Molecular orbital study of the conformational energies $(-\Delta G^{\circ})$ or A values) of 2-alkyltetrahydro-2H-thiopyrans (tetrahydrothiopyrans, thiacyclohexanes, thianes)

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ABSTRACT: *Ab initio* 6–31G* and MP2/6–31G*// 6–31G* methods and density functional (pBPDN**) theory were used to calculate the geometries and relative energies of the chair, boat, and twist-boat conformations of tetrahydro-2*H*-thiopyrans (tetrahydrothiopyrans, thiacyclohexanes, thianes). The chair conformation of thiacyclohexane is 5.3 and 8.0 kcal mol⁻¹, respectively, (1 kcal = 4.184 kJ) more stable than the twist-boat and boat structures at the MP2/6–31G*// 6-31G* level. *Ab initio* methods were used to calculate the relative energies of the rotamers in the chair conformations of 2-alkylthiacyclohexanes (CH₃, C₂H₅, *i*-C₃H₇, *t*-C₄H₉, *neo*-C₅H₁₁, SiMe₃). The MP2/6–31G*// 6-31G* conformational energies ($-\Delta G^{\circ}$ or *A* values, kcal mol⁻¹) of the 2-alkylthiacyclohexanes (Me = 1.00; Et = 1.16; *i*-Pr = 1.02; *t*-Bu = 3.56; *neo*-Pent = 1.04; SiMe₃ = 2.05) are smaller than those calculated for the corresponding alkylcyclohexanes and 4-alkylthiacyclohexanes. Plots of the calculated conformational energies ($-\Delta G^{\circ}$) for the 2-alkylthiacyclohexanes versus the calculated conformational energies for the corresponding alkylcyclohexanes are linear (slope = 0.680 and r = 0.983 for 6–31G* and slope = 0.667 and r = 0.989 for MP2/6–31G*// 6–31G*). The C(2)—S(1) bond lengths are in the range 1.815–1.819 Å. With the exception of the axial 2-*t*-Bu substituent (103.0°), the C—S—C angles vary from 98.6° to 100.4°. The S(1)—C(2)—C(7) angle in the most stable axial conformer is larger that the corresponding angle in its most stable equatorial conformer. Copyright © 1999 John Wiley & Sons, Ltd.

 $KEYWORDS: conformational\ energy;\ 2-alkyltetrahydro-2 \textit{H-} thiopyrans;\ tetrahydrothiopyrans;\ thiacyclohexanes;\ thianes;\ molecular\ orbital\ study$

INTRODUCTION

Substituents in the chair conformation of cyclohexane $^{1-3}$ or in the 4-substituted half-chair conformation of cyclohexene (F. Freeman and W. J. Hehre, unpublished work) prefer the equatorial position. In monosubstituted cyclohexanes (1), the axial substituents have steric interactions with the synaxial hydrogens at C(3) and C(5) [1,3-diaxial interactions; Eqn 1]. The difference in energy between the most stable axial and equatorial conformers is designated the conformational energy $(-\Delta G^{\circ} \text{ or } A \text{ value})$. The free-energy difference between

conformers is referred to as the conformational energy,^{4,5} or sometimes as the A value.⁶ For substituted cyclohexanes it is conventional to specify the value of $-\Delta G^{\circ}$ for the equilibrium

$axial \rightleftharpoons equatorial$

Since $-\Delta G^{\circ}$ will be negative when the equatorial conformation is more stable than the axial, the value of $-\Delta G^{\circ}$ is positive for substituents that favor the equatorial position. The larger the conformational energy, the greater is the preference for the equatorial position. The replacement of carbon by another element in cycloalkanes or cycloalkenes (F. Freeman, Z. M. Tsegai and W. J. Hehre, submitted; N. L. Allinger and K. H. Chen, personal communication) produces changes in several structural parameters and consequently affects the conformational characteristics of the molecules. For example, introduction of a nitrogen, an oxygen or a sulfur atom for a methylene group in cyclohexane will lead to changes in bond lengths (C—C 1.54 Å; C—N 1.47 Å; C—O 1.43 Å; C—S 1.82 Å) and in bond angles. These new elements may drastically change the non-

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bonded interactions and conformational flexibility of the six-membered ring [Eqn 2].

The conformational properties of six-membered rings containing sulfur [3, Eqn 2] have been studied much less than the oxygen systems. This study was undertaken in order to help elucidate the influences of replacing a methylene group in cyclohexane with a sulfur atom by comparing the calculated geometries $(6-31G^*)$ and single point energies $(MP2/6-31G^*)/(6-31G^*)$ of the conformers [chair, C_s point group (2); boat, C_s point

group (4); boat (5); twist-boat (skew-boat, C_2 point group (6); planar $C_{2\nu}$ point group (7)] of tetrahydro-2H-thiapyran (tetrahydrothiopyran, thiacyclocylohexane, thiane; 2). The calculated 6–31G* and density functional theory (pBPDN**) results from the chair conformer 2

were compared with the structural parameters from other calculational methods and from microwave spectroscopic and x-ray diffraction data (Table 1). *Ab initio* methods were used to calculate the conformational energies $(-\Delta G^{\circ})$ or *A* value; see earlier) of the rotamers of 2-alkylthiacyclohexanes [3, R = CH₃, C₂H₅, *i*-C₃H₇, *t*-C₄H₉, *neo*-C₅H₁₁, SiMe₃; Eqn 2] and to compare these values with those for the corresponding substituted alkylcyclohexanes (F. Freeman, Z. M. Tsegai and W. J. Hehre, unpublished work) and 4-alkylthianes [8; Eqn 3].

Another objective of this study was to analyze energy differences among conformers and rotamers by considering the *gauche* (synclinal) interactions and repulsive nonbonded interactions in the equatorial conformer rather than solely in terms of the 1,3-diaxial repulsive nonbonded hydrogen-hydrogen interactions in the axial conformer ¹⁰ (F. Freeman, Z. M. Tsegai and W. J. Hehre, unpublished work) [*gauche*: in A—B—C—D, ligands A and D are *gauche* if the torsion angle (ABCD) about the B—C bond is near +60° or -60°. Synclinal (*sc*): in X—A—B—Y, ligands X and Y are synclinal if the torsion angle (XABY) about the A—B bond is between +30° and +90° or between -30° and -90°^{1,12}].

Table 1. Comparison of calculated and experimental bond lengths (Å), bond angles (°) and dihedral angles (°) for thiacyclohexane (2)



Parameter	$MM4[^{11}]$	pBPDN**	6–31G*	$MW[^{20}]$	$ED[^{21}]$
Bond length					
С—Н	1.113	_	_	1.095	1.114
C_2 — H_{ax}	_	1.111	1.085		_
C_2 — H_{eq}	_	1.107	1.083		_
$C_2 - C_3$	1.531	1.532	1.528	1.533	1.528
C ₂ —H _{eq} C ₂ —C ₃ C ₃ —C ₄	_	1.539	1.532	1.533	1.528
C_2 — S_1	1.814	1.836	1.817	1.832	1.811
Bond angle					
H—C—H	106.4	-	_	108.5	105.9
H — C_2 — H	_	107.4	107.4		_
H — C_3 — H	_	107.1	107.1	_	_
C_2 — C_3 — C_4	112.7	112.9	112.8	107.9	112.3
$C_3 - C_4 - C_5$	112.9	113.2	113.2	109.2	113.8
$C_3 - C_2 - S_1$	112.8	97.7	112.9	114.1	112.7
C_2 — S_1 — C_6	97.4		98.5	99.2	97.6
Torsion angle					
C_2 — C_3 — C_4 — C_5	_	59.2	59.4	_	58.6
C_4 — C_3 — C_2 — S_1	_	60.4	60.0	_	60.8
$C_5 - C_6 - S_1 - C_2$	_	53.7	53.4	_	55.4

Table 2. Calculated energies (6–31G*) and energy differences (MP2/6–31G*//6–31G*)^a of conformations of thiacyclohexane



		Energy (hartree)		
Conformation	Dipole moment (D)	6-31G*	MP2/6-31G*/ /6-31G*	
Chair $(2, C_s)$	2.0	-592.681604	-593.456783	
Boat $(4, C_s)$	2.0	-592.668742	-593.444105	
Twist-boat (6 , C_2)	2.2	-592.673231	-593.448390	

^a Energy difference = $\Delta E = E_{\text{chair}} - E_{\text{boat}} = 8.0 \text{ kcal mol}^{-1}$; $\Delta E = E_{\text{chair}} - E_{\text{twist-boat}} = 5.3 \text{ kcal mol}^{-1}$; $\Delta E = E_{\text{twist-boat}} - E_{\text{boat}} = 2.7 \text{ kcal mol}^{-1}$.

COMPUTATIONAL METHODS

The optimized geometries (6–31G*, pBPDN**) were performed with the MacSpartan Plus 1.1.6 and/or Spartan 4.1 and 5.0 computational programs and single point energy calculations (MP2/6–31G*//6–31G*) were performed with the Spartan 4.1 and 5.0 programs. Frequency calculations (Supplementary Material) were computed on geometry optimized structures at the 6–31G* level with the MacSpartan Plus 1.1.6 and/or Spartan 4.1 and 5.0 programs. (MacSpartan Plus and Spartan computational programs are available from Wavefunction, Inc., 18401 Von Karman Avenue, Suite 370, Irvine, CA 92612, USA).

Conformers which contain an alkyl group directed inwards toward the ring are not included in the discussion since they are energetically unfavorable and do not contribute significantly to the population.

RESULTS AND DISCUSSION

Thiacyclohexane (2) has a chair structure similar to cyclohexane (1), although the heterocycle is more puckered in order to accommodate the bond angles and bond lengths characteristic of sulfur. [The 6-31G*calculated flagpole hydrogen to hydrogen distance in the boat conformation of cyclohexane (1) is 2.357 Å. Electron diffraction studies in the gas phase reveal a slight flattening of the chair conformation of cyclohexane. The C—C—C—C torsion angles are 55.9°, compared with 60° for the ideal chair conformation. ¹⁹] Microwave spectroscopy (although the chair conformation of 2 was deduced from the microwave spectroscopic investigation, the rotational constants proved insufficient to specify the molecular conformation uniquely²⁰), gasphase electron diffraction studies,²¹ molecular mechanics calculations 11,21,22 and molecular orbital calculations indicate that the chair conformation of 2 is the most stable form (Table 2). Frequency calculations show that the boat conformer 4 has one imaginary frequency, that the twist boat structure **6** does not have an imaginary frequency and that the planar $C_{2\nu}$ structure **7** has three imaginary frequencies. This suggests that the boat conformer **4** may be a transition state between the chair conformer **2** and the twist boat conformer **6** (Fig. 1). The high energy hypothetical planar form **7** would suffer from 10 eclipsing non-bonded hydrogen-hydrogen eclipsing interactions and bond angle strain.

The experimental inversion barrier for thiacyclohexane (2) is 9.4 kcal mol⁻¹ (1 kcal = 4.184 kJ). [The experimental inversion barriers for cyclohexane (10.3 kcal mol⁻¹), oxacyclohexane (10.3 kcal mol⁻¹) and azacyclohexane (10.1 kcal mol⁻¹) are essentially identical. As one goes down the Periodic Table (O, S, Se, Te), the barriers become lower, probably because the torsional potentials diminish.²³] MP2/6–31G*//6–31G* single point energy calculations show that the chair conformer 2 (Table 2) is 8.0 and 5.3 kcal mol⁻¹, respectively, more stable than the boat (4) and twist-boat (skew-boat, 6) structures.²¹ (*R* values have been proposed

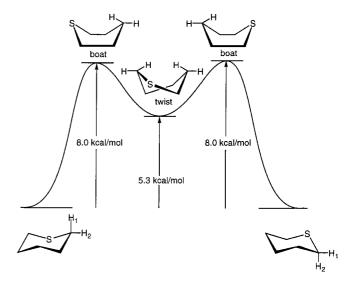


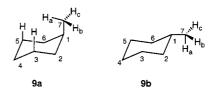
Figure 1. Energy diagram for ring inversion of thiacyclohexane

as a qualitative measure of the distortions of six-membered rings from the chair conformation of cyclohexane^{24–26}). This suggests that the twist-boat conformer **6** is 2.7 kcal mol⁻¹ more stable than the boat structure **4**. Molecular mechanics calculations indicate that the twist-boat conformer **6** is 3.8²¹ or 4.03^{11,22} kcal mol⁻¹ higher in energy that the chair conformer **2**, while the chair structure (**2**) is 5.2 kcal mol⁻¹ more stable than the boat conformer (**4**).²¹ No experimental value is available for the chair–twist-boat energy difference, but it has been estimated to be 4.02 kcal mol⁻¹.^{11,27}

The boat conformer 4 is less stable than the chair form 2 in part because of the eclipsing interactions of the hydrogens on C(2) and C(3) and on C(5) and C(6). The 6-31G* calculated C—S—C bond angle in 4 is 97.0°, which is smaller than the value (98.5°) for thiacyclohexane (2). The boat conformer 3 is expected to be more stable than the boat form 5 owing to an absence of nonbonded interactions between flagpole hydrogens [the transannular strain resulting from steric crowding of the inside (flagpole) C(2) and C(5) hydrogens]. In addition, the boat conformation 5 is less stable than the boat conformer 4 owing to the eclipsing of four hydrogens. As in cyclohexane (1), the boat conformer (4) of 2 is flexible. Twisting of one of the carbon-carbon bonds relative to another leads to a twist-boat conformation (6), which is stabilized by removal of the transannular interaction. There is no transannular strain resulting from steric crowding of the inside (flagpole) C(2) and C(5) hydrogens in 6 since the distance between these two hydrogens is 2.655 Å, which is greater than the sum of their van der Waals radii $(2 \times 1.20 \text{ Å} = 2.40 \text{ Å})$. (The van der Waals radius, which is determined from interatomic distances in crystals, is the effective size of the atomic cloud around a covalently bonded atom as perceived by another atom. The van der Waals radius is not the distance at which repulsive interactions of the electrons on two atoms outweigh the attractive forces between them.^{28–34})

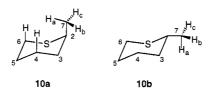
The 6-31G* calculated dipole moment of 2 is 2.0 D while the dielectric absorption measurement³⁵ and microwave spectroscopic (see earlier²⁰) values are 1.71 D and 1.78 D, respectively. In the chair conformer 2, the C(2)—C(3)—C(4)—C(5), C(6)—S(1)—C(2)—C(3) and S(1)—C(2)—C(3)—C(4) torsion angles are 59.4°, 53.4° and 60.0° , respectively. The C(6)—S(1)—C(2)— H_{ax} torsion angle in 2 is 69.7° . The C(2)—C(3)—C(4), C(3)—C(4)—C(5), C(5)—C(6)—S(1) and C—S—Cbond angles are 112.8°, 113.3°, 112.9° and 98.5°, respectively. The C—S—C angle in dimethyl sulfide is $98.8^{\circ}.^{36}$ The $6-31G^{*}$ calculated carbon-sulfur bond length in 2 is 1.817 Å. Electron diffraction gives a carbon–sulfur bond length (r_g) of 1.811 Å and a C—S— C angle of 97.6° for 2 (Table 1).²¹ The 6–31G* calculated C(2)— H_{ax} and C(2)— H_{eq} bond lengths in **2** are essentially the same (1.085 and 1.083 Å), which suggests an absence of anomeric effects. 1,7,9 [The calculational methods used here give some approximation to the equilibrium bond lengths $(r_{\rm e})$, whereas the experimental values cited in the references from electron diffraction $(r_{\rm g})$ and microwave $(r_{\rm s})$ spectroscopic studies give vibrationally averaged values. These numbers are different physical quantities and have different numerical values. Thus, $r_{\rm e}$ needs to be converted to $r_{\rm g}$ and $r_{\rm s}$ in order to make valid comparisons.]

Calculations were performed on the axial [9a, two gauche (synclinal) butane interactions] and equatorial [9b, zero gauche (synclinal) butane interactions] conformers of methylcyclohexane (9) (F. Freeman, Z. M. Tsegai and W. J. Hehre, submitted)^{38–46} [molecular mechanics calculations give a $-\Delta G^{\circ}$ value of 1.77 kcal mol^{-1} for methylcyclohexane $(\mathbf{8}, \mathbf{9})^{39}$] (the gauche interactions in cyclohexanes and thiacyclohexanes may not be comparable to those of gauche butane because the hydrogen-hydrogen interactions are minimized in the cyclic structures. 43-45) for comparison purposes with 2-methylthiacyclohexane (10; see Table 3). At the 6-31G* level, the equatorial diastereomer 9b $(C_s \text{ point group, energy} = -273.243665 \text{ hartree, } \mu = 0.09$ D) is more stable $(-\Delta G^{\circ} = 2.30 \text{ kcal mol}^{-1})$ than its axial isomer **9a** (C_s point group, energy = -273.239998hartree, $\mu = 0.05$ D). At the MP2/6-31G*/6-31G* level, the equatorial diastereomer **9b** (E = -274.160811 hartree) is more stable $(-\Delta G^{\circ} = 1.96 \text{ kcal mol}^{-1})$ than its axial isomer **9a** $(E = -274.157688 \text{ hartree})^{.38-44} \text{ An}$ experimental $-\Delta G^{\circ}$ value of 1.74 kcal mol⁻¹ has been obtained for methylcyclohexane (9) in CFCl₃CDCl₃ via low-temperature carbon-13 NMR spectroscopy. 40,41



Although steric effects are considered to be responsible for equatorial preferences in cyclohexanes, natural bond orbital analysis (NBO)^{38,47} suggests that bond– antibond interactions of the exocylic C(1)—C(7) bond in mainly responsible for the equatorial preference. However, bond-antibond stereoelectronic interactions may not be the sole contributor to the equatorial preference in methylcyclohexane (9). At the $6-31G^*$ level, the C(1)— C(2), C(1)—C(7) and C(1)— H_{eq} bond distances in **9a** are 1.539, 1.534 and 1.088 Å, respectively. The corresponding distances in the equatorial conformer 9b are 1.534, 1.539 and 1.091 Å, respectively. The C(1)—C(2)—C(3) and C(2)—C(3)—C(7) angles in 9a are 113.2° and 112.5°, respectively, while the same angles in the equatorial conformer 9b are 112.2° and 111.6°, respectively. The torsion angles C(3)—C(2)—C(1)—C(7) for 9a and 9b are 73.5° and 179.1°, respectively. Compared with cyclohexane, which has an H_{ax} —C(1)—C(2)—C(3)

Table 3. Comparison of calculated bond lengths (Å), bond angles (°) and dihedral angles (°) for 2-methylthiacyclohexane (10)



	10a	a	101	b
Parameter	MM4 $(r_g)[^{11}]$	6-31G*	MM4 $(r_g)[^{11}]$	6-31G*
Bond length				
C_2 — H_{ax}	_	_	1.114	1.086
C_2 — H_{eq}	1.115	1.085	_	_
C_6 — H_{ax}	1.111	1.084	1.112	1.085
C_6 — H_{eq}	1.112	1.083	1.112	1.083
C_2 — C_3	1.539	1.536	1.537	1.531
$ \begin{array}{cccc} C_{3} & & & \\ C_{4} & & & \\ C_{4} & & & \\ \end{array} $	1.527	1.533	1.526	1.532
C_4 — C_5	1.525	1.531	1.526	1.531
S_1 — C_2	1.820	1.830	1.820	1.830
S_1 — C_6	1.816	1.818	1.813	1.827
Bond angle				
H—C ₆ —H	105.9	107.2	106.2	107.4
C_2 — C_3 — C_4	114.7	114.8	113.0	113.7
S_1 — C_2 — C_3	111.2	111.2	111.1	111.5
C_2 — S_1 — C_6	99.1	100.3	97.7	99.4
Torsion angle				
C_2 — C_3 — C_4 — C_5	59.0	59.7	60.0	60.0
$S_1 - C_2 - C_3 - C_4$	59.2	57.9	62.3	59.4
C_2 — S_1 — C_6 — C_5	54.5	52.8	55.7	53.6

torsion angle of 65.9°, the geometry of the ring in $\bf 9a$ is slightly distorted since the C(3)—C(2)—C(1)—C(7) torsion angle is increased by 7.6° compared with cyclohexane. This outward bending of the methyl group in $\bf 9a$ helps to minimize the repulsive non-bonded hydrogen–hydrogen interactions (see earlier). $^{28-34}$

In the axial diastereomer $\bf 9a$, the C(7)- H_a distance from both C(3)- H_{ax} and C(5)- H_{ax} is 2.401 Å while C(7)- H_b and C(7)- H_c are both 2.453 Å from C(1)- H_{eq} . There are two *gauche* interactions in $\bf 9a$ with respect to the C(3)—C(2)—C(1)—C(7) (73.5°) and C(5)—C(6)—C(1)—C(7) (73.5°) bonds. ¹² In the equatorial diastereomer $\bf 9b$, the methyl group is staggered relative to the H_{ax} —C(1) bond, and the H_{ax} —C(1)—C(7)— H_a torsion angle is 180°. The distances from C(1)- H_{ax} to C(7)- H_b and C(7)- H_c are 2.481 and 2.481 Å, respectively. The equatorial conformer $\bf 9b$ has an anti arrangement with respect to the C(3)—C(2)—C(1)—C(7) (179.1°) and C(5)—C(6)—C(1)—C(7) (179.1°) bonds. In this type of analysis, it is generally assumed that the repulsive *gauche* butane interactions dominate the overall energetics. ^{38,40–46}

The enthalpy and entropy contributions can also play important roles in determining the free energy difference between axial and equatorial conformations. ^{40,41,48,49}

In addition, the chair conformation of cyclohexane is viewed as a structure with no strain and is compared with butane. However, the *gauche* interactions in cyclohexane may not be comparable to those of *gauche* butane because the hydrogen–hydrogen interactions are minimized in the cycloalkane.⁴⁵ A value no greater than 0.28 kcal mol⁻¹ per cyclohexane gauche interaction has been suggested.⁴³

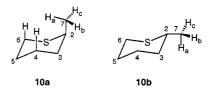
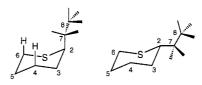


Table 3 shows a comparison of calculated bond lengths, bond angles and dihedral angles for 2-methylthiacyclohexane (10) and Table 4 shows that the equatorial conformer 10b is more stable than the axial conformer 10a ($-\Delta G^{\circ} = 1.00 \text{ kcal mol}^{-1}$). The 6–31G* calculated conformational energy for 10 is similar to the values ($-\Delta G^{\circ} = 1.03 \text{ or } 1.18 \text{ kcal mol}^{-1}$) obtained by Allinger and co-workers^{11,22} using molecular mechanics, but smaller than the value of 1.42 kcal mol⁻¹ obtained from the respective low-temperature ¹³C NMR studies of Barbarella *et al.*⁵⁰ in CD₂Cl₂ and of Eliel and Willer^{51,52} in CDCl₃. The MP2/6–31G*//6–31G* calculated ($-\Delta G^{\circ}$ value of 1.00 kcal mol⁻¹) is smaller than the

Table 4. Calculated energies (hartree), conformational energies $(-\Delta G^{\circ}, kcal \, mol^{-1})^{a}$ and dipole moments (D) of 2-alkylthiacyclohexanes

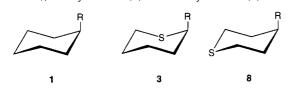


			6–31G*			MP2/6-31G*// 6-31G*	
R	Compound	Energy	$-\Delta G^{\circ}$	Dipole moment	Energy	$-\Delta G^{\circ}$	
Н	2	-592.681604	_	2.0	-593.456783	_	
CH _{3ax}	10a	-631.714339	_	2.0	-632.624102	_	
CH _{3eq}	10b	-631.717384	1.91	2.0	-632.626705	1.00	
C_2H_{5ax}	11a	-670.747960	_	2.0	-671.790164	_	
C_2H_{5ax}	11b	-670.747190	(0.48)	2.0	-671.789499	(0.42)	
C_2H_{5eq}	11c	-670.750466	1.57	1.9	-671.792007	1.16	
$^{2}H_{5eq}$	11d	-670.750092	[1.34]	1.9	-671.791505	[0.84]	
$^{2}H_{5eq}$	11e	-670.748815	[0.54]	1.9	-671.790715	[0.35]	
$-C_3H_{7ax}$	12a	-709.778975		2.0	-710.956741	_	
$-C_3H_{7eq}$	12b	-709.781704	1.71	1.9	-710.958375	1.02	
$-C_3H_{7eq}$	12c	-709.781695	[1.71]	1.9	-710.958195	[0.91]	
$-C_3H_{7eq}$	12d	-709.781191	[1.39]	1.9	-710.957850	[0.70]	
$-C_4H_{9ax}$	13a	-748.805254		1.9	-750.120533		
$-C_4H_{9eq}$	13b	-748.811801	4.11	1.9	-750.126213	3.56	
eo-C ₅ H _{11ax}	14a	-787.844745	_	1.9	-789.289880	_	
eo-C ₅ H _{11eq}	14b	-787.847327	1.62	1.8	-789.291532	1.04	
SiMe _{3ax}	15a	-999.890557	=	1.9	-1001.148292	_	
SiMe _{3eq}	15b	-999.895280	2.96	1.8	-1001.151559	2.05	

^a Conformational energy $(-\Delta G^{\circ})^{1-6}$ refers to the energy difference between the most stable axial conformer and its most stable equatorial conformer. The other values in the $-\Delta G^{\circ}$ columns are the energy differences (in brackets) between the most stable axial conformer and its other equatorial conformers and the energy differences (in parentheses) between axial conformers.

calculated values for methylcyclohexane (9, $-\Delta G^{\circ} = 1.96 \text{ kcal mol}^{-1}$) and 4-methylthiane (3, R = CH₃, $-\Delta G^{\circ} = 1.94 \text{ kcal mol}^{-1}$; Table 5). It is known that there is little variation in conformational energy between

Table 5. Comparison of calculated (MP2/6–31G*//6–31G*) conformational energies ($-\Delta G^{\circ}$, kcal mol $^{-1}$) for alkylcyclohexanes (1) (F. Freeman, Z. M. Tsegai and W. J. Hehre, submitted), 2-alkylthianes (3) and 4-alkylthianes (8) 10



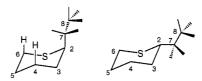
	$-\Delta G^{\circ}$				
R	1	3	8		
CH ₃	1.96	1.00	1.94		
C_2H_5	1.80	1.16	1.75		
i-C ₃ H ₇	1.59	1.02	1.62		
t - C_4H_9	5.45	3.56	5.49		
neo-C ₅ H ₁₁	1.32	1.04	1.39		
SiMe ₃	2.69	2.05	2.76		

different heterocycles and cyclohexane at the 4-position. The smaller $-\Delta G^{\circ}$ value for $\mathbf{10}$ is reasonable since the carbon–sulfur bond is longer than the carbon–carbon bond. The longer bond increases the distance between the axial methyl group and the synaxial hydrogen at C(6) in $\mathbf{10}$ [Eqn 4].



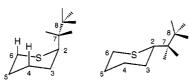
The dipole moments of **10a** and **10b** are nearly equivalent since the exocyclic carbon–carbon bonds are not polar (Table 4). In **10a**, the distances from C(7)- H_a to C(4)- H_{ax} and C(6)- H_{ax} are 2.413 and 2.521 Å, respectively, $^{28-34}$ and C(7)- H_b and C(7)- H_c are 2.487 and 2.439 Å, respectively, from C(2)- H_{eq} . Also in **10a**, C(7)- H_b is 2.496 Å from C(3)- H_{eq} . There are two *gauche* interactions in **10a** and its enantiomer (Table 6). $^{43-46}$ Compared with thiacyclohexane (**2**), which has H_{ax} —C(2)—C(3)—C(4) and H_{ax} —C(2)—S(1)—C(6) dihedral angles of 62.4° and 69.7°, respectively, there is an outward bending of the methyl group in **10a** by 7.9° and 8.1°, respectively. Thus, as in methylcyclohexane (**9**),

Table 6. Calculated torsion angles in 2-alkylthiacyclohexanes



		Torsion angle (°)				
R		C_4 — C_3 — C_2 — C_7	C_6 — S_1 — C_2 — C_7	S ₁ —C ₂ —C ₇ —C ₈	C_3 — C_2 — C_7 — C_8	
CH _{3ax}	10a	70.3	77.8	=	_	
CH_{3eq}	10b	178.6	176.6	_	_	
C_2H_{5ax}	11a	70.2	77.5	66.9	165.8	
C_2H_{5ax}	11b	69.0	78.3	172.2	60.5	
C_2H_{5eq}	11c	177.7	176.6	65.4	171.3	
C_2H_{5eq}	11d	179.3	177.4	170.2	66.5	
C_2H_{5eq}	11e	174.0	178.8	66.6	59.3	
i-C ₃ H _{7ax}	12a	68.2	78.0	63.6 (173.9)	46.9 (169.4)	
i-C ₃ H _{7eq}	12b	172.0	178.0	61.2 (65.8)	64.8 (168.2)	
i-C ₃ H _{7eq}	12c	174.4	178.0	71.7 (162.6)	53.9 (71.9)	
i-C ₃ H _{7eq}	12d	178.7	177.7	61.3 (176.0)	52.7 (175.5)	
t-C ₄ H _{9ax}	13a	82.0	92.4	56.8 (63.8) (179.2)	76.4 (163.0) (46.0)	
t-C ₄ H _{9eq}	13b	172.3	176.8	65.0	60.6	
neo-C ₅ H _{11ax}	14a	68.7	76.1	95.6	137.9	
neo-C ₅ H _{11eq}	14b	178.2	176.7	90.2	137.9	
SiMe _{3ax}	15a	76.4	87.2	68.6 (51.9) (173.9)	158.4 (81.1) (40.9)	
SiMe _{3eq}	15b	175.3	179.8	174.4 (55.9) (65.6)	59.8 (178.3) (60.2)	

Table 7. Calculated (6–31G*) bond angles in 2-alkylthiacyclohexanes



	Bond angle (°)						
R	Compound	C_2 — S_1 — C_6	C_2 — C_3 — C_4	S_1 — C_2 — C_3	S_1 — C_2 — C_7	C_3 — C_2 — C_7	
H	2	98.5	112.8	112.9	_	_	
CH _{3ax}	10a	100.3	114.8	111.2	112.6	113.5	
CH_{3eq}	10b	99.4	113.7	111.5	108.5	112.1	
C_2H_{5ax}	11a	100.3	114.8	110.6	113.6	113.2	
C_2H_{5ax}	11b	100.4	114.7	110.4	111.9	115.2	
C_2H_{5eq}	11c	99.4	113.9	111.0	109.7	111.5	
C_2H_{5eq}	11d	99.7	113.7	111.1	107.6	113.5	
C_2H_{5eq} i — C_3H_{7ax}	11e	98.8	113.1	111.3	110.2	113.9	
i — C_3H_{7ax}	12a	100.4	114.7	109.2	113.0	115.8	
i — C_3H_{7ea}	12b	98.6	113.2	110.7	111.8	113.4	
i — C_3H_{7eq}	12c	99.1	113.1	110.8	109.3	115.3	
i — C_3H_{7eq} i — C_3H_{7eq}	12d	100.1	114.1	110.2	108.8	113.7	
t — C_4H_{9ax}	13a	103.0	117.7	109.9	115.0	118.8	
t — C_4H_{9eq}	13b	99.2	113.4	109.7	111.3	115.3	
neo—C ₅ H _{11ax}	14a	100.2	114.5	110.1	113.8	112.9	
neo — C_5H_{11eq}	14b	99.6	113.9	110.6	109.4	111.8	
SiMe _{3ax}	15a	100.9	114.8	111.0	113.5	118.6	
SiMe _{3eq}	15b	99.8	113.6	111.2	110.4	113.8	

Table 8. Calculated (6–31G*) bond distances (Å) of 2-alkylthiacyclohexanes

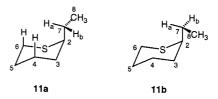
R	Compound	C_2 — C_7	C_2 — C_3	C_2 — S_1	C_6 — S_1
H	2	_	1.528	1.817	1.817
CH _{3ax}	10a	1.531	1.536	1.830	1.818
CH _{3eq}	10b	1.567	1.531	1.827	1.817
C_2H_{5ax}	11a	1.537	1.536	1.832	1.818
C2H50r	11b	1.538	1.537	1.832	1.818
C_2H_{5eq}	11c	1.584	1.532	1.830	1.816
$C_{2}H_{5eq}$ $C_{2}H_{5eq}$ $C_{2}H_{5eq}$ i — $C_{3}H_{7ax}$	11d	1.535	1.532	1.830	1.816
C_2H_{5eq}	11e	1.536	1.532	1.831	1.817
i — C_3H_{7ax}	12a	1.549	1.539	1.836	1.818
i — C_3H_{7eq}	12b	1.545	1.533	1.834	1.816
i — C_3H_{7eq} i — C_3H_{7eq}	12c	1.546	1.533	1.833	1.816
i — C_3H_{7eq}	12d	1.547	1.535	1.836	1.815
t — C_4H_{9ax}	13a	1.566	1.545	1.839	1.819
t — C_4H_{9eq}	13b	1.560	1.537	1.840	1.815
neo — C_5H_{11ax}	14a	1.547	1.538	1.834	1.817
neo — C_5H_{11eq}	14b	1.546	1.533	1.832	1.816
SiMe _{3ax}	15a	1.921	1.540	1.837	1.818
SiMe _{3eq}	15b	1.919	1.539	1.826	1.816

this outward bending of the methyl group in 10a helps to minimize the repulsive nonbonded hydrogen-hydrogen interactions. ^{28–34} In the equatorial conformation 10b, there are no significant non-bonded repulsive interactions involving hydrogens (no distance between two hydrogens of less than 2.500~Å). ^{28–34}

The C(3)—C(2)—C(1)—C(7) torsion angle in **9a** is 73.5° and the C(4)—C(3)—C(2)—C(7) and C(6)— S(1)—C(2)—C(7) torsion angles in **10a** are 70.3° and 77.8°, respectively (Table 6). Hence the steric repulsion is minimized in 10a by an outward bending of the methyl group away from the interior of the thiane ring. The C(2)—S(1)—C(6) bond angles in **10a** (100.3°) and **10b** (99.4°) are larger than the corresponding angle in thiacyclohexane (2, 98.5°; Table 6). The difference (0.90°) in axial and equatorial C(2)—S(1)—C(6) bond angles in 10 is comparable to the difference (0.97°) between the axial and equatorial C(1)—C(2)—C(3) bond angles in methylcyclohexane (9). However, there is a larger difference (4.1°) in axial (112.6°) and equatorial $(108.5^{\circ}) S(1)$ —C(2)—C(7) bond angles in **10** than in the axial and equatorial C(2)—C(1)—C(7) bond angles in methylcyclohexane (0.90°; Table 7). The C(2)—C(7) and C(2)—S(1) bond lengths in **10a** and **10b** are comparable (Table 8).

Table 4 shows that the three equatorial rotamers of 2-ethylthiacyclohexane [11c (methyl group is on sulfur side of ring), 11d (methyl group is opposite sulfur side of ring) and 11e] are more stable than the axial conformer 11a $(-\Delta G^{\circ} = 1.16 \text{ kcal mol}^{-1})$. The MP2/6-31G*// 6-31G*

calculated conformational energies for ethylcyclohexane and 4-ethylthiacyclohexane are 1.80 and 1.75 kcal mol⁻¹, respectively (Table 5).



Each rotamer (11a, 11b) of axial 2-ethylthiacyclohexane has three gauche interactions. In 11a, the methyl group is pointed toward the sulfur side of the ring whereas in 11b the methyl group is pointed away from the sulfur side of the ring. In rotamer 11a, the distances from C(7)- H_a to C(4)- H_{ax} and to C(6)- H_{ax} are 2.445 and 2.477 Å, respectively and the C(7)- H_b to C(3)- H_{eq} distance is 2.379 Å. The C(6)—S(1)—C(2)—C(7) angle of 77.5° in **11a** shows that the ethyl group is tilted away from the interior of the ring. In rotamer 11b, the distances from C(7)- H_a to C(4)- H_{ax} and to C(6)- H_{ax} are 2.404 and 2.500 Å, respectively, the C(7)- H_b to C(2)- H_{eq} distance is 2.435 Å and the C(8)-H to C(3)- H_{eq} distance is 2.288 Å. The stronger repulsive non-bonded hydrogen-hydrogen interactions make rotamer 11b of higher energy (0.42 kcal mol⁻¹) than **11a**. In particular, the C(8)-H to C(3)-H_{eq} repulsive interaction in **11b** is absent in **11a** since its methyl group points toward the sulfur side of the ring.

In the three equatorial 2-ethylthiacyclohexanes there is one *gauche* interaction in **11c** and its enantiomer, one

gauche interaction in **11d** and its enantiomer and two gauche interactions in **11e**. Coincidentally, conformer **11c** is $0.82 \, \text{kcal mol}^{-1}$ (one gauche interaction) more stable than conformer **11e**^{51,52} (in solution, the entropy and enthalpic terms must be considered^{40,41}). In conformer **11c**, the C(7)-H_b to C(2)-H_{ax} and C(3)-H_{eq} to C(7)-H_{ax} distances are 2.498 and 2.388 Å, respectfully. In conformer **11d** the distances from C(2)-H_{ax} to C(7)-H_a and from C(3)-H_{ax} to C(7)-H_b are 2.495 and 2.288 Å, respectively. In **11e**, the distances from C(2)—H_{ax} to C(7)-H_a and from C(2)-H_{ax} to C(7)-H_b are 2.382 and 2.411 Å, respectively, and the C(3)-H_{ax} to C(8)-H distance is 2.364 Å. Thus the greater stability of **11c** versus **11e** is due to one less gauche interaction and fewer repulsive non-bonded hydrogen interactions.

The C(2)—S(1)—C(6) bond angles in **11a** (100.3°) and **11c** (99.4°) are the same as the respective values for **10a** and **10b** and are larger than the corresponding angle in thiacyclohexane (**2**, 98.5°; Table 7). The large difference (3.9°) in axial (**11a**, 113.6°) and equatorial (**11c**, 109.7°) or (3.4°) in axial (**11a**, 113.6°) and equatorial (**11d**, 110.2°) S(1)—C(2)—C(7) bond angles in **11** gives additional support for the increased steric repulsion in **11a** which leads to an outward bending of the ethyl group (Table 7).

The conformational energy of 2-isopropylthiacyclohexane (12) is $1.02 \text{ kcal mol}^{-1}$ (Tables 4 and 5). The axial conformer exists as the single rotamer 12a with four *gauche* interactions, whereas there are two chiral equatorial conformers (12b, 12c) and an achiral rotamer 12d. The equatorial rotamers 12b and 12c and their enantiomers each has three *gauche* interactions and the other rotamer 12d has two *gauche* interactions. Conformer 12b is 0.11 and 0.32 kcal mol⁻¹ more stable than equatorial conformers 12c and 12d, respectively. One difference between 12b and 12c is that one methyl group of the isopropyl group in the former points toward the sulfur side of the ring.

In the axial conformation 12a, the distances from C(7)-

 H_a to C(4)- H_{ax} and C(6)- H_{ax} are 2.493 and 2.443 Å, respectively, the distance from C(9)-H to C(3)-H_{eq} is 2.220 A and the distance from C(8)-H to C(2)-H_{eq} is 2.499 Å. The C(4)—C(3)—C(2)—C(7) and C(6)— S(1)—C(2)—C(7) torsion angles are 68.2° and 78.0° , respectively (Table 6). The value of 78.0° in 12a is consistent with the isopropyl group bending away from the interior of the ring in order to minimize non-bonded repulsive interactions. In the equatorial conformer 12b, the C(2)- H_{ax} to C(7)-H, C(2)- H_{ax} to C(8)-H, C(3)- H_{eq} to C(7)-H and C(3)-H_{ax} to C(9)-H distances are 2.417, 2.416, 2.383 and 2.293 Å, respectively. In the equatorial conformer **12c**, the C(2)- H_{ax} to C(7)-H, C(2)- H_{ax} to C(8)-H, C(2)-H_{eq} to C(8)-H and C(3)-H_{ax} to C(9)-H distances are 2.456, 2.405, 2.308 and 2.399 A, respectively. Although there are no non-bonded interactions between C(7)-H_a and the ring hydrogens in the equatorial conformation 12d, the distance from C(8)-H to C(3)- H_{ax} is 2.174 A.

The C(2)—S(1)—C(6) bond angles in **12a** (100.4°) and **12d** (100.1°) are similar and are larger than the corresponding angle in unsubstituted thiacyclohexane (**2**, 98.5°; Table 7), while this angle in **12c** (98.8°) is close to that in **2**. The large difference (4.2°) in axial (**12a**, 113.0°) and equatorial (**12d**, 108.8°) S(1)—C(2)—C(7) bond angles in **12** gives additional support for the increased steric repulsion in **12a** which leads to an outward bending of the isopropyl group (Table 7). A similar analysis (111.8°) between **12a** and the higher energy equatorial rotamer **12c** gives a value of 1.2°.

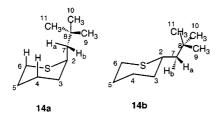
The axial conformer of 2-tert-butylthiacyclohexane (13a) has six gauche interactions and one methyl group pointing into the ring. Tables 4 and 5 show that the equatorial conformation 13b is more stable than the axial conformer 13a ($-\Delta G^{\circ} = 3.56 \text{ kcal mol}^{-1}$). In 13a, the C(8)-H to C(4)-H_{ax} and to C(6)-H_{ax} distances are 2.161 and 2.318 Å, respectively, and the distances from C(9)-H to C(3)-H_{ax} and C(10)-H to C(2)-H_{eq} are 2.212 and 2.331 Å, respectively. The C(4)—C(3)—C(2)—C(7) and C(6)—S(1)—C(2)—C(7) torsion angles in 13a are 82° and 92.4°, respectively.

In the equatorial conformation **13b**, the C(3)- H_{ax} to C(8)-H, C(3)- H_{ax} to C(9)- H_a , C(2)- H_{ax} to C(9)- H_b and C(2)- H_{ax} to C(10)-H distances are 2.325, 2.141, 2.442, and 2.455 Å, respectively. The C(2)—S(1)—C(6) bond angles in **13a** (103°) and **13b** (99.2°) are larger than the corresponding angle in unsubstituted thiacyclohexane (**2**, 98.5°; Table 7). The large difference (3.7°) in axial (**13a**, 115.0°) and equatorial (**13b**, 111.3°) S(1)—C(2)—C(7)

bond angles in 13 gives additional support for the increased steric repulsion in 13a which leads to an outward bending of the *tert*-butyl group.

The equatorial conformer (14b) of 2-neopentylthiacy-clohexane (14) is more stable than its axial conformer 14a ($-\Delta G^{\circ} = 1.04 \text{ kcal mol}^{-1}$; Tables 4 and 5). The conformational energy for 14 is similar to the values for 2-ethylthiacyclohexane (11, $-\Delta G^{\circ} = 1.16 \text{ kcal mol}^{-1}$) and 2-isopropylthiacyclohexane (12, $-\Delta G^{\circ} = 1.02 \text{ kcal mol}^{-1}$).

In the axial conformer **14a**, the distance from C(6)- H_{ax} to C(7)- H_a is 2.413 Å and the distance from C(4)- H_{ax} to C(7)- H_b is 2.531 Å. The distance from C(3)- H_{eq} to C(7)- H_b is 2.339 Å, and the distances from C(2)- H_{eq} to C(9)-H and C(11)-H are 2.193 and 2.430 Å, respectively. In the equatorial conformer **14b**, the C(3)- H_{ax} to C(7)- H_a distance is 2.411 Å and the C(2)- H_{ax} to C(9)-H and C(11)-H distances are 2.203 and 2.376 Å, respectively.



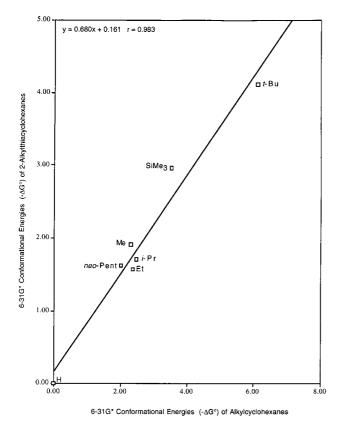


Figure 2. Plot of the 6–31G* conformational energies of 2-alkylthiacyclohexanes versus the conformational energies of the corresponding alkylcyclohexanes

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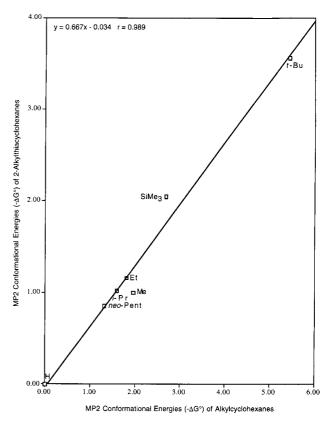


Figure 3. Plot of the MP2/6–31G*//6–31G* conformational energies of 2-alkylthiacylohexanes versus the conformational energies of the corresponding alkylcyclohexanes

The axial conformation of 2-(trimethylsilyl)thiacyclohexane (**15a**) has six *gauche* interactions and the equatorial conformations **15b** has four *gauche* interactions. There are no non-bonded interactions in **15b** and one such interaction (the C(6)- H_{ax} to C(8)-H distance is 2.415 Å) in **15a**. The conformational energy for **15** is 2.05 kcal mol⁻¹, which is smaller than the value of 3.56 kcal mol⁻¹ for 2-tert-butylthiacyclohexane (**13**) owing to the longer carbon–silicon bonds (Table 8).

CONCLUSIONS

The conformational energies of 2-alkylthiacyclohexanes (3) are smaller than the conformational energies of the corresponding alkylcyclohexanes (1) and 4-alkylthiacyclohexanes (8). The $-\Delta G^{\circ}$ values in 3 may be accounted for in terms of *gauche* (synclinal) interactions, repulsive non-bonded interactions, and steric effects in both the

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axial and the equatorial conformer (F. Freeman, Z. M. Tsegai and W. J. Hehre, submitted). $^{38,40,41,43-46,48}$ Although 2-alkylthiacyclohexanes (3) show a smaller preference for the equatorial conformation than the corresponding alkylcyclohexanes (1), the conformational energies are influenced by similar nonbonded interactions and steric factors (Figs 2 and 3). Plots of the calculated conformational energies for the 2-alkylthiacyclohexanes (3) versus the calculated conformational energies for the corresponding alkylcyclohexanes (1) are linear [slope = 0.680 and r = 0.983 for 6–31G* (Fig. 2) and slope = 0.667 and r = 0.989 for MP2/6–31G*/ /6-31G*, Fig. 3].

The 1,3-diaxial interactions which destabilize the axial conformer are weaker in alkylthiacyclohexanes (3) because the carbon–sulfur bond is longer than carbon–carbon bond, which places the synaxial hydrogen and alkyl group farther apart than they would be in the corresponding alkylcyclohexanes [(1); Eqn 4]. The C(2)—S(1) bond lengths in (3) are in the range 1.827–1.840 Å and the C(6)—S(1) bond lengths are in the range 1.815—1.819 Å. With the exception of the axial *t*-Bu substituent (103.0°), the C—S—C angles vary from 98.6° to 100.4°. The S(1)—C(2)—C(7) angle in the most stable axial conformer is larger that the corresponding angle in its most stable equatorial conformer (Table 7).

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