-1.71273000
H	-3.11415400	-0.30548300	2.26691000
H	-1.54599900	-0.66047800	1.60504500
H	-2.82098600	2.08813600	2.08752300
H	-1.32219800	1.47661500	2.78131100
H	-0.16534300	1.67986300	0.61076900
H	-0.99455000	3.19137300	0.89870400
C	-3.97703900	-1.16606200	-0.05987100
H	-3.59444100	-2.10857500	0.33926400
H	-4.90091100	-0.92159800	0.48520300
H	-4.21218400	-1.29661500	-1.11651000

MP2, TS2a, -834.74626254 hartree, 9.0 debye

N	-0.16131800	1.30122100	1.40781600
N	1.05079600	0.98257400	1.87593300
N	1.26829500	-0.35038500	1.83657000
C	0.17107200	-0.94185100	1.28518500
N	-0.72972100	0.12263500	1.05576600
C	-1.96425800	0.09221400	0.37383100
C	-2.48660200	1.28164200	-0.15653100
C	-3.70103000	1.24873200	-0.83919300
C	-4.38259600	0.04242400	-1.01828000
C	-3.84933100	-1.13388000	-0.48900800
C	-2.64217600	-1.12380600	0.21085100
H	-1.95876900	2.21047900	0.00759100
H	-4.11048900	2.16838700	-1.23824500
H	-5.32282700	0.02254700	-1.55409800
H	-4.37697900	-2.07117700	-0.61392700
H	-2.21470400	-2.02438400	0.62452300
O	0.00091500	-2.13502500	0.92039300
C	1.14751700	-1.07124400	-1.24339700
C	1.05365900	0.30982600	-1.21284100
C	2.11117800	1.06741500	-0.70573200
C	3.38321400	0.41233800	-0.30157000
C	3.61665600	-0.90736700	-1.02823600
C	2.38676900	-1.79587500	-0.88996500
H	0.27154400	-1.64752400	-1.51443100
H	0.12492600	0.79503300	-1.48250100
H	4.21258400	1.11073000	-0.42605400
H	3.25957100	0.22530700	0.77463800
H	3.81215200	-0.71424300	-2.08663400
H	4.49423000	-1.40775700	-0.62022500
H	2.22749700	-2.11241400	0.15071700
H	2.45821500	-2.70933700	-1.48244300
C	2.00670900	2.53985400	-0.57328400
H	2.33802200	2.82620100	0.42390600
H	2.68376200	2.99769900	-1.30134600
H	0.99288900	2.89580400	-0.72819700

MP2, TS2a', -834.73903699 hartree, 10.3 debye

N	-1.23362300	2.45921400	0.31699700
N	-0.07549900	3.09294800	0.21374200
N	0.84755800	2.33603800	-0.43099000
C	0.26312100	1.15663500	-0.76286200
N	-1.04648900	1.24994300	-0.28249800
C	-2.05247400	0.26317500	-0.26121800
C	-3.10439500	0.37488400	0.65960000
C	-4.09219900	-0.60699800	0.68713100
C	-4.02529800	-1.70870600	-0.17021000
C	-2.96851200	-1.81215100	-1.07583400
C	-1.97593900	-0.83237600	-1.13303100
H	-3.14196500	1.23450800	1.31334200
H	-4.91068100	-0.51572500	1.39020800
H	-4.79381100	-2.47036500	-0.13932000
H	-2.91608800	-2.65631200	-1.75235200
H	-1.15726600	-0.88759500	-1.83462000
O	0.78774700	0.13214100	-1.29893600
C	3.10796900	0.62147200	-0.07930800
C	2.40388800	0.38732000	1.09116000
C	1.77405700	-0.84576700	1.18636100
C	2.17801100	-1.99024000	0.32216300
C	3.57378600	-1.80691300	-0.25489900
C	3.65810400	-0.44929600	-0.93478300
H	3.22014600	1.65359700	-0.39141900
H	2.05965100	1.22492900	1.68047300
H	2.08627800	-2.91869300	0.88998900
H	1.42687200	-2.01423500	-0.47627100
H	4.30703900	-1.85504400	0.55392600
H	3.80630400	-2.60667000	-0.95684900
H	3.05278100	-0.41405600	-1.84682200
H	4.68028200	-0.18300600	-1.21802900
C	0.56381300	-0.99864700	2.02955100
H	-0.29748000	-0.98983300	1.35143600
H	0.56100400	-1.95815700	2.54643700
H	0.44772900	-0.18183000	2.73767000

MP2, TS2b, -834.76908917 hartree, 5.8 debye

N	0.17028500	-2.07420000	0.25558000
N	-1.09295600	-2.25921300	-0.13154000
N	-1.47774400	-1.28121700	-0.99178200
C	-0.40276100	-0.44428600	-1.18367500
N	0.60709100	-0.95881000	-0.35510600
C	1.92360700	-0.47724900	-0.13945900
C	2.59101800	-0.83739200	1.03678100
C	3.88598500	-0.36897900	1.24708800
C	4.49603600	0.47298100	0.31364200
C	3.80948800	0.83328200	-0.84689300
C	2.51845500	0.36224300	-1.08749100
H	2.10408200	-1.49020000	1.74730400
H	4.41274500	-0.65188200	2.14939700
H	5.49941000	0.83939100	0.48731700
H	4.27915500	1.48126500	-1.57576300
H	1.97587300	0.63198000	-1.98086800
O	-0.35850300	0.61866200	-1.85251500
C	-1.57700900	1.96481200	-0.59521900
C	-2.68290000	1.13944700	-0.64309600
C	-2.78048600	0.05609600	0.23599400
C	-2.03017900	0.09796600	1.54247800
C	-1.46546500	1.48391300	1.84049100
C	-0.71463300	2.02994600	0.63005900
H	-1.45488200	2.72680500	-1.35394100
H	-3.32410100	1.16256500	-1.51485500
H	-2.71817400	-0.21236600	2.33391600
H	-1.23817500	-0.65902600	1.54311700
H	-2.28937200	2.16051800	2.08000400
H	-0.80919800	1.44528300	2.71117100
H	0.20213500	1.45677500	0.46662000
H	-0.40100000	3.06258000	0.79638300
C	-3.93073100	-0.89558700	0.15588500
H	-3.62503300	-1.87871300	0.50869500
H	-4.73322900	-0.53031700	0.80392500
H	-4.30148900	-0.98303300	-0.86187800

MP2, TS2b', -834.76579356 hartree, 5.1 debye

N	0.87139900	2.06979900	-0.50458500
N	-0.41879800	2.25886700	-0.75550500
N	-1.04836500	1.06226800	-0.91245300
C	-0.12240500	0.07430700	-0.74105700
N	1.07122100	0.73755300	-0.46563900
C	2.34666900	0.19494700	-0.16893700
C	3.26177300	0.97128900	0.54957800
C	4.51846200	0.44119700	0.83325700
C	4.84940300	-0.85422900	0.42783400
C	3.91874400	-1.61846500	-0.27783500
C	2.66172000	-1.10052600	-0.59017300
H	2.98612100	1.96948800	0.85855400
H	5.23514600	1.03844600	1.38213800
H	5.82451800	-1.26287000	0.65892800
H	4.17192500	-2.62086200	-0.59848600
H	1.93253000	-1.68025300	-1.13558400
O	-0.32412000	-1.17510800	-0.69153900
C	-1.66620900	-1.51783000	0.89504800
C	-1.89132300	-0.21017000	1.30478300
C	-2.65931700	0.62544500	0.49247500
C	-3.62831700	-0.00738800	-0.48030200
C	-3.96162700	-1.43880400	-0.07194700
C	-2.68363400	-2.25757800	0.07370900
H	-0.93385900	-2.10911000	1.43169900
H	-1.24602400	0.23792100	2.05147900
H	-4.53236900	0.60476900	-0.52377900
H	-3.18254500	0.01094800	-1.47509900
H	-4.49163400	-1.42499300	0.88346900
H	-4.62597900	-1.89581900	-0.80669600
H	-2.25068700	-2.47888000	-0.90330700
H	-2.89309800	-3.21587600	0.55595600
C	-2.90633400	2.05023100	0.87229700
H	-3.05528400	2.66146000	-0.01467300
H	-3.81773800	2.08689900	1.47606200
H	-2.08189900	2.46563100	1.44654200

DFT, TS3a, -955.18072885 hartree, 7.8 debye

N	2.37017500	-1.16957200	-1.70231000
N	1.46656600	-2.08055700	-1.63850800
N	0.91412500	-2.14403000	-0.41332700
C	1.50696900	-1.20943800	0.37976800
N	2.42948100	-0.58832500	-0.44717200
C	3.33976500	0.45852800	-0.17044300
C	4.18929000	0.91645200	-1.18758200
C	5.08321400	1.95006700	-0.92009000
C	5.14207900	2.53292800	0.34754700
C	4.29304200	2.06823800	1.35216900
C	3.39012000	1.03437900	1.10619600
H	4.13799500	0.45870300	-2.16728500
H	5.73907200	2.30035300	-1.71203300
H	5.84214000	3.33815900	0.54902400
H	4.32968300	2.51131700	2.34343800
H	2.72833000	0.66480600	1.87921300
O	1.25736100	-0.97983000	1.58476200
C	-0.49873700	-2.88657300	1.55220100
C	-1.61925800	-2.06773700	1.46422900
C	-2.47621700	-2.07019300	0.37659200
C	-3.60412700	-1.09113400	0.32227200
C	-3.25301800	0.13231400	-0.59020700
H	0.16824400	-2.80921700	2.39845500
H	-1.78281800	-1.33738900	2.25184500
H	-4.49830900	-1.57906400	-0.08543800
H	-3.84527300	-0.72100600	1.32301300
H	-2.99820700	-0.24020200	-1.59023800
C	-2.21945400	-2.92681100	-0.81654400
H	-2.23483100	-3.99125000	-0.55592200
H	-2.94878800	-2.75791800	-1.61033100
H	-1.19984900	-2.71817600	-1.18335900
H	-0.34453200	-3.71059800	0.87184500
H	-2.35315600	0.61202800	-0.19829000
C	-4.40620100	1.09124100	-0.68610800
H	-5.28243400	0.69600900	-1.20193600
C	-4.48020300	2.34545600	-0.21254400
C	-5.73031000	3.16758800	-0.41370600
H	-6.15172100	3.48325400	0.54960500
H	-5.50972800	4.08683600	-0.97147300
H	-6.50100000	2.61621900	-0.95870700
C	-3.37758100	3.05709200	0.53103900
H	-2.48982700	2.44386300	0.69314200
H	-3.06706300	3.95790700	-0.01334000
H	-3.73549700	3.39700700	1.51116900

DFT, TS3a', -955.17940164 hartree, 8.1 debye

N	1.88538300	-1.88034700	0.06183600
N	0.65159900	-1.71157800	-0.25586100
N	0.42151900	-0.46110100	-0.69050800
C	1.58922600	0.23283900	-0.65692600
N	2.51918000	-0.67109100	-0.17482400
C	3.90390100	-0.51162400	0.06689500
C	4.63763300	-1.58565500	0.58983500
C	6.00070500	-1.43150400	0.82951400
C	6.64137800	-0.22139600	0.55532000
C	5.90104800	0.84028500	0.03490800
C	4.53523300	0.70759600	-0.21316400
H	4.13334900	-2.52058700	0.79863300
H	6.56445200	-2.26709900	1.23444100
H	7.70462800	-0.10860200	0.74463600
H	6.38714000	1.78680500	-0.18421600
H	3.95173300	1.52499400	-0.61776700
O	1.75422500	1.42981800	-0.99220000
C	-0.72721800	1.58811100	-1.55840600
C	-1.06660400	2.33374100	-0.42777900
C	-1.95929400	1.90244200	0.53288600
C	-2.61385500	0.55737700	0.48508900
C	-3.89694800	0.53311800	-0.40875100
H	-1.27082200	0.69940400	-1.83619700
H	-0.55869300	3.28222000	-0.27796300
H	-1.89221400	-0.18232600	0.11546100
H	-2.90595100	0.26219000	1.49806700
H	-3.61105100	0.82015800	-1.42906600
C	-2.27622100	2.77578600	1.70434600
H	-3.36171700	2.85283600	1.84419000
H	-1.88552400	2.31288400	2.62090700
H	-1.85266000	3.77803600	1.61417300
H	-0.02858200	1.98735400	-2.27906200
H	-4.59839800	1.29647400	-0.05880300
C	-4.52426500	-0.83205200	-0.42292000
H	-3.88920900	-1.61271200	-0.84259100
C	-5.73695000	-1.19407200	0.02520600
C	-6.19368300	-2.62949800	-0.07509400
H	-6.42850800	-3.03639100	0.91712400
H	-7.11376000	-2.70905000	-0.66868600
H	-5.43573600	-3.26927600	-0.53392300
C	-6.74652000	-0.26141700	0.64710500
H	-6.39693100	0.76901700	0.73286000
H	-7.67432000	-0.25221600	0.06080000
H	-7.02020000	-0.60829600	1.65170300

DFT, TS3a'', -955.17905429 hartree, 7.5 debye

N	-1.32575600	-1.56439200	-0.80763100
N	-0.19357100	-1.14421800	-1.24927800
N	-0.12571100	0.19782800	-1.24441000
C	-1.29977900	0.68555500	-0.77026100
N	-2.06177800	-0.43537200	-0.48910000
C	-3.37616800	-0.53926400	0.02394300
C	-3.94301700	-1.80767700	0.21115400
C	-5.23603100	-1.91173700	0.71768000
C	-5.97095900	-0.76871200	1.03911100
C	-5.39666200	0.48817400	0.84803900
C	-4.10335100	0.61536600	0.34243800
H	-3.36693500	-2.68868800	-0.04179800
H	-5.67052200	-2.89704700	0.86035000
H	-6.97896300	-0.85762300	1.43288000
H	-5.95762600	1.38563200	1.09315900
H	-3.64880800	1.58609300	0.18931800
O	-1.59512300	1.89675300	-0.62360600
C	0.68498700	2.57625900	-1.51867900
C	1.20948600	2.97810000	-0.28883500
C	2.33205600	2.43136000	0.30284100
C	3.03843100	1.22145300	-0.21940500
C	2.78076800	-0.03231500	0.68980000
H	-0.18239300	3.07825700	-1.92103300
H	0.67896800	3.76208500	0.24479700
H	2.71725800	0.97230900	-1.22995800
H	4.11910100	1.40696100	-0.24523200
H	1.70310200	-0.10233400	0.86500900
C	2.83146800	3.00459800	1.59293200
H	2.69198100	2.27981600	2.40572100
H	3.90951600	3.19727600	1.53729100
H	2.31558000	3.92694200	1.86646600
H	1.21663100	1.90510400	-2.17419400
H	3.26777500	0.11808900	1.65763600
C	3.24121000	-1.29580100	0.01647900
H	2.54243200	-1.67939600	-0.72584300
C	4.38161700	-1.97065200	0.23171800
C	4.67926200	-3.23889800	-0.53071000
H	5.61518200	-3.14813100	-1.09816300
H	4.81300400	-4.08700600	0.15373900
H	3.87817000	-3.49006800	-1.23012700
C	5.44822700	-1.56740600	1.21994800
H	5.21469400	-0.64846700	1.76188300
H	5.61186200	-2.36110800	1.96044900
H	6.40942500	-1.42007600	0.71027700

DFT, TS3b, -955.19733638 hartree, 4.5 debye

N	-1.06364400	-0.85350000	-1.52797400
N	-0.07486300	-0.08925000	-1.84624900
N	-0.22505800	1.15411200	-1.32620200
C	-1.37627900	1.15656000	-0.60233400
N	-1.89984300	-0.12795000	-0.73177200
C	-3.08186900	-0.70266700	-0.19547100
C	-3.30016000	-2.07889900	-0.34277400
C	-4.45913100	-2.64733300	0.17913700
C	-5.39818600	-1.86117500	0.85003300
C	-5.16876300	-0.49276400	0.99214600
C	-4.01732100	0.09787300	0.47214000
H	-2.56725100	-2.68110700	-0.86471300
H	-4.62565400	-3.71392800	0.06058000
H	-6.29900200	-2.31112200	1.25603100
H	-5.89326600	0.12985700	1.50894000
H	-3.83358000	1.15882600	0.57695000
O	-1.84009800	2.09933300	0.09334400
C	-0.31594600	2.65224700	1.56326300
C	0.74250000	1.83924100	1.21825100
C	1.46990900	2.00619200	0.02893900
C	2.56312800	1.04666200	-0.36386700
C	2.64568700	-0.30081700	0.37436500
H	-0.91970000	2.42374200	2.43403300
H	0.86415300	0.90724800	1.76001200
H	2.46600300	0.86827300	-1.44034800
H	3.51455600	1.58904000	-0.24299500
H	1.67255900	-0.80123900	0.29087000
C	1.48538000	3.31330700	-0.71605600
H	2.20854000	3.98447500	-0.23249800
H	1.80075500	3.15978500	-1.74885100
H	0.51078400	3.79847400	-0.72908300
H	-0.44954700	3.63920900	1.14574200
H	2.83265700	-0.13580600	1.43955800
C	3.69919200	-1.19080000	-0.23019900
H	3.48650800	-1.49743100	-1.25489000
C	4.83740500	-1.63427300	0.32487200
C	5.77234400	-2.53197800	-0.45011400
H	6.76926800	-2.08047300	-0.54065000
H	5.91319100	-3.49213700	0.06377500
H	5.39946000	-2.73802200	-1.45681700
C	5.30022500	-1.31475500	1.72530200
H	4.62604300	-0.64782700	2.26582500
H	5.40749500	-2.23364400	2.31640800
H	6.29197600	-0.84446400	1.70495600

DFT, TS3b', -955.19685072 hartree, 4.4 debye

N	1.78123700	1.46002100	-1.20802800
N	0.49645600	1.52927700	-1.12631400
N	-0.01655400	0.44601300	-0.49362700
C	1.02676700	-0.34439300	-0.12604000
N	2.16799100	0.31350300	-0.57874800
C	3.53886100	-0.04043500	-0.47767800
C	4.51471100	0.89190700	-0.85426200
C	5.86090400	0.54900500	-0.75992300
C	6.24414600	-0.70850100	-0.28901200
C	5.26364400	-1.62738400	0.08480200
C	3.90953200	-1.30674500	-0.00791700
H	4.20886000	1.86469600	-1.21818900
H	6.61398900	1.27357600	-1.05535700
H	7.29568400	-0.96871500	-0.21615000
H	5.54896100	-2.61007900	0.44886200
H	3.14365500	-2.01530600	0.27806400
O	0.98018700	-1.41484100	0.53722800
C	0.00130600	-0.97346900	2.44058400
C	-0.34326900	0.35311900	2.29939100
C	-1.34167500	0.80613400	1.41977800
C	-2.44171500	-0.10509100	0.92992700
C	-3.23613800	0.37043300	-0.30324500
H	0.84618300	-1.25480600	3.05845700
H	0.32953700	1.10330300	2.70778300
H	-2.03043500	-1.09586400	0.72710100
H	-3.13925200	-0.22278000	1.77470500
H	-2.51422900	0.56325900	-1.10508400
C	-1.54641200	2.28393800	1.27187500
H	-1.62405800	2.56310800	0.21883300
H	-2.48823700	2.56949100	1.75988500
H	-0.73300800	2.85463500	1.72359800
H	-0.65778500	-1.77882600	2.15191800
H	-3.74009000	1.31621700	-0.08692800
C	-4.21951900	-0.67563500	-0.75500800
H	-3.75863200	-1.60514300	-1.09313400
C	-5.55927300	-0.60767800	-0.79202500
C	-6.37649300	-1.77499400	-1.29219300
H	-7.07466700	-2.12791200	-0.52144200
H	-6.98993600	-1.48607800	-2.15594200
H	-5.74623700	-2.61677800	-1.59109100
C	-6.37756700	0.58730600	-0.36806700
H	-5.77542000	1.41508500	0.01066100
H	-6.97393800	0.96512000	-1.20873000
H	-7.09320000	0.30572300	0.41546100

MP2, TS3a, -952.24896144 hartree, 7.7 debye

N	-0.33396400	-1.63861700	-1.88482000
N	-1.65666700	-1.73330200	-1.80942800
N	-2.07120100	-1.75934300	-0.51758200
C	-0.97600200	-1.65913600	0.28731400
N	0.10384400	-1.60869400	-0.59694700
C	1.46661600	-1.40952000	-0.29309100
C	2.28515700	-0.74226400	-1.21274700
C	3.63215400	-0.55266400	-0.90817300
C	4.15499300	-1.00134100	0.30862300
C	3.32133900	-1.65250100	1.22109000
C	1.97436100	-1.86726000	0.92810700
H	1.86311000	-0.40383900	-2.14839100
H	4.27300700	-0.04968100	-1.62188200
H	5.20177300	-0.84955900	0.53979300
H	3.72153400	-2.00658000	2.16295100
H	1.31157200	-2.35823200	1.62527900
O	-0.92829400	-1.53694400	1.54253600
C	-3.46086500	-0.86157100	1.53724600
C	-2.99440200	0.43029000	1.45465900
C	-3.08372300	1.14467000	0.25925200
C	-2.09427200	2.22126500	0.04305300
C	-0.79998300	1.49805300	-0.47205100
H	-3.20558100	-1.48081400	2.38219700
H	-2.33426700	0.79506600	2.22917100
H	-2.42909900	2.95295600	-0.69147400
H	-1.84447200	2.72546500	0.97711700
H	-1.02859200	0.97626600	-1.40464600
C	-3.91846400	0.71085200	-0.88487300
H	-4.87637400	0.30314400	-0.57203600
H	-4.07570800	1.52728300	-1.58465200
H	-3.36676300	-0.09833400	-1.38734100
H	-4.09737400	-1.28676900	0.77810300
H	-0.53669300	0.73970100	0.25979600
C	0.32788100	2.45679200	-0.66813400
H	0.25860100	3.11130800	-1.53281700
C	1.41501500	2.55260100	0.12278200
C	2.52171300	3.51090200	-0.21817500
H	2.70932800	4.20076600	0.60734800
H	3.44985500	2.96224100	-0.39304600
H	2.29291000	4.09287200	-1.10975900
C	1.66160100	1.72978200	1.35595200
H	0.84753200	1.05539300	1.61060400
H	2.55863500	1.12336900	1.22139500
H	1.84062700	2.38952300	2.20843000

MP2, TS3a', -952.24252078 hartree, 8.3 debye

N	-1.02983800	-1.42897100	-1.10388400
N	0.05535000	-0.88315800	-1.63123100
N	-0.01386100	0.46689700	-1.56713800
C	-1.18467900	0.81617900	-0.97771200
N	-1.81185300	-0.38967500	-0.69016800
C	-3.05748900	-0.62624000	-0.06150700
C	-3.36391900	-1.91847600	0.38208900
C	-4.59379400	-2.15003600	0.99393500
C	-5.50482200	-1.10696400	1.17824600
C	-5.18000300	0.17641000	0.73722300
C	-3.95943700	0.42975500	0.11069000
H	-2.64900300	-2.71442300	0.23315000
H	-4.83629500	-3.14908300	1.33362600
H	-6.45641200	-1.29385900	1.65901200
H	-5.88141400	0.98999300	0.87351200
H	-3.69001400	1.41705100	-0.23374700
O	-1.57759100	1.99363300	-0.73046200
C	0.60932600	2.83463100	-1.38281400
C	0.88622800	2.99546900	-0.02537600
C	1.89896200	2.32000300	0.62286300
C	2.69193500	1.22016100	0.00968600
C	2.36975100	-0.13211000	0.70524000
H	-0.17998900	3.40839100	-1.83926300
H	0.24419300	3.65365500	0.54476700
H	2.48525900	1.10501100	-1.05015800
H	3.75876700	1.42374000	0.13940900
H	1.28680700	-0.26169400	0.71812600
C	2.13479700	2.59088400	2.07347000
H	1.65691600	1.80833000	2.66812200
H	3.19969200	2.56500000	2.30411500
H	1.71822900	3.54840400	2.37838300
H	1.29096200	2.32329500	-2.03890100
H	2.71681200	-0.10474600	1.73841900
C	2.98627700	-1.26461600	-0.05952800
H	2.38601300	-1.62449900	-0.88860900
C	4.19325400	-1.81549800	0.16950200
C	4.69408700	-2.94300800	-0.68975500
H	5.63650800	-2.67536600	-1.17320600
H	4.88867300	-3.83274400	-0.08663000
H	3.97267000	-3.20336500	-1.46199600
C	5.13016600	-1.39366300	1.26741600
H	4.75226900	-0.55874500	1.85363200
H	5.32226000	-2.22585100	1.94822700
H	6.09540600	-1.10049200	0.84865000

MP2, TS3a'', -952.23962938 hartree, 9.1 debye

N	1.74110400	-1.90293000	-0.13424800
N	0.49399700	-1.72512300	-0.53514300
N	0.31533500	-0.45193600	-0.95931900
C	1.48128700	0.22726100	-0.82225200
N	2.37074300	-0.70218800	-0.30335500
C	3.73283600	-0.54345000	0.04666600
C	4.36486200	-1.54004900	0.80019200
C	5.70780500	-1.38494700	1.13649100
C	6.41378700	-0.24585200	0.74174200
C	5.76642600	0.74115000	-0.00238500
C	4.42534000	0.60218700	-0.36105700
H	3.80567000	-2.41449200	1.09904900
H	6.20086400	-2.15582700	1.71523700
H	7.45576100	-0.13044700	1.01114400
H	6.30649100	1.62622300	-0.31457500
H	3.90773800	1.35758300	-0.93331100
O	1.67309700	1.44681900	-1.10044800
C	-0.69107100	1.67024100	-1.59124100
C	-0.88218500	2.29852600	-0.35842800
C	-1.71761300	1.77968600	0.60657400
C	-2.44349000	0.48993100	0.43514700
C	-3.76969600	0.66353700	-0.34940200
H	-1.31881700	0.85303500	-1.89920400
H	-0.30891800	3.19057600	-0.14616600
H	-1.79075800	-0.22519300	-0.07590900
H	-2.68628600	0.08297700	1.41813400
H	-3.54517800	1.08975000	-1.33074100
C	-1.89858700	2.49559600	1.90238600
H	-2.96179600	2.58538700	2.13330800
H	-1.45278500	1.90000200	2.70297000
H	-1.44221600	3.48271600	1.90236400
H	-0.07218100	2.13092100	-2.34281200
H	-4.40529000	1.38536500	0.16488900
C	-4.45410000	-0.65801100	-0.51245700
H	-3.93268700	-1.36370400	-1.15352500
C	-5.59549600	-1.06771300	0.07226600
C	-6.13077000	-2.44876300	-0.18825700
H	-6.21522200	-3.01200600	0.74341000
H	-7.13177800	-2.39975200	-0.62210100
H	-5.48868200	-3.00617400	-0.86776300
C	-6.43451100	-0.23942800	1.00501500
H	-6.01759800	0.74466700	1.20309100
H	-7.43722800	-0.10556300	0.59411600
H	-6.55109200	-0.75477700	1.96042900

MP2, TS3b, -952.27651013 hartree, 4.2 debye

N	-0.86315400	-0.79764700	-1.52363900
N	0.13396100	0.03410700	-1.80789800
N	-0.11148000	1.26019900	-1.26795500
C	-1.30697100	1.20069100	-0.61457900
N	-1.73877200	-0.11314800	-0.76301900
C	-2.90513800	-0.73365300	-0.24723400
C	-2.89800600	-2.11685900	-0.04245200
C	-4.04578700	-2.72832800	0.45693700
C	-5.17420300	-1.96683500	0.77162400
C	-5.15726500	-0.58540400	0.57226200
C	-4.02610700	0.04509600	0.05376900
H	-2.01308300	-2.68740000	-0.28642500
H	-4.05273100	-3.79952900	0.61162700
H	-6.05942300	-2.44719900	1.16761600
H	-6.03087500	0.00760100	0.81032600
H	-3.99961000	1.11232100	-0.10705100
O	-1.84749300	2.10112100	0.10117600
C	-0.55148900	2.53927300	1.58373400
C	0.55474900	1.75773000	1.28933300
C	1.31267800	1.98704200	0.13738200
C	2.45582900	1.07445100	-0.21766400
C	2.42561100	-0.33344600	0.38171500
H	-1.18904000	2.27001500	2.41519700
H	0.62900600	0.78941400	1.76545100
H	2.50035900	1.00517300	-1.30608300
H	3.37839000	1.58034200	0.09422800
H	1.44223400	-0.77235200	0.20142500
C	1.38970400	3.35520200	-0.48369000
H	2.06895600	3.97473000	0.10778600
H	1.77876100	3.27774900	-1.49547800
H	0.41863300	3.83804600	-0.53200800
H	-0.60329300	3.57168600	1.27839100
H	2.56736200	-0.28495300	1.46188300
C	3.46299500	-1.20080000	-0.26557600
H	3.22110800	-1.51018500	-1.27888600
C	4.64400600	-1.59200400	0.24794200
C	5.58039100	-2.45398700	-0.55353800
H	6.54397300	-1.95826600	-0.69133400
H	5.77971600	-3.39534100	-0.03640300
H	5.17002300	-2.68385000	-1.53542500
C	5.14169000	-1.23590900	1.62139400
H	4.46456900	-0.58170000	2.16479300
H	5.29498800	-2.13797200	2.21786500
H	6.10990600	-0.73543300	1.55252500

MP2, TS3b', -952.27503700 hartree, 4.1 debye

N	1.81398300	1.34726300	-1.39240800
N	0.49229600	1.47560100	-1.37445000
N	-0.06551600	0.46374500	-0.65481400
C	0.94136200	-0.33034000	-0.19470100
N	2.11226400	0.26490100	-0.64755200
C	3.45741100	-0.12608100	-0.42507500
C	4.46219300	0.84360900	-0.49728900
C	5.78440900	0.45858200	-0.28676500
C	6.09589800	-0.86925600	0.01633100
C	5.07733700	-1.82034600	0.09614500
C	3.74877100	-1.46123500	-0.13128100
H	4.20024200	1.86650100	-0.72762600
H	6.57000500	1.20055200	-0.34818200
H	7.12428200	-1.15968700	0.18733500
H	5.31461900	-2.85110000	0.32584800
H	2.95139200	-2.18683100	-0.07436700
O	0.85035300	-1.31324600	0.60923400
C	0.07160900	-0.60418200	2.30888400
C	-0.24648800	0.71254200	2.00802700
C	-1.22113400	1.02735900	1.05753300
C	-2.34805300	0.06677200	0.75968500
C	-3.17658600	0.38233100	-0.48972500
H	0.91107700	-0.81181200	2.95912400
H	0.47275100	1.48431800	2.25747400
H	-1.94724900	-0.94213800	0.66957400
H	-3.01437300	0.07108500	1.63153700
H	-2.49261900	0.49080100	-1.33329400
C	-1.45317300	2.47039200	0.72341800
H	-1.68560200	2.59935900	-0.32991000
H	-2.30138100	2.83296100	1.31104300
H	-0.58025900	3.07497800	0.96006100
H	-0.67373300	-1.38144400	2.25819900
H	-3.69605800	1.33251700	-0.36863200
C	-4.14735400	-0.72652300	-0.76818000
H	-3.70833000	-1.62444900	-1.19602300
C	-5.46945000	-0.74239400	-0.51483900
C	-6.30100800	-1.95152000	-0.84443600
H	-6.78761300	-2.34548300	0.05069000
H	-7.09497100	-1.69570800	-1.54969700
H	-5.69755500	-2.74456000	-1.28319900
C	-6.23760000	0.39904100	0.09180300
H	-5.60660500	1.24182900	0.36233200
H	-7.00293600	0.75574200	-0.60097100
H	-6.75796900	0.06545200	0.99216000