AMBIENT ENERGY MATRIX ISOLATION SPECTROSCOPY

A Molecular Study of p-Toluenesulfonic Acid Solvation



A Senior Honors Thesis

submitted by

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Abstract

Clathrate hydrates are water cages that incase hydrophobic gases, such as methane and CO_2 . They form under extreme conditions (pressure ranging from 9 to several thousand atm, and at cryogenic temperature). They occur naturally in the permafrost and on the continental shelf of the seabed (700 m) where the pressure is about 70 atm. Clathrates are estimated to contain a substantial amount of energy. Extraction of this energy is dangerous and not environmentally favorable, since clathrate hydrates are not stable without its guest gas and at lower pressure environment. Careless extractions could release all the trapped hydrocarbon at once, thus accelerate global warming. Nonetheless, they have the potential to be a good system for cleaning greenhouse gases such as CO_2 and as a gas storage and transfer media. To produce these gas hydrate cost-effectively, promoters, such as the hydrotrope pToluenesulfonic acid (pTSA), are used. The molecular mechanism of how promoters interact with water and the guest gas molecule is studied here using a method called Ambient Energy Matrix Isolation Spectroscopy, or AE-MIS.

AE-MIS uses carbon tetrachloride (CCl₄) matrix to simplify the water spectrum in the mid-IR region. The vibrational signatures of water in CCl₄ – the symmetric stretch, asymmetric stretch, and rotation about the symmetry axis – are present because water exists as monomers in CCl₄ with quenched rotation about the axes perpendicular to the symmetry axis. Monitoring and interpreting changes of these features when studying water interaction with other molecules can give a molecular insight to puzzling mechanisms of bigger systems. Herein, experimental spectra of small (pTSA)•(H2O)_n clusters ($n \le 10$) and large clusters ($n \ge 20$) are presented. Peak assignments and a proposed bimodal distribution model are aided by Density Functional Theory (DFT) calculations.

1. Introduction:

1.1 Motivation: Clathrate Hydrates

Clathrate hydrates are water cages that encapsulate hydrophobic gases, such as methane (CH₄) or carbon dioxide (CO₂), inside. Gas-hydrates deposits (GHDs) occur naturally at the continental shelf of the seabed and in the permafrost due to its extreme formation conditions. For example, methane hydrates are stable from 20 MPa to 2 GPa (~9-20,000 atm) at temperature 70 K to 350 K.² It was estimated by Kwenvolden³ and Makogon⁴ in 1988 that the amount of gas stored in clathrate hydrates contained twice the energy of fossil fuel reserves. Indeed, Makogon estimated in 2007 that with modern energy consumption rate, if just 15% of the amount of energy available in natural GHDs could be produced, the world would have enough energy for 200 years.² Unfortunately, extraction of the natural gases in naturally occurring clathrate hydrates is dangerous, for these gas hydrates are unstable at lower pressure environment. The extraction process may alter the pressurized environment. If done recklessly, a large amount of methane could be released at once. Given that methane is more effective at absorbing infrared radiation than is CO₂,⁵ a sudden increase in methane concentration in the atmosphere could worsen the greenhouse gas effect and accelerate global warming. Furthermore, such an event, or the existence of a "mining" facility at the location of these gas-hydrates deposits, could greatly alter the marine environment around it. Thus, extraction of the gases from naturally occurring clathrate hydrates is not environmentally favorable.

Clathrates also have the potential to be a green energy source by being the gas carrier and storage medium as well as an atmospheric cleanser by trapping CO_2 . A method of reducing CO_2 concentration in the atmosphere is by a process called sequestration, which is performed by releasing the CO_2 in the ocean from 400-2000 m, where they are trapped by dissolution in water.⁶⁻⁸ The pressure and temperature environment fitting for CO_2 hydrates to start appearing is

at 500 m, where they sink due to density and stabilize in the long term on the seabed.^{8,9} CO₂ hydrates density depends on the amount of CO₂ dissolve in the water cage. For CO₂ hydrate to sink in ambient water, the mole ratio threshold of CO₂/water at 128.3 atm and 4 °C (1.3 km deep in the ocean) is 0.145.⁹ Note that this is relative density, as many other parameters factor into CO₂ hydrate density, such as the pressure and temperature of formation. Thus, many aspects of CO₂ marine sequestration, such as CO₂ solubility, CO₂-hydrate formation mechanism, and CO₂-hydrate stability in different environment need to be studied further before putting it to practical use.

The gas storage of natural gas hydrate, specifically methane hydrate, can be achieved by mixing gas and water under hydrate forming condition: 275-283 K and 8-10 MPa (~79-99 atm).¹⁰ Natural gas hydrate is more ideal than liquefied natural gas as a transportation medium since it was found that hydrates can store a large quantity of gas (180 SM³ per M³ of hydrate) and could be stored for 15 days at atmospheric pressure and just below freezing temperature at -15 °C.¹¹ It is estimated that using clathrate hydrates to transport natural gases from the North Sea to Central Europe could save up to 24% in cost compare to liquefied natural gases.¹² In reality, slow hydrate formation rate affects the cost-effectiveness of the formation process, since it requires keeping a cold bath at relatively high pressure. To increase the rate of hydrate formation and storage capacity of the gas hydrates, thereby increasing cost efficiency, promoters could be added to the solution.^{11,13,14}

Several clathrate hydrate promoters have been studied. Tetrahydrofuran (THF), for example, is known to have successfully separate a large quantity CO_2 (more than 99 mol%) from flue gas exhaust at 273-283 K by lowering the equilibrium hydrate formation pressure in a process called hydrate base gas separation.¹⁵ Surfactants such as sodium dodecyl sulfate (SDS)

and dodecyl polysaccharide glycoside (DPG) have also been studied to test their potential in increasing storage capacity of clathrate hydrates¹¹ with the conclusion that anionic surfactant (SDS) is more efficient than nonionic surfactant (DPG).

Another study has been done on similar organic compounds with properties comparable to surfactants called hydrotropes. Surfactants are amphipathic molecule with a long hydrophobic hydrocarbon chain and a hydrophilic end. They are known to self-assemble into micelles in aqueous solutions. Hydrotropes are similarly amphipathic, but instead of a long hydrocarbon chain, they contain a short and bulky hydrophobic end, usually an aromatic group. Gnanendran and Amin studied a common hydrotrope's, para-Toluenesulfonic acid (pTSA), effect on gas hydrate formation and compared it to surfactants.^{16,17} They concluded that pTSA is a better promoter than surfactants due to higher solubility of hydrate forming gases achieved in water.

Although the aforementioned studies have shown that THF and the hydrotrope pTSA have the potential to improve clathrate hydrate formation efficiency, they were all done at a macroscopic level. A main area of interest in clathrate hydrate studies, the study of clathrate hydrate formation mechanism with a promoter, remains largely unexplored. Understanding how the hydrate promoters help gas hydrate formation could provide substantial insights into improving gas hydrate formation rate and efficiency while providing fundamental understanding of molecular-level water-gas-promoters interaction. This potential insight, therefore, could have a large impact in industry and environmental research as well as in fundamental research. This project focuses on pTSA's interaction with water on a molecular level using the CCl₄ matrix, since the same matrix have been used to study THF.¹⁸

1.2 Ambient Energy Matrix Isolation Spectroscopy

Clathrate hydrates, surfactants-water, and hydrotropes-water systems can all be classified as water clusters. Water clusters have been studied in the IR region since the 1970s¹⁹⁻²³ and are important in many biological systems,²⁴ organic chemistry mechanisms,^{25,26} ice,^{27,28} and clathrate hydrate nucleation.²⁹ Water clusters are stabilized through hydrogen bond networks. Infrared absorption spectroscopy (IR) is perhaps the most useful method to study hydrogen bonded (Hbonded) water systems due to the high sensitivity of the OH stretch to the local chemical environment. Unfortunately, broadening in the H-bonded region, which extends hundreds to thousands of wavenumbers, is a major limitation for assigning stretch resonances and diagnosing interactions. Despite the challenges, there have been pioneering theoretical³⁰⁻³⁴ and experimental³⁴⁻³⁸ works providing glimpses into H-bonded network structures in the mid-IR region. Among these pioneering works is a method developed by Kuo and Shultz¹ called Ambient Energy Matrix Isolation Spectroscopy (AE-MIS). AE-MIS uses a carbon tetrachloride (CCl₄) matrix, which operates at in the temperature range of CCl₄'s liquid phase (-22.96°C – 76.72°C) and atmospheric pressure. It simplifies the rotational activity of water in the mid-IR region by restricting certain water's rotational modes as well as preventing hydrogen bonding between water molecules. The more specific molecular picture is explored further below.

a) Water Rotational and Vibrational Signatures in AE-MIS

Water has three rotational modes: about its symmetry axis (z-axis) and perpendicular to its symmetry axis (y-axis and x-axis). Water also as three fundamental vibrational modes: symmetric stretch (3650 cm⁻¹), asymmetric stretch (3760 cm⁻¹), and bend (1600 cm⁻¹).³⁹ Note that the frequencies given above are for water vibrations in gas phase. The symmetric stretch of water in CCl₄ is 3615.5 ± 1 and the asymmetric stretch is 3708.5 ± 0.5 , redshifted due to CCl₄'s dielectric constant (2.2379).¹ The bending mode of water cannot be observed because its

absorption frequency overlaps with CCl₄'s own absorption, and since there are more CCl₄ than water molecules in the sample, the bending mode is swallowed. In IR spectroscopy, a molecule's vibration transition is coupled with a rotational transition. Additionally, water is an asymmetrical top, thus has a complex rotational spectrum extending for hundreds of wavenumbers.³⁹ These complex rotational lines, in CCl₄ matrix, collapse and only the rotational features that are associated with the asymmetric stretch is observed (Figure 1.1). This rotational feature can be identified using symmetry.

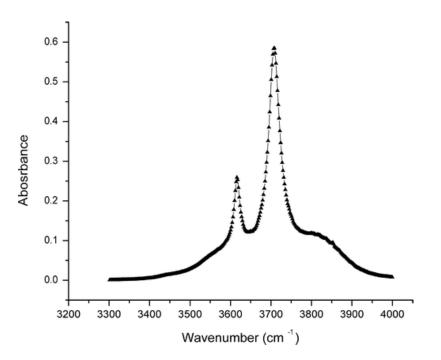


Figure 1.1. FTIR spectrum of water in CCl4. The concentration of water is 7.5 mM. Reproduced from reference 1.

A single water molecule has C_{2v} symmetry. The IR active Γ_{vib} of a C_{2v} point group contains the 2 A₁ and B₂ representations.⁴⁰ Since the A₁ representation is symmetric with respect to the rotation principle axis (z-axis), one must be the symmetric stretch for water and the other the bending mode. The B₂ representations are asymmetric with respect to rotation of the principle axis with subscript 2 denoting the asymmetric mode with respect to the C_n principle axis, thus is the asymmetric stretch of water. The bending mode (A_1) is irrelevant in this context because it cannot be observed due to CCl₄ absorption below 1650 cm⁻¹ as mentioned above.

C_{2v}	Ε	$C_2(z)$	$\sigma_v(xz)$	$\sigma_v(xz)$	Linear functions, rotations	Quadratic functions	Cubic functions
			+1			x^2, y^2, z^2	$z^{3}, x^{2}z, y^{2}z$
			-1			ху	xyz
B_1	+1	-1	+1	-1	x, R _y	XZ	xz^{2}, x^{3}, xy^{2}
B_2	+1	-1	-1	+1	y, R _x	yz	$yz^{2}, y^{3}, x^{2}y$

Table 1.1 – C_{2v} character table.

Looking at a C_{2v} character table (Table 1.1), the symmetric stretch (A₁) must be coupled with the rotations around the x (B₁) and y (B₂) axes. The absence of rotational wings around the symmetric stretch means that the rotation about the x and y axes must be restricted. The asymmetric stretch (B₂) couples with rotation about the y and z axis. Since the rotation about the y-axis is quenched, the rotational wings associated with the asymmetric stretch must be the rotation about the z-axis, or the symmetry axis. The absence of the broad band from 3000-3500 cm⁻¹, generally referred to as the hydrogen-bonded region, also suggests that water monomers, and not hydrogen-bonded water, exists in CCl₄. A possible model that fits these observations is that the slightly negatively charge oxygen atom of the water molecule interacts with the slightly positively charged carbon atom of CCl₄, the conclusion that Shultz and Kuo came to in reference 1. The electromagnetic potential map of water-CCl₄ modeling their interaction is shown in Figure 1.2.

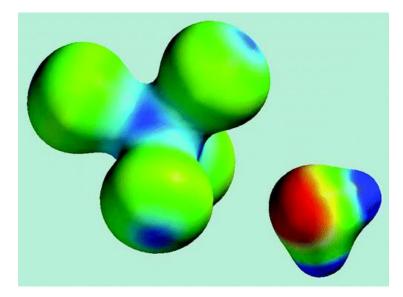


Figure 1.2 A molecular picture of water's interaction with CCl4. Density-potential surface for a water–carbon tetrachloride complex calculated at the DFT- B3LYP 6-31G level of theory. Reproduced from reference 1.*

b) Water as a Pseudo Prolate Top

To confirm the interaction above, water was treated as a near pseudo prolate top for calculation and modeling.¹ Here, we define the rotational constant of the rotation about the z-axis as A, x-axis as B, and y-axis as C. If the oxygen is weakly interacting with the carbon in CCl₄, then it has only one rotation (about its symmetry axis). The other two rotations are restricted (rotational constants equal zero), thus we have A > B = C = 0. Since the rotational constants and their principal moments of inertia have an inverse relationship, I_B and I_C are infinite while I_A is a finite number. Therefore, $I_A < I_B = I_C = \infty$, which is the definition of a prolate top. Quantum mechanics calculations of a prolate top³⁹ were applied to the system and detailed in reference 1. From this model, it is concluded that the vibrational origin is composed of $\Delta J = 0$, $\Delta K = 0$ transitions and $\Delta J = \pm 1$, $\Delta K = 0$ transitions. The former is associated with the vibration origin with no rotational excitation. As a near pseudo prolate top model, the symmetric stretch is a parallel mode, thus has no Q band. Due to the short-lived rotation in the x and y axis, their

rotational signals contributed to make a Q-like band. The latter results in a P-Q-R type structure (P when $\Delta J = -1$, R when $\Delta J = +1$, Q when $\Delta J = 0$, and $\Delta K = 0$ for all). This approximation results in an excellent fit for the spectrum of water in CCl₄ (Figure 1.1) and a calculated rotational constant of 14.8 cm⁻¹ which is the same as water's rotational constant in the gas phase of 14.504 cm⁻¹ (within experimental error).¹

Kuo also reported the upper limit on the rotational lifetime of 0.83 ps, compare to 0.7-1.3 ps in liquid water,¹ which results in rotational bands rather than sharp lines. In the liquid state, water molecules are hydrogen-bonded to each other, thus restricting rotational motions and resulting in a short rotational lifetime. Since water exists as monomers in CCl₄, the only molecule that interacts with water to cause the short rotational life time is CCl₄.

	in CCI ₄	gas phase (ref	Li ⁺ (ref	Na ⁺ (ref	K ⁺ (ref 14)	Cs+ (ref 13)
		37)	15)	14)		
A ₀ (cm ⁻¹)	14.8 (±0.2)	14.5074	13.9	14.3	14.1	14.0
V _{sym} (cm ⁻¹)	3615.5 (±1)	3657	3629	3634	3636	3635
V _{asym} (cm ⁻	3708.5	3756	3691	3707	3710	3711
¹)	(±0.5)					
l _a /l _s	14.8	18	1.53	1.57	2.44	2.04

Table 1.2. Ia/Is of water interaction with different cations. Reproduced from reference 1.

Furthermore, Kuo included a table comparing the ratio of the intensities between the asymmetric and symmetric stretch (I_a/I_s) in systems where water is coupled with different cations (Table 1.2).¹ It is observed that the above ratio decreases with the more positive cation, meaning that the symmetric stretch signal increases with increasing ionic strength. This enhancement makes sense because in cation-water complex, the slightly negative oxygen interacts strongly with the positively charged cation. The oxygen is "pinned down," increasing the oscillator strength of the symmetric stretch. In water-CCl₄ complex, the symmetric stretch is slightly

enhanced compare to the symmetric stretch of gas phase water (ratio I_a/I_s of 14.8 compare to 18, respectively), though not as much as the positive cation-water complex (ratios around 1.5-2.5). This behavior is consistent with the interaction of the oxygen in water with the carbon in CCl₄. Only a slight enhancement is observed because the carbon in CCl₄ is not a full positive charge. All three results: the rotational constant calculated from the near pseudo prolate top approximation, the short rotational lifetime, and the slight enhancement of the symmetric stretch are consistent with the oxygen-carbon interaction model.

Using this simple picture, the rest of this report is organized as follows: chapter 2contains a discussion of experimental and theoretical methods, chapter 3 results from these methods of $(pTSA) \cdot (H_2O)_n$ for small clusters (n ≤ 10) and chapter 4 for large clusters (n ≥ 20). Chapter 5 focuses the hydrocarbon impurities and their implications in this system, and chapter 6 outlines future works and summary.

2. Experimental Methods

2.1 Preparing dried CCl₄

Anhydrous carbon tetrachloride (CCl₄, Sigma-Aldrich, \geq 99.5% anhydrous) is dried further with silica gel for at least 2 days prior to use. The dried silica gel is held in a round bottom flask connected to a vacuum line with 1/4 inch stainless steel tubing (Figure 2.1). The flask is sealed with a greased rubber septum. CCl₄ comes from a bottle with a septum cap, and is removed using a 60 mL syringe and long needle. When not in use, the syringe is kept in a desiccator to prevent moisture absorption from the atmosphere.

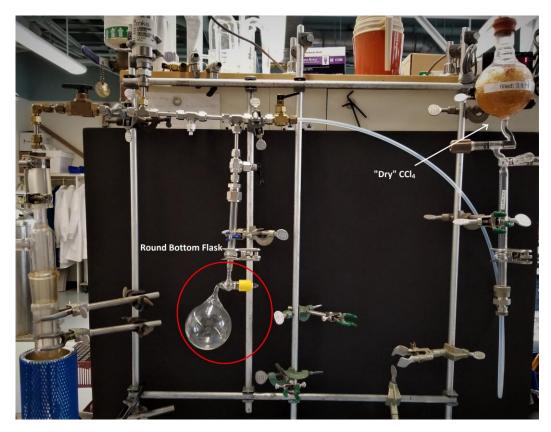


Figure 2.1 – Experimental setup: vacuum line, "dried" CCI_4 in silica gel, and a round bottom flask for sample addition.

2.2 Glass Cell

The glass cell that is used to contain the CCl₄ and acid sample has 25 cm path length and 30 cm diameter. Calcium fluoride (CaF₂) IR windows with 40 cm diameter are used. The windows and cell are held together via Torr Seal epoxy (Torr Seal base resin, 82 grams, Agilent Technologies) to withstand the cold (-20°C) temperature since the temperature range of Torr Seal is -45°C to 120°C (Figure 2.2).



Figure 2.2 – A typical glass cell with CaF_2 windows attached by Torr Seal Expoxy.

2.3 Sample Preparation

a) Aqueous Stock pTSA Solutions

Aqueous stock solutions of different p-toluenesulfonic acid monohydrate (pTSA•H₂O, Sigma-Aldrich, \geq 98.5%, ACS reagent) are prepared by mixing the solid pTSA•H₂O crystals with Nanopure water (18 M Ω , Barnsted GenPure Pro with UV). The acid:water ratios are determine by first measure about 0.50 grams of acid monohydrate, then convert that amount to mole, then decide the desired acid:water mole ratio (1:6, 1:8, 1:10, 1:26, 1:51), and finally convert the desired mole of water to mL by using water density of approximately 1 g/mL. The same process was used to make (pTSA)•(D₂O)_n (D₂O, Signma-Aldrich, 99.9 atom % D) and (pTS⁻Na⁺)•(H₂O)_n (pTS⁻Na⁺, Sigma-Aldrich, 95%) samples.

b) Experimental Procedure

The cell with windows glued on with epoxy are kept in a desiccator overnight prior to sample addition. The empty cell is first evacuated and weighted. A spectrum of the empty cell is acquired using a Nicolet Magna-IR 760 spectrometer with DTGS KBr detector at 64 scans and 1

 cm^{-1} resolution. The spectrometer is purged with nitrogen gas overnight to remove atmospheric moisture prior to use. All FTIR spectra were taken using the same instrument and parameters. Then, dried CCl₄ is added to the cell. Residual CCl₄ usually remains on top of the cell valve after the cell is closed. Thus, the closed cell is evacuated to remove any residual CCl₄ in the valve.

The now filled cell is weighted, and its background spectrum at room temperature is taken. A difference spectrum of dried CCl₄ sample against the empty cell spectrum is also taken. The prepared pTSA solutions can now be added to the CCl₄-filled cell. Microliters are taken from the stock solutions prepared as described above using a 100-microliter syringe. The CCl₄ and acid sample are mixed in an evacuated round bottom flask (the flask is kept in a desiccator when not in use) and is sonicated for thirty minutes to ensure mixing before transferring it back to the cell. The sample is then allowed to equilibrate for 2 days at room temperature.

After 2 days, a difference spectrum of the acid sample in CCl₄ is taken against the CCl₄ background spectrum acquired earlier. The cell is then chilled in a -20 °C freezer for two days before the IR spectrum of the sample at -20 °C is acquired against the CCl₄ background spectrum. More water or aqueous acid solutions could be added in the same manner. All additions of extra water or acid solutions are performed at room temperature. All acquired spectra are then graphed using Origin Lab 9.0.

2.4 Theoretical Methods⁺

Calculations were performed on a Parallel Quantum Solutions Linux box using the Parallel Quantum Solutions (PQS) suite of programs.⁴¹ Density functional theory (DFT) with the hybrid method B3LYP was chosen since it has been reported to give trustworthy results for many water cluster studies,^{42,43} specifically for predicting energies and spectra for gas-phase

⁺ This section along with all other theoretical calculations are credited to Dr. David J. Anick.

water clusters $(H_2O)_n$ and for $X(H_2O)_n$ clusters, where X is an ion or a polar solute, when a triple- ζ basis is employed.⁴⁴⁻⁵³ The triple- ζ basis includes both polarizations and diffuse functions. Diffuse basis functions, used in the PQS programs, caused overlap problems when applied on the aromatic ring which then led to nonconvergence of the self-consistent field. This is not to say that the diffuse basis functions caused this problem, since the causation was never identified, but since diffuse functions are usually not necessary for aromatic optimization, the B3LYP with a "mixed basis" was adopted. The "mixed basis" consisted of 6-311++G(d,p) on the water molecules and S and O atoms of the sulfonyl group but also consisted of 6-311[++]G(d,p), which is the method used for all calculations in this report.

Calculated IR frequencies of the clusters were generated with the harmonic approximation and were scaled by 0.965⁵⁴ prior to comparison with experimental results. Free energies at -20°C and 1 atm were calculated using standard formula, which does not adequately account for the free rotation (treated as low-frequency vibration) of the methyl group. This treatment has minimal effect on the zero point energy (ZPE), but could have a significant effect on the calculated entropy. Methods for calculating rotations are complex. To minimize errors, the principal results of this approach are the free energy differences, which then approximately cancels errors.

Ab initio molecular dynamics (AIMD) studies were also ran for the di- and trihydrate $(pTSA) \cdot (H_2O)_2$ and $(pTSA) \cdot (H_2O)_3$. Each run was of the NVE type (fixed moles (N), volume (V), and energy (E)) starting at the global minimum with kinetic energy added at random to approximate -20°C. In this case, NVE simulates dynamics for a single isolated cluster with no solvent, moving under its B3LYP –computed potential conserving total energy. In these runs,

kinetic energy is defined as the total kinetic energy divided by $\frac{k_B(3N_{at}-6)}{2}$ where $(3N_{at}-6)$ is the number of degrees of freedom. The NVE method was employed as a compromise since it requires less computation time than a highest-quality simulation including solvent molecules filling out a periodic box along with a thermostat. Runs used a step size of 0.5 fs and lasted at least 12 ps after a 1 ps equilibration period.

3: Small (pTSA)•(H₂O)_n Clusters (n \leq 10)

3.1 Experimental Results

The gas-phase IR spectrum of water is complex because water is an asymmetric rotor, but it is simplified in CCl₄ since water exists as monomers in the CCl₄ matrix. The isolated monomers' vibrations and rotations can be modeled as a pseudo prolate top¹ having free rotations about its symmetry axis and quenched rotations about the other two orthogonal axes. A typical water spectrum in CCl₄ shows two fundamental vibrations in the free –OH stretching region: the symmetric and asymmetric stretches at 3617 and 3709 cm⁻¹, respectively (Figure 1.1). The rotational wings resulted from the water's rotation about the symmetry axis accompanies the asymmetric stretch. These features are labeled in figure 3.1 (pure water spectrum shown in reference 1 and Figure 1.1). These features are redshifted relative to gas-phase water due to the dielectric constant of CCl₄ (2.2379). Monitoring changes in these features give insights to the molecular bonding interactions of molecules in the matrix, in this case water and pTSA. For example, the presence of a weak hydrogen bond acceptor restricts the rotation of water monomer about its symmetry axis, which leads to the collapse of the rotational wings. On the other hands, a strong interaction gives redshifted stretch bands and removes equivalence of the two O-H stretches.⁵⁵ Additionally, interaction with the oxygen atom lone pair enhances the symmetric

stretch oscillator strength (hence enhances the intensity of the symmetric stretch peak) relative to the asymmetric stretch.⁵⁶⁻⁶¹ The results reported in this chapter is reproduced from reference 62.

a) $(pTSA) \bullet (H_2O)_n$

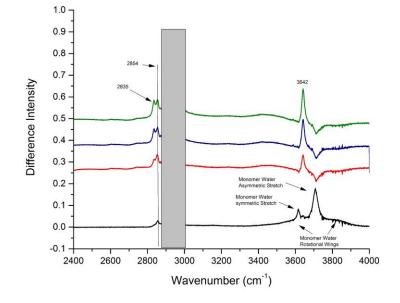


Figure 3.1 – Infrared spectra of $pTSA \cdot H_2O$ in CCI_4 : injected solution mole ratio acid:water 1:6 at room temperature (black line) and -20 °C (red line). Adding 14, and 28 μ L water and cooling to – 20 °C (acid to water mole ratio 1:6, 1:8, 1:10) (red, blue, and green, respectively; spectra offset for clarity). At 1:6, the injected water is more than 20 times that of saturated neat water. The grey region is inaccessible due to hydrocarbons in CCl₄. The low intensity peak at 2854 cm-1 is also an impurity. All spectra are difference spectra subtracting the room temperature, nominally dry CCl₄ spectrum.

The presence of acid in the CCl₄-water system alter the IR spectrum (Figure 3.1). The stock solutions used for the spectra in Figure 3.1 have the acid:water ratio of 1:6 (black at room temperature, red at -20 °C), 1:8 (blue), and 1:10 (green). At 1:6, the water concentration present in CCl₄ is more than 20 times the saturated amount of neat water in CCl₄ (7.5 mM).¹ The symmetric and asymmetric stretch of water are visible at room temperature in Figure 3.1 along with an additional peak at 3642 cm⁻¹.

The spectrum changes when the sample is cooled to -20 °C. The water monomers peaks diminish significantly, shown by the "negative" asymmetric stretch. Since the sample is taken against the "dry" CCl₄ background, the "negative" peak indicates that the concentration of monomer water after the cooling process is less than at room-temperature in the presence of acid. This is consistent with the dehydrating capacity of sulfuric acid and its derivatives, meaning monomer water molecules are hydrogen bonding to each other and to the acid molecule. The small but broad band from 3000-3550 cm⁻¹ is further evidence for this interpretation, since this band signifies hydrogen bonding. The 3642 cm⁻¹ peak enhances as the temperature is cooled and increases further as more water is added. Additionally, a new peak appears at 2835 cm⁻¹ and also increases in intensity as more water is added at -20 °C. This peak is red of the impurity peak at 2854 cm⁻¹, which is a hydrocarbon impurity that appears in all spectra including the CCl₄ solution without acid-water addition. This impurity peak, along with the grey-out region from 2900-3000 cm⁻¹, appears in both 99.5% and 99.9% CCl₄ solutions from Sigma-Aldrich, although less intense in 99.9% solution, which is characteristic of impurities (See Appendix). The impurities peaks are discussed in more details in chapter 5. Changes from cooling are entirely reversible: reheating and recooling the sample produce the same spectra. Therefore, within this acid:water ratio (n \leq 10), the 2835 cm⁻¹ and 3642 cm⁻¹ cannot be distinctly observed at room temperature.

b) $(pTSA) \bullet (D_2O)_n$ and $(pTS^-Na^+) \bullet (H_2O)_n$

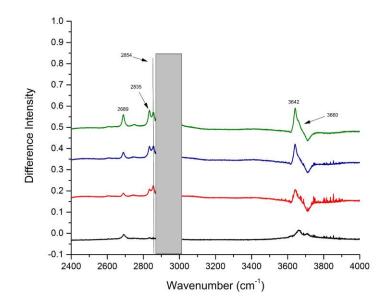


Figure 3.2 – Substitution of D_2O for H_2O to dissolve the acid monohydrate. In CCl₄ at room temperature (black line) and at -20 °C with increasing D_2O (red, blue, and green, respectively). Spectra are difference spectra subtracting the room temperature, nominally dry CCl4 spectrum. Note that pTSA is a monohydrate, hence there are three ordinary hydrogen atoms per acid anion. Mole ratios acid ordinary water to heavy water are 1:1:5 (red), 1:1:7 (blue), and 1:1:9 (green). Spectra are offset for clarity.

To further aid in the identification of the 2835 and 3642 cm⁻¹ resonances, $D_2O^{31,32,63}$ and

the salt version of pTSA, sodium p-toluenesulfonic acid (pTS⁻Na⁺), were used. pTSA•H₂O was dissolved in D₂O using the same method. D₂O stretches are redshifted by $\sqrt{1/2}$ relative to normal water (Figure 3.2). Normal OH modes remain from the normal hydrogen bond of the monohydrate to the acid. Peaks blue of the 3550 cm⁻¹ appears broader than the corresponding peaks in normal water (Figure 3.1 and 3.2). The monomer normal water peaks are less intense in Figure 3.2 than in Figure 3.1, consistent with using D₂O to dissolve the acid rather than H₂O. Due to deuterium substitution, the 3600-3800 cm⁻¹ region is that of the OH resonances of the HOD species. The OD stretch of this species is at 2689 cm⁻¹.

The purpose for using the salt experiment is to test whether the 2835 and 3642 cm⁻¹ are acid-water resonances. Since $pTS^{-}Na^{+}$ is the salt version, the lack of both aforementioned

resonances would be adequate evidence to conclude that they are indeed acid-water peak. The salt-water solutions were prepared in the same way and their spectra are displayed in Figure 3.3. The salt lacks the acid OH on the sulfonyl group, hence fewer features were observed in Figure 3.3. At room temperature, the water monomer peaks are present with lower intensity than in the corresponding acid experiment despite higher water concentration in the stock solution. The 2854 cm⁻¹ peak is more intense in the salt. The reason for this enhancement is discussed in chapter 5. Since the 2835 cm⁻¹, 3642 cm⁻¹, and the broad hydrogen bonded region from 3000-3550 cm⁻¹ features are absent in the salt experiment, they are concluded to be acid-water features.

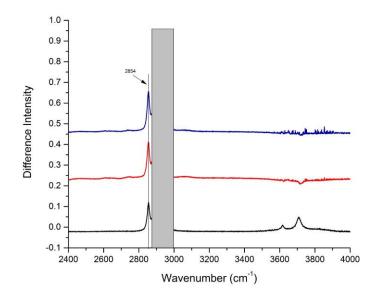


Figure 3.3. Infrared spectra of pTS-Na+ and water in CCl_4 at room temperature (black) and at -20 °C with 0 and 10 μ L added water (salt to water mole ratios 1:20, 1:21; (red and blue, respectively)). Spectra are difference spectra subtracting the room temperature, nominally dry CCl_4 spectrum. Note absence of the 2835 cm⁻¹ peak (the sharp peak at 2854 cm⁻¹ is the impurity mentioned above).

3.2 Computational Results[‡]

a) Low Energy Clusters

Table 3.1 gives the lowest energy structure found at -20 °C for each *n*, $1 \le n \le 12$, and others found within 2 kcal•mol⁻¹ of the lowest. This is given in terms of the free energy of formation from widely separated pTSA and *n*H₂O units, denoted as ΔG^{253} . Unionized clusters are named pTSA.WnX where X is a letter. Ionized clusters are named pTS–.H+WnX. Only the lowest energy structure is listed for isomers having one or more "DA" (single donor-acceptor) water molecules having a free H with two possible orientations. Similarly, only the lowest structure is listed when several local minima occur with respect to rotation around the C–CH₃ or the C–S bonds. Table 3.1 also lists the calculated electronic energy of formation ΔE° as well as ΔG^{0K} (defined as $\Delta E^{\circ} + \Delta ZPE$). There are three clusters whose energy lies beyond the 2 kcal•mol⁻¹ limit but are of interests due to occurring in the AIMD simulation or having a water molecule that donates into the aromatic ring are also included in Table 3.1.

[‡] This section along with all other theoretical calculations are credited to Dr. David J. Anick.

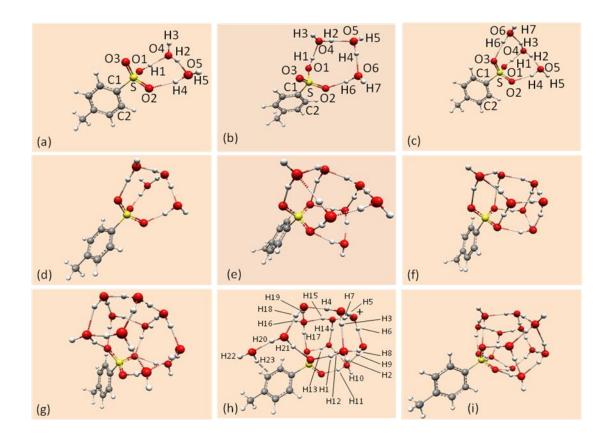


Figure 3.4 – A selection of the optimized (pTSA)(H2O)n clusters listed in Table . (a) pTSA.W2A (b) pTSA.W3A (c) pTSA.W3B (d) pTS-.H+W4A (e) pTS-.H+W7A (f) pTS-.H+W7D (g) pTS-.H+W10A (h) pTS-.H+W11X (i) pTS-.H+W12A.

-				0, 1			
CLUSTER	ΔE°	ΔG^{0K}	ΔG^{253}	CLUSTER	ΔE°	ΔG^{0K}	ΔG^{253}
pTSA.W1	-11.64	-9.10	-2.04	pTSH+W7B	-85.89	-66.97	-13.76
				pTSH+W7C	-87.10	-67.48	-13.57
pTSA.W2A	-24.25	-19.40	-5.50	pTSH+W7D	-87.73	-67.87	-13.38
				pTSH+W7E	-86.33	-66.49	-12.16
pTSA.W3A	-35.70	-28.50	-7.56				
pTSA.W3B	-35.93	-28.84	-6.93	pTSH+W8A	-100.02	-77.84	-16.16
pTSA.W3C	-35.47	-28.03	-6.30	pTSH+W8B	-98.00	-75.97	-14.70
pTSA.W3D	-31.00	-24.55	-4.55				
pTSH+W3A	-35.14	-28.36	-6.11	pTSH+W9A	-112.26	-87.28	-17.66
				pTSH+W9B	-111.80	-86.72	-17.27
pTSH+W4A	-48.70	-38.56	-8.91	pTSH+W9C	-112.30	-87.55	-16.75
pTSH+W4B	-48.56	-38.66	-8.82	pTSH+W9D	-112.74	-87.47	-16.68
pTSA.W4A	-46.60	-37.09	-8.24	pTSH+W9E	-110.38	-85.50	-16.19
pTSA.W4B	-45.14	-35.74	-7.80	pTSH+W9X	-104.96	-80.79	-12.21
pTSA.W4C	-45.89	-36.22	-7.39				
				pTSH+W10A	-123.48	-95.85	-19.04
pTSH+W5A	-59.34	-47.01	-10.49	pTSH+W10B	-122.89	-95.34	-18.52
pTSH+W5B	-60.51	-47.48	-9.70	pTSH+W10C	-123.16	-95.32	-18.29
pTSH+W5C	-59.32	-46.24	-8.83	pTSH+W10D	-122.37	-94.93	-17.77
pTSH+W5D	-61.46	-47.39	-8.74	pTSH+W10E	-119.92	-92.95	-17.59
pTSA.W5A	-57.93	-46.14	-9.58				
pTSA.W5B	-56.28	-44.76	-8.91	pTSH+W11A	-136.05	-105.69	-20.92
				pTSH+W11B	-135.44	-105.03	-19.98
pTSH+W6A	-75.37	-58.21	-11.86	pTSH+W11C	-134.85	-104.48	-19.86
pTSH+W6B	-70.70	-55.34	-11.82	pTSH+W11X	-128.97	-99.38	-15.92
pTSH+W6C	-74.28	-57.47	-11.06				
				pTSH+W12A	-148.03	-114.86	-22.33
pTSH+W7A	-86.60	-67.41	-13.82	pTSH+W12B	-148.10	-114.68	-22.02

Table 3.1 – Selected cluster binding energy (kcal·mol-1) via B3LYP/6-311[++]G(d,p), including electronic energy, ZPE-corrected electronic energy (denoted ΔGOK), and free energy at 253 K.

Selected clusters from Table 3.1 are illustrated in Figure 3.4. The coordinates for these clusters can be found in xyz format in the Appendix. Clusters with the letter "A" are local minima. Breaking an H-bond could result in greater binding energy in some cases. For instance, "pTS⁻H+W7A" and "pTS⁻H+W7D" fall into this category (Figure 3.4e,f): one H-bond in Figure 3.4f is broken in 3.4e. From Table 3.1, the electronic and free energy of 3.4f is lower at 0K, but 3.4e is lower at 253K since the entropy gained from the breaking of that bond lowers the free energy more than the lost enthalpy raises it. Global minima for clusters with $n \ge 8$ have an outer surface consisting of the Eigen ion and DDA waters (Figure 3.4g,i for n = 10 and n = 12, respectively) besides the DAA waters that donate to the negatively charged O atoms of the sulfonyl group. This picture is consistent of the "hydrophobic water"⁶⁴⁻⁶⁶ that tends to repel other similar surfaces or free water.

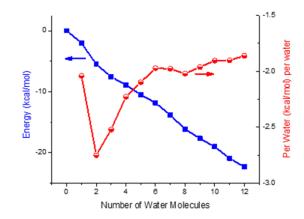


Figure 3.5 - Cluster binding energy as a function of the number of water molecules (blue line/symbols). Note the larger stabilization gain from n = 1 to n = 2 than any other step, shown dramatically in the binding energy per water plot (right axis, red line/symbols).

Free energy of formation of the lowest energy clusters shown in Table 3.1 is graphed in Figure 3.5 for each *n* as a function of *n*. A large gap in binding energy of 3.5 kcal•mol⁻¹ occurs between the mono- and dihydrate. Subsequent water molecules additions have a consistent binding energy of ~1.7 kcal•mol⁻¹. The result is a "kink" at n = 2 due to a slope change. Enhanced binding per water for n = 2 is emphasized by the per molecule binding energy, shown in Figure 3.5. Stabilization at n = 2 implies that the "preferred" number of water per acid molecule is two, meaning the population of the pTSA dihydrate is expected to be overrepresented in a system with populations of various (pTSA)•(H₂O)_n clusters. This interpretation is similar to the situation that causes the n = 21 to be a "magic number" among the H⁺(H₂O)_n clusters.^{38,67-69} Details of the source of this unusual binding energy is under investigation.

Note that the binding energies in Table 3.1 are much smaller at 253K than at 0K due to entropy contribution to water cluster free energy near ambient temperature, which is a known phenomenon as discussed by Shields et al.⁷⁰ for solute-free (H₂O)_n clusters with $2 \le n \le 10$. In Shields' Table 8, ΔE_e , ΔE_0 , and ΔG_T° (T = 10, 200, 298) notations correspond to ΔE^0 , ΔE^{0K} , and ΔG^T (T = 253) in this report. For (H₂O)_n clusters, Shields et al. found that binding energies become significantly smaller at 200 K than at 0 K and positive at 298 K, meaning small water clusters are unstable at 298 K and evaporate over time to dissociate to water monomers. ΔG^{253} values in Table 3.1 are negative, thus the pTSA-water clusters are predicted to be stable at 253 K. It is also found that the entropy term greatly affects both the absolute and relative binding energies (i.e., relative ordering for a fixed *n*) of pTSA-water clusters.

b) Identification of the 2835 and 3642 cm⁻¹ Resonances

2835 cm⁻¹. After 150 (pTSA)•(H₂O)_n cluster optimizations and frequency calculations, we reasoned that the identity of this peak was either somewhere among the OH stretch signals already computed or from an OH stretch with a local bonding environment that is similar to one computed. (The exact frequency of an OH stretch is extremely sensitive to its donor types (DA, DAA, etc.) and (if H-bonded) of its acceptor.⁷¹⁻⁷⁴)

Modes in the range of 2835 ± 15 cm⁻¹ are called "matches" and in the range of 2835 ± 10 cm⁻¹ are "strong matches." In total, 28 matches are found, 12 of which are strong matches ranging from 2830 to 2845 cm⁻¹ and are the O-H stretch of the sulfonic acid H donating to the O of a water molecule in pTSA.W2A, the dihydrate. This stretch is seen in various dihydrate clusters because there are separate optimizations for the pTSA.W2A where either toluene is rotated or the direction of the free H atom is flipped. The other 16 matches occur in clusters with $n \ge 5$ and are the threefold symmetric stretch of a H₃O⁺ embedded in an Eigen ion. Since the Eigen ion is known to have a very broad signal experimentally, even at cryogenic temperature, so the idea of the existence of an Eigen ion at ambient energy can be ruled out. Therefore, the 2835 cm⁻¹ is tentatively assigned the S–O–H--- O of the dihydrate. As described above in Figure 3.5 and Table 3.1, the dihydrate has an unusually strong binding energy and is likely to be overrepresented, thus its signal is predicted to be unexpectedly strong. This assignment of the 2835 cm⁻¹ to the S–O–H--- O stretch is thus consistent with the binding energy calculation. The other O-H stretch of the two water molecules in the dihydrate were either low intensity free O-H bonds or water-water H-bonds, which are too broad at 253 K to be distinctly detected. The full predicted IR spectrum of the dihydrate is included in the Appendix.

3642 cm⁻¹. The same approach is taken for the 3642 cm⁻¹ signal, which resulted in 37 matches. One match is the free O–H stretch of dry pTSA at 3645 cm⁻¹, but it was ruled out because it would imply that pTSA, which exists as the monohydrate, lost *all* its water to the CCl₄ solvent. Fourteen of the matches are from bonds of a DDA water to one O of the SO₃⁻. These bonds are strained (large H–O–O angle), or coupled with other stretches, or both. The 3642 cm⁻¹ for the range of DDA---O–S stretches is on the blue end of the specified range and is expected to occur only if the H---O distance is unusually long or if the frequency is altered by strong coupling, or both. These bonds are one of the weakest and intermittently break and reform at -20 °C, making them poor candidates for the narrow signal observed in the experimental data. Eight other matches are strained and long DDA-DAA bonds, seven are DDA---DDAA or DDAA---O–S are also dismissed as weak and/or strained, thus unlikely to generate a strong, narrow peak.

Finally, there are six strong matches ranging from 3641 to 3651 cm⁻¹ and are the water donating to the benzene ring, denoting π -OH. The benzene π -OH resonance has been reported to be at 3660 cm⁻¹ and is expected to be redshifted about 20 cm⁻¹ when attached with an electron withdrawing group,⁷⁵ such as the sulfonate substituent. An earlier work⁶² hypothesized that the mechanism for the formation of the π -OH is through a "spillover" mechanism. Water H-bonded more favorably with the sulfonyl group than the benzene ring, and thus attempts at H₂O---benzene interaction optimizations resulted in water being ripped away to form bonds at the sulfonyl group. Thus, one way for water to reach the benzene ring in pTSA is for the cluster at the sulfonyl group to get relatively big, then some water molecule *could* spillover and interacts with the benzene ring, forming something like a scorpion tail in Figure 3.4h. The predicted number of water molecules needed for this mechanism to occur was between 16 and 20. More recent computational results did not support this hypothesis. The energy of the "spillover" clusters are not the global minima in terms of binding energy for a given *n*. Thus, a second hypothesis was proposed. Ionic liquids could exist in aqueous solution as totally dissociated ions or as ionic pairs: molecularly dispersed or forming parts of aggregates. In this sense, pTSA was found to be forming aggregates in aqueous solutions.⁷⁶ On a molecular level, this means that the water cluster on pTSA could be a multi-acid cluster. This picture would allow some water molecules to come in contact with the benzene ring of another pTSA molecule while bonding with the sulfonyl group of a pTSA molecule. Since pTSA are bulky groups, it is predicted that the number of water needed for this to occur would need to be large, similarly predicted in the first hypothesis, although the exact number remains to be explored. Thus, the 3642 cm⁻¹ is assigned the π -OH interaction of a water to a benzene ring for a "large, multi-acid cluster" with unspecified size or geometry.

Since the "matching" method compares the gas-phase computation results to the liquidphase experimental observations, one could ask whether solvent interaction, in this case CCl₄, could alter the frequencies significantly. The features of dangling H atoms exposed to the CCl₄ are redshifted by 35 to 50 cm⁻¹ according to both experimental and theoretical result and is shown in the monomer's symmetric and asymmetric stretches. Since all dangling H's in all clusters in this calculation resonate at least 90 cm⁻¹ bluer than 3642 cm⁻¹, solvent effect cannot be the explanation.

c) AIMD Results

NVE AIMD runs for the dihydrate pTSA.W2A and trihydrates pTSA.W3B were performed in the gas-phase. The pTSA.W2A and pTSA.W3B data were collected for 12.2 ps and the pTSA.W3A for 20.3 ps following a 1 ps equilibration interval. Temperature for the dihydrate based on kinetic energy was 268 ± 26 K. From Figure 3.4a, the O1—O4 H-bond remained intact throughout while the O4—O5 H-bond present during 98.1% of the snapshots. Following Kumar's geometric criteria⁷⁷ for O_D-H*---O_A to be deemed H-bonded,[§] the O5—O2 bond broke and reformed several times. Starting at 2.84 ps, it remained severed for 55 ps and become attached to O3 via H4 instead of bonding via H4 to O2 when reformed. This pattern persists for the rest of the run. Overall, an H-bond was present from O5 to either O2 or O3 during 93.2% of snapshots (98.1% if $\beta_{cut} = 40^{\circ}$). The dangling H's flips often.

pTSA.W2

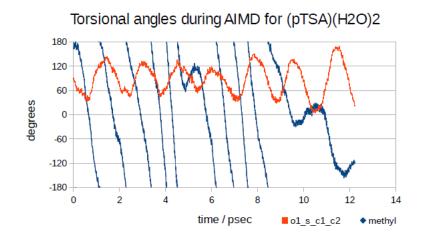


Figure 3.6 – Evolution of the torsional angles – measuring internal rotations – during the AIMD run. The methyl group (blue) spins freely. In contrast, there is no net rotation about the S-O bond.

 $d(H^*, O_A) < 2.5 \text{ Å}, d(O_D, O_A) < 3.4, \text{ and } \beta = angle(H^*-O_D-O_A) < \beta_{cut} = 30^{\circ}.$

Figure 3.6 shows the instantaneous torsional angles, measuring internal rotations of the methyl (blue) and sulfonic (red). The methyl group spun freely (180° to -180°, making a 360° rotation) completing almost 10 net revolutions during the run. On the contrary, no net rotation was observed for the S—C bond but only mere oscillation across a mean position of ~90°. The transition state for rotation about the S—C has $\Delta G^{ts253} = 3.1$ kcal•mol⁻¹ and occurs at orientations of -6° and 174° for the O1—S—C1—C2 dihedral angle. The dihedral angle of the optimized geometry is 81°.

The trihydrate run (pTSA.W3A) lasted for 20.36 ps and temperature 241 ± 29 K revealed a fairly stable bonding pattern. From Figure 3.4b, the O1—H1---O4 bond stays connected for the entire run and O4—H2---O5 was present for 99.9% of snapshots (100.0% if $\beta_{cut} = 35^{\circ}$). An Hbond from O5 to O6 appeared in 98.0% of snapshots, which flipped once from using H4 to H5 midway through the simulation. The bond from O6 to O2 flipped from using H6 to H7 at 17.3 ps and "walked" to an O6—H6---O3 bond at 19.5 ps then "walked" back to O6—H7---O2 at 20.3 ps right before the run completion.

From 2.3 to 3.6 ps, an additional OS—H5---O3 H-bond formed, turning the structure to that of pTSA.W3C and the system also became topologically pTSA.W3C during 19.9 to 20.1 ps when an O5—H4---O2 bond occurred. Comparing the total of 1.5 ps of the additional bond with the 18.8 ps without it yields a ratio of 12.5 favoring pTSA.W3A over pTSA.W3C. This result is consistent with the ratio predicted in Table 3.1. The ΔG^{253} difference between pTSA.W3A and pTSA.W3C of 1.26 kcal•mol⁻¹ equals 2.5 kT and predicts an occurrence ratio of 12.2. Although other trihydrate topologies including the pTSA.W3B would also occur in a long enough run, but the barriers to interchange are higher and were not overcome in this limited simulation.

Another simulation with the similar parameters were performed starting with the pTSA.W3B topology (Figure 3.4c) at 238 ± 23 K. This topology shows less bond stability than the pTSA.W3A and the From Figure 3.4c, the O1-H1---O4 bond was only intact for 6 fs when β crested at 31.2°. H1 "almost" transferred once when d(O1,H1) surged briefly to 1.17 Å. The O5 side was more stable during the run: an O4—H2---O5 bond (respectively, an O5—O2 bond via either H4 or H5) was present in 88.9% (respectively 84.9%) of snapshots. This percentage rose to 98.1% (respectively, 93.3%) when β_{cut} was raised to 40°, suggesting similar behavior to the dihydrate. The O6 water, on the other hand, is freer than the ones described above. The O4— H3---O6 bond was present for 69.2% and 76.0% for $\beta_{cut} = 40^{\circ}$ (47.0%, respectively). The configuration topology match the cluster pTSA.W3D, a high energy cluster, 43.2% of the time with the entire final 3.5 ps when the O6 water appeared to swing wildly on the end of the O4— O6 bond and remained far removed from the sulfonate, matching the low-energy pTSA.W3B (Figure 3.4c) just 14.6% of the time. Similarly, with the dihydrate, the methyl group spun freely, whereas the S—O bond only slowly oscillated. The methyl group completed seven revolutions in this run within the first 4 ps, then 10 more in the reverse direction prior to the run completion. This result is consistent with a classical calculation of the moment of inertia around the C—C bond of the methyl group, yielding 1.31 THz as the frequency when rotational kinetic energy is 1/2kT.

Although bond rearrangements are expected to happen more slowly in solvent than in gas phase, the simulations described still provides relevant insights to the matrix: the AIMD run support the idea of the unusual stability of the dihydrate. While the trihydrate with the pTSA.W3A topology (with ~8% as pTSA.W3C) seems stable, this picture has the possibility to change bond directions if it turns to pTSA.W3B. When converted, the flanking waters (i.e. O5 or O6) is held very loosely by the trihydrate that they could be prone to "evaporation" into the matrix monomer population at this temperature. Contrastingly, when a dihydrate cluster interacts with a monomer, the monomer may join the cluster for a while. The general picture is a dynamic equilibrium where water molecules are exchanged among $(pTSA)(H_2O)_n$ clusters via the monomer population. Investigations on how the dynamics may change once $n \ge 4$ and the cluster dissociates are in progress. Preliminary data on n = 7 cluster shows frequent interchange among top four geometries (Figure 3.4e,f) but no proton transfers that affect the embedded H₃O⁺.

d) D and H Position Preferences in D₂O

When the system contains both ¹H and D in which these isotopes are allowed to freely distribute into the various available positions for H, a "positional isotope effect" can be observed. For example, the central H of the Zundel cation is 2.3 times more likely to be H than D at 80 K.³¹ Another well documented phenomenon of the HOD monomer binding to benzene is its preferred orientation with the O—D making the H-bond.⁷⁸⁻⁸⁰ A previous computational studies of the "positional isotope effects" for a large database of (H₂O)_n clusters found that a J-shaped curved accurately predicted the ZPE of replacing a single position with D as a function of the O—H* distance in the optimized structure.⁸¹ This study also shows that there exist mild cooperativity effects when replacing two H by D atoms on the same or adjacent water molecules. This cooperation is small, thus a reasonable approximation is that the effect of multiple substitution is additive.

The reported experiments involved addition of D_2O followed by observing the $O^{-1}H$ stretch resonances. In these experiments, a few exchangeable H atoms are ¹H from the original pTSA monohydrate (3 ¹H atoms per acid molecule) and a small amount of adventitious water in "dry" CCl₄, but the remainder (10, 14, or 18 hydrogen atoms per acid molecule) is D. Thus, the

water clusters in these systems consist of mainly D_2O with some small number of H substitutions. To investigate the positional isotope effects on the results (or, perhaps, lack thereof), computation on the free-energy difference between the cluster with all D and the cluster with all H were performed in both the pTS-.H+W11X (Figure 3.4h) as a representation of a cluster having a H-bond to the benzene ring and for the dihydrate. The procedure for this computation is outlined in reference 81.

Table 3.2. For two clusters, calculated ZPE change and free energy change (at -20°) from switching a single position from D to H, along with the Boltzmann probability for that position to be H when there is just one H in the cluster.

		ΔG^{253}			ΔΖΡΕ	ΔG^{253}	
	$\Delta ZPE (D \rightarrow H)$	(D→H)	prob		(D→H)	(D→H)	prob
pTS.H+				pTS.H+			
W11X				W11X			
H1	2.152	2.205	0.042	H16	2.032	2.154	0.046
H2	2.036	2.155	0.046	H17	2.156	2.213	0.041
H3	2.179	2.238	0.039	H18	2.183	2.238	0.039
H4	2.158	2.200	0.042	H19	2.168	2.235	0.039
H5	1.978	2.009	0.062	H20	2.147	2.219	0.041
H6	2.120	2.155	0.046	H21	2.157	2.222	0.040
H7	2.038	2.069	0.055	H22	1.955	2.130	0.048
H8	2.176	2.235	0.039	H23	2.007	2.142	0.047
H9	2.173	2.231	0.040	pTSA.W2A			
H10	2.031	2.152	0.046	H1	2.043	2.078	0.234
H11	2.163	2.214	0.041	H2	2.152	2.211	0.179
H12	2.148	2.220	0.040	H3	2.017	2.153	0.201
H13	2.145	2.217	0.041	H4	2.145	2.214	0.179
H14	2.176	2.226	0.040	H5	1.986	2.139	0.207
H15	2.180	2.238	0.039	L	1	1	1

The results from this study is listed in Table 3.2: $\Delta ZPE(D \rightarrow H)$ is the effect of single Dto-H substitution at the indicated hydrogen while the $\Delta G^{253}(D \rightarrow H)$ is the calculated free-energy difference at -20 °C. After one ¹H substitution with a Boltzman distribution at -20 °C, the last column shows the probability of each position having the ¹H. Hydrogen labels are the same as in Figure 3.4a,h. Without the positional isotope effect, these probabilities would be 1/23 = 0.043and 1/5 = 0.2 for the two-water clusters.

These results are consistent with the theory in reference 81: positions in short H-bonds associated with the Eigen cation have the smallest substitution energy, followed by dangling H atoms, and lastly those in H₂O—H₂O H bonds. Figure 3.4h shows the H₃O⁺ with a "+" sign with its three hydrogens H5, H6, and H7. The benzene-bonding H is H23, which is one of the more favored positions to be ¹H in the cluster overall. When the temperature is at -20 °C, the effect is small, thus it is only ~10% more likely to be ¹H than pure chance prediction. As for the dihydrate, the sulfonic acid is 30% more likely to be ¹H than the other bonding H atoms (H2 and H4). This result is uncertain for it is suspected that many pTSA molecules never dissociate in the low-water environment and those that becomes a dihydrate may keep the protium atom that they started with. For pTSA.W2A, a Boltzmann distribution may determine ¹H and D at H2 through H5, but not at H1.

3.3 Implications of the 3642 and 2835 cm⁻¹ resonances

The experimental results of the aforementioned systems – pTSA-(H₂O)_n, pTSA-(H₂O)-(D₂O)_{n-1}, and pTS⁻Na⁺-(H₂O)_n – along with the theoretical results support the existence of at least two structures in the CCl₄ matrix: a dihydrate with two water molecules interacting with the nonionized acid giving rise to the 2835 cm⁻¹ and the other a large water cluster d—OH donating into the π ring giving rise to the 3642 cm⁻¹. Both of these resonances grow with addition of water, suggesting a source consisting of relatively dry acid. Since the amount of water in this system is beyond the saturation point of water in CCl₄, there must be many water-water bonds. Thus, the method for assigning these resonances is as follows. The most intense water resonances are expected to be the Zundel and Eigen associated, but they are known to be very broad. Additionally, the nonhydronium water—water bonds are not observed because they are also known to be very broad under ambient thermal conditions. For nonwater-water interactions in these systems, the SO—H—OH₂ stretch is overrepresented, high-intensity, and is at the right frequency matching the resonance observed at 2835 cm⁻¹. Similarly, the water— π resonance is known⁸² and would redshifts about -20 cm⁻¹ with an electron withdrawing substituent such as the sulfonate⁷⁵ to be at the right frequency (3642 cm⁻¹) and to be narrower than water—water bands. Thus, this section focuses on discussing these two resonances and the smaller 3660 cm⁻¹ shoulder. Then, the section concludes with a possible three-structure model: the dihydrate, the multiwater single-acid cluster π donation, and a relatively dry acid source that is consistent with observations.

a) 2835 cm⁻¹

Strong evidence presented in this studies support the assignment of this peak as an acid resonance. Experimentally, this resonance is not observed in the salt solution even though it contains many times more water in the system than in the acid solution. Since it is only present in the acid system, the possible assignment of this peak to any C—H stretching resonances of the toluene group or any hydrocarbon impurities is eliminated. Additional experimental observations narrow potential assignments: the frequency is unaffected by added water or with substitution of D₂O for H₂O to dissolve the acid, and the intensity increased with added H₂O and D₂O. Therefore, the associated structure must be stable.

The AIMD simulations of the dihydrate structure (Figure 3.4a) supports the claimed stability of this structure at -20 °C. Consistently, both the acid and salts⁶¹ show that a two-water structure is IR-observable. When only two water molecules bind with the acid, a five-member ring is formed, thus stabilizing the dihydrate. Comparing the calculated binding energy of pTSA- $(H_2O)_n$ isomers at -20 °C for n = 1-12 (Table 3.1 and Figure 3.5), the dihydrate structure is preferred over other cluster sizes from n = 1-12 on a per water basis. Thus, there is strong experimental and theoretical evidence of the dihydrate existence in the acid solution.

Furthermore, the 2835 cm⁻¹ resonance also increases in intensity when using D_2O to dissolve the acid monohydrate relative to the intensity gain using normal water. This observation suggests that the 2835 cm⁻¹ is site specific in both normal water and D_2O . Absence of H/D exchange in the D_2O system does not make the ionization impossible at various points in the process as long as the H never departs from the S—O—H---O H-bond in which it starts.

b) 3642 and 3660 cm⁻¹ Shoulder

Similarly, there is strong experimental and theoretical evidence that supports the assignment of the 3642 cm⁻¹ resonance. This peak and the broad hydrogen bonded feature are not detected in the salt system, thus it must be associated with the acid system. Comparing the 3642 cm⁻¹ resonance in normal and D₂O water, one could see that it is somewhat more intense in the normal water system than in deuterated system. The broad resonance is also more intense and structured in normal water. The broad feature from 3000-3550 cm⁻¹ is non-detectable in the salt but noticeable in the acid, which is could be explained from the stronger polarization of the water by the acid.

The strongest d-OH resonances in this system is one donating into the aromatic π cloud, consistent with the π cloud polarizing OH bond, thus enhancing the oscillator strength. There is

ample precedent for this assignment,⁸²⁻⁹⁰ especially the 4 cm⁻¹ wide benzene—water stretch at 3642 cm⁻¹ for n = 5 reported by Pribble and Zwier.⁸² The assignment to a sulfonate donating OH resonance is eliminated since the intensity grows with n. Formation of this cluster is reversible as indicated by its disappearance on warming and reappearance in recooling, consistent with the weak OH— π bond.

The 3660 cm⁻¹ shoulder is assigned the d-OH of the HOD water. A 3660 cm⁻¹ resonance was observed previously in a CCl₄ matrix for mixed isotope water.⁹¹ The heavy water system shows a larger 3660 cm⁻¹ peak due to the contribution of both the d-OH of the cluster and of the HOD monomers. Consistently, the d-OD of this species is observed at 2689 cm⁻¹ that grows with added D_2O .

c) A Possible Model Tying the 3642 and 2835 cm⁻¹

The assignments above – 2835 cm⁻¹ to the dihydrate and 3642 cm⁻¹ to the OH- π of a larger water cluster – suggest that there are at least two species existing simultaneously in the solution. Consider, then, a model with a multi-acid hydrated cluster reservoir, breaking into smaller units as water is added. The size and nature of the units show an interesting aspect of the CCl₄ matrix, since injection of neat water only results in monomers but larger clusters could exist when aqueous acid solutions are injected. The existence of the dihydrate (2:1 water/acid) in a 6:1 water/acid injected solution would require the existence of some units with larger water/acid ratios. This bimodal distribution is consistent with the asymmetric fission model of a water-organic droplet argued by Donaldson, Tuck, and Vaida^{92,93} based on thermodynamics. The unexpected stability of the dihydrate is likely the driving force for its separation from the multi-acid multi-water cluster. Alternately, various sizes units may initially separate, then exchange with the monomer population, as suggested by the AIMD run of the trihydrate pTSA.W3B,

which then leads to an equilibrium distribution among cluster sizes in which dihydrates dominates for aforementioned reasons. The stability of the 2835 cm⁻¹ resonance suggests that the H stays with the sulfonate and does not exchange with D. Additions of water to the pool caused more dihydrates detachments, increasing the 2835 cm⁻¹ peak. D₂O is more efficient in this mechanism, thus the larger 2835 cm⁻¹ enhancement with D₂O addition.

The larger-water clusters are somewhat of a mystery, as there are multiple possible interpretations of them. Previous work⁶² mentioned that it could be a larger cluster detachment from the multi-acid multi-water cluster reservoir as a single acid multi-water cluster with a "scorpion tail" d-OH— π feature causing the 3642 cm⁻¹. Recent computational studies could not find a low energy cluster like one described above as *n* increases, therefore the existence of this cluster is not supported. Another model could be the shedding of a larger multi-acid cluster with a d-OH from a water molecule of one pTSA- $(H_2O)_n$ donating to the π ring of another pTSA- $(H_2O)_n$ cluster, resulting in the 3642 cm⁻¹ resonance. Yet the 3642 cm⁻¹ could also be the d-OH— π from the multi-acid multi-water reservoir. Although there are equivocal pictures on the structure of the larger acid cluster, it is unambiguous that the 3642 cm⁻¹ is the result of the d-OH— π resonance. Polarization from the donation increases the oscillator strength and a slight bias for a d-OH instead of a d-OD in the mixed isotope system coupled with enhanced couple splitting with D_2O leads to a nearly isotope independent growth of the 3642 cm⁻¹ resonance. Additional tuning of experimental methods are currently being conducted in order to elucidate the complete spectrum of pTSA-(H₂O)_n without impurities. This spectrum could provide more information about the system as a whole as well as the structure of the larger acid cluster.

4: Large (pTSA)•(H₂O)_n Clusters (n \ge 20)

An interesting question to ask about this system is if this type of bimodal distribution would persist at lower acid concentration (more water than acid). Another question is whether the H of the SO—H---OH₂ bond would exchange when dissolve with larger amount of D₂O and how much D₂O is needed. It is also of interest to study larger water clusters near the "magic" number n = 21, since it has interesting stability properties that could provide insights into clathrate hydrates formation mechanisms. Furthermore, most gas hydrates are big water clusters $(n \ge 20)$,²⁹ therefore it makes sense to go toward this direction for this project. This chapter discusses the findings of large pTSA•(H₂O)_n and pTSA•(D₂O)_n clusters and promising new leads at 3420 and 1717 cm⁻¹.

4.1 Experimenta Results of Large H₂O and D₂O Clusters Spectra

a) 1:26 pTSA-H₂O and 1:1:25 pTSA-H₂O-D₂O

Figure 4.1 shows the 1:26 pTSA•(H₂O) solution in CCl₄. The method for this experiment is the same as described above. There are some notable new features about this spectrum. First, the 2835 cm⁻¹ and the 3642 cm⁻¹ are not only present, but also extremely intense and sharp even at room temperature. The 3642 cm⁻¹ also seems like it has saturated. Several new features appear: a small "bump" at 3420 cm⁻¹, a broad H-bonded band at 3500 cm⁻¹, and some new features that overlap with the hydrocarbon impurities (covered but are discussed in Chapter 5).

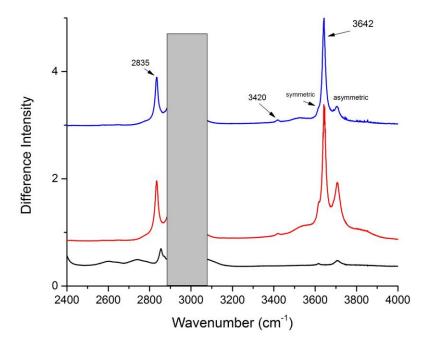


Figure 4.1 - Infrared spectra of pTSA and water in CCl_4 at room temperature (red) and at -20°C (blue). Spectra are difference spectra subtracting the room temperature, nominally dry CCl_4 spectrum. The black line is CCl_4 against cell background spectrum at room temperature, showing adventitious water. Note the intense 2835 cm⁻¹ and 3642 cm⁻¹ peaks along with the 3420 cm⁻¹ bump. Spectra are offset for clarity.

The intense and sharp 2835 cm⁻¹ and 3642 cm⁻¹ peaks are consistent with the previous small clusters studies. Previous studies show that both the 2835 and 3642 cm⁻¹ increase with increasing water concentration, thus these high intensity peaks are consistent with the trend. Nevertheless, it is quite unexpected that their intensities are this high. They imply large populations of both the dihydrate and d-OH— π clusters, both stable at room temperature and -20 °C. They also answer the question that the bimodal distribution persists at this acid/water ratio.

The 3420 cm⁻¹ and 3500 cm⁻¹ are H-bonded features. The 3420 cm⁻¹ implies the existence of the hydronium ion in solution. The vibrational feature of the hydronium ion is known to be

broad and has not been observed at ambient energy conditions. Thus, the 3420 cm⁻¹ feature is an important insight and is discussed in more details in Section 4.2. The overlapping features observed at the hydrocarbon impurities region are discussed in Chapter 5.

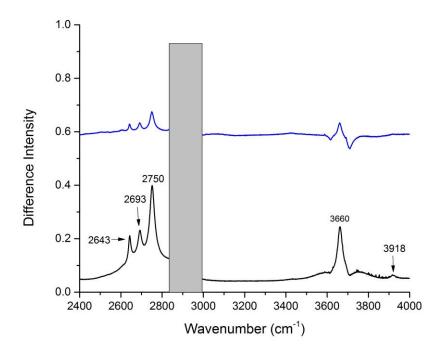


Figure 4.2 - Infrared spectra of pTSA and D_2O in CCl_4 at room temperature (black) and at -20 °C (blue). Spectra are difference spectra subtracting the room temperature, nominally dry CCl_4 spectrum. Spectra are offset for clarity.

Figure 4.2 shows the 1:1:25 (pTSA)-(H₂O)-(D₂O)₂₅ experiment. Two familiar resonances appear at 2693 and 3660 cm⁻¹, the OD and OH stretch of the HOD species, respectively. Additionally, the 2643 and 2750 cm⁻¹ resonances are also observed. These are the symmetric and asymmetric stretches of the D₂O species. No H₂O monomer features, beside its rotational wings, are observed in the OH-stretching region. This observation shows that enough D₂O have been added to exchange virtually all H₂O species to the HOD species with D₂O monomers leftover. When the system is cooled down to -20 °C, the 2835 and 3642 cm⁻¹, which are the SO—H---OH₂ stretch of the dihydrate and the OH--- π stretch of a bigger cluster, are not observed. This observation shows that the H in SO—H---OH₂ that is responsible for the 2835 cm⁻¹ had exchanged to become the D species. The same could be said about the 3642 cm⁻¹: its absence shows that the responsible H donation had become D. This is consistent with the model proposed in chapter 3.

A new feature shows up in this spectrum at 3918 cm⁻¹. This feature is assigned to be the first overtone of the S—O—D---OD₂ of the deuterium analogue of dihydrate, further evidence that deuterium exchange occurs at this acid-water-deuterium ratio. The lack of an O—D--- π feature implies that it is less favorable for the OD stretch to donate into the π ring. More experimental and theoretical work must be done to elucidate this phenomenon.

b) Experimental Results for the 1:51 pTSA-H₂O Cluster

Figure 4.3 shows the spectra for the 1:51 pTSA-H₂O experiment against room temperature CCl₄ background. The sample was taken at room temperature (black line) and -20 °C (red line). Per usual, the symmetric stretch, asymmetric stretch, and the rotational wings are observed at the free OH stretching region at room temperature. Several new observations are also observed in this experiment. The 3420 cm⁻¹, like the 1:26 pTSA-H₂O experiment, is also observed here at both temperatures, accompanies by a 3480 cm⁻¹ feature. When this sample is cooled to -20 °C, a very broad and structured H-bonded band is observed from 3000-4000 cm⁻¹. The 3420 and 3480 cm⁻¹ features ride on this broad band at -20 °C. The monomer symmetric and asymmetric stretches decreased significantly, but did not disappear completely. This observation signals the loss of monomers to the H-bonded cluster and also that the cluster is at its equilibrium size and thus some monomers are left over. The 2835 cm⁻¹ and 3642 cm⁻¹ can be observed, but they are very weak and can only be seen as shoulders and not sharp, distinct peaks.

The well-structured H-bonded feature is typical of the existence of large water clusters⁹⁴⁻⁹⁶ and ice.^{97,98} This large cluster is likely to have multiple hydration layer with 3- and 4-coordinate water, and perhaps this is the reason that the 3642 cm⁻¹ is quenched: there are many different d-OH on this cluster, thus multiple ways for the d-OH— π to interact which then leads to a broad distribution and therefore resulted in broader resonances instead of a sharp peak.

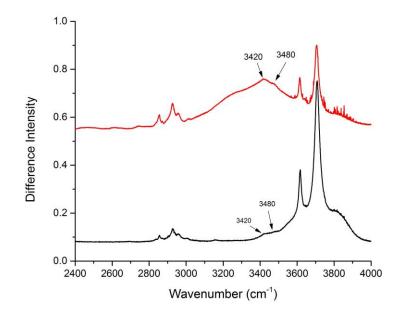


Figure 4.3 – Infrared spectra of the 1:51 pTSA-H₂O sample against room temperature CCl_4 spectrum. The sample was taken at room temperature (black line) and at -20 °C (red line). The features from 2850-3000 cm⁻¹are hydrocarbon impurities, which are discussed in more details in Chapter 5. Spectra are offset for clarity.

The lack of the 2835 cm⁻¹ and 3642 cm⁻¹ as well as the appearance of a much more intense H-bonded structure signify that the distribution of water clusters in this system no longer follows the bimodal theory proposed above. This result answers the question: does the bimodal distribution persists in bigger cluster system presented earlier in the chapter. It was determined, by experimental results, that this distribution persists at 1:26 pTSA:H₂O ratio. At 1:51, the bimodal distribution collapses, giving rise to a large cluster with multi-hydration layers. The threshold ratio at which this bimodal distribution collapsed and the structure as well as geometry of this multi-hydration cluster are still unknown. Experimental data with acid:water ratio between 1:26 and 1:51 are needed to see whether there is a pattern of gradient (i.e. a slow decrease of the 2835 and 3642 cm⁻¹ and a slow increase of the H-bonded band) or a sudden collapse at specific acid:water ratio. Theory could then be applied once a clear pattern, or lack thereof, could be established. Nevertheless, some observations from this experiment are consistent behaviors of larger clusters ($n \ge 20$): the 3420 cm⁻¹ "bump" and the 1717 cm⁻¹ sharp peak.

4.2 A Promising Lead for Observing the H_3O^+ at Ambient Energy: 3420 cm⁻¹ and 1717 cm⁻¹

3420 cm⁻¹. The small "bump" at 3420 cm⁻¹ is analogous to previous studies of salt ions and water in CCl₄.⁶¹ A small "bump" is also observed in this study at 3440 cm⁻¹ and is assigned the water-water bond of a dimer induced by the cation. With only a 20 cm⁻¹ difference and in the H-bond region, the 3420 cm⁻¹ is probably the water-water bond, but whether it is of the dihydrate or not is unclear. A similar peak at around 3450 cm⁻¹ is shown in the predicted spectrum of the dihydrate (see Appendix), which is the d-OH donating to one of the =O on the sulfonate group.

The region between 3400 and 3450 cm⁻¹ have shown up in multiple SFG studies on ice^{99,100} and spectroscopic studies on aqueous ionic solutions,^{101,102} to name a few. Despite the multiple reports, few have given a definitive assignment on this feature. On the SFG study of the prism face of ice,⁹⁹ Bisson and Shultz reported and assigned a resonance at 3415 cm⁻¹ to three coordinate water that has only a single donor to the 4-coordinate water in the bottom half of the top bilayer. Gopalakrishman *et al.*¹⁰² reasoned that the increase of the 3450 cm⁻¹ resonance in a salt ionic solution is from the increased ordering of the interfacial water molecules under the influence of a local weak electric field induced by dianions in aqueous Na₂SO₄ and (NH₄)₂SO₄

and/or an increase in the interfacial depth. Shultz and Allen *et al.*¹⁰¹ reported the same 3450 cm^{-1} in acidic solutions, claiming that its enhancement is caused by a short range polar ordering due to the negative ion at the interface.

The 3420 cm⁻¹ resonance falls in this range and is consistent with the aforementioned studies. It is only observed when the acid/water ratio decreases (more water than acid), similar to the "bulk" amount of water in ice and aqueous solutions. The aforementioned studies observed the 3400-3450 cm⁻¹ features at the water/air interface, which is a water/hydrophobic interface, analogous to that of water/CCl₄ interface. Therefore, there is precedent to assigning the 3420 cm⁻¹ feature as the 3-coordinate water molecules with a single H donating to a 4-coordinate water molecules at or near the interface.

This interpretation provides insights into the structure of the acid/water cluster. First, for the above interpretation to work, the cluster size has to be quite large, which is consistent with the acid/water ratio used for both aforementioned experiments (1:26 and 1:51 pTSA:H₂O). Second, increased ordering induced by the local electric field of the anion would require that the pTSA molecule in the cluster be ionized. According to theoretical calculations, as mentioned above, the acid proton would be ionized when n = 8. Although the exact threshold is experimentally unknown, the ratios of acid:water in the large cluster experiments are much larger than 8, thus the acid is expected to be ionized. Therefore, the interpretation that this peak was caused by the increased ordering due to the local electric field of the anion is consistent.

Enhanced ordering of the water on the interfacial surface could also be caused by the hydronium ion on the exterior. The hydronium ion is amphiphillic⁶⁶ with three covalently bonded hydrogen to the oxygen and one lone pair. In a bulk water solution, it is pushed to the exterior, similar to the hydrophobic effect.^{64,65} It has a solvation asymmetry, donating all three hydrogen

atoms when H-bonded to other water molecules while not accepting any H-bond, which in turns caused increased ordering of the water molecules in its vicinity. Thus, the rise of the 3420 cm⁻¹ could have been caused by the water molecule in the vicinity of the hydronium ion in the water/CCl₄ interface.

The above interpretations of the 3420 cm⁻¹ give us a simple picture: water is sandwiched between the ionized pTSA molecule and the H₃O⁺. This is consistent with the findings in Bisson and Shultz's studies of salts in CCl₄⁶¹ as well as our theoretical results. The 3420 cm⁻¹ also confirms the existence of the hydronium in solution and that the acid is ionized at 26 water to 1 acid solution. The threshold at which the acid is ionized is still experimentally unknown, although theoretical calculation suspected that ionization of pTSA occurs at n = 8.

1717 cm⁻¹. Figure 4.4 shows the 1717 cm⁻¹ resonance for the 1:26 and 1:51 $(pTSA) \cdot (H_2O)_n$ experiments. The 1717 cm⁻¹ for the 1:26 experiment is quite intense – to the point of saturation in the spectrum. The same peak for the 1:51 experiment is less intense, although is still quite sharp. Both are observed at room temperature and at -20 °C and are not observed in the small cluster experiments discussed in Chapter 3.

This 1717 cm⁻¹ peak is quite promising, for it falls in the region where bending modes of water in different bonding environment exist. Notably, resonances assigned to be the bending mode of the hydronium ion have been reported in the region between 1700 and 1800 cm⁻¹.¹⁰³ Since this resonance only is only observed in bigger clusters, it is consistent with the picture provided above from the interpretation of the 3420 cm⁻¹ where the hydronium ion is on the interface of water/CCl₄. Preliminary calculations suspect that this resonance might be due to the coupling of the stretching and bending motion of the Zundel cation. Many obstacles, both

computationally and experimentally, are still needed to be overcome before a definitive assignment of this peak.

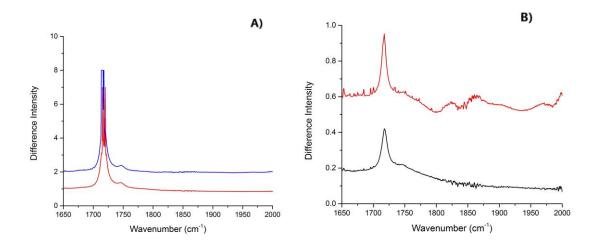


Figure 4.4 – A) The 1717 cm⁻¹ peak of pTSA-H₂O with ratio 1:26 against room temperature CCl₄ background spectrum. The sample was taken at room temperature (red line) and at -20°C (blue line). B) The 1717 cm⁻¹ peak of pTSA-H₂O with ratio 1:51 against room temperature CCl₄ background spectrum. Notice the different scales of A and B and that the 1717 cm⁻¹ in A is much more intense than in B. The reason for this is under investigation. Spectra are offset for clarity.

Experimentally, the region from 1650 to 2000 cm⁻¹ is where some unknown absorption features from CCl₄ are observed. These features could be seen in the cold spectrum of the 1:51 (Figure 4.4b), and they could be from the CCl₄ absorption or from hydrocarbon impurities. Therefore, "cleaning" this region would be ideal for more information. Computationally, the bending mode of the hydronium ion has not been studied extensively in the literature, thus it is difficult to claim any particular bending mode to be responsible for this feature. Additionally, due to the bimodal distribution nature of this system, it is impossible to know the size and structure of this cluster that is responsible for the 1717 cm⁻¹. Thus, more experimental data are needed. For example, CsOH could be added to test whether the 1717 cm⁻¹ is indeed a feature of an acidic species. CsOH is a base, thus is expected to balance the positive charge from the acid. Cs⁺ is a large and polarizable cation, thus is not suspected to be interacting with the system

significantly. If the 1717 cm⁻¹ diminished after the introduction of CsOH, then it could indeed be concluded to be an acidic feature. The 1717 cm⁻¹ is quite promising, for if it is a feature caused by the hydronium or Zundel cation, it would be the first time it is observed at ambient energy conditions.

5: Hydrocarbon Impurities

Hydrocarbon impurities are inherent in CCl₄, probably residues from its synthesis. Their features range from 2800-3100 cm⁻¹. Although these residues are minimal – CCl₄ use in the experiments above are 98.5% and 99.9% pure – their resonances can be seen in IR spectroscopy. Other methods besides FTIR have been used, namely H-NMR and C-NMR, without prevail. This section shows their FTIR features, some attempts of identifying them, and their implications in the presence of pTSA.

5.1 Hydrocarbon Impurities Experimental Spectra and What They Are Not

Figure 5.1 shows typical hydrocarbon impurities vibrational resonances from 2800-3000 cm⁻¹ (black line). A Lorentzian deconvolution of this spectrum shows at least five resonances: 2854 cm⁻¹, 2872 cm⁻¹, 2903 cm⁻¹, 2926 cm⁻¹, and 2958 cm⁻¹. These peaks could be from one molecule with multiple C—H stretches or multiple molecules with different C—H stretches. According to the U.S. National Library of Medicine,¹⁰⁴ CCl₄ is synthesized by reacting carbon disulfide with chlorine and a catalyst, chlorination of hydrocarboms (namely CH₄), or pyrolysis of perchloroethane (C₂Cl₆). The first reaction does not produce any hydrocarbon, but the second and third reactions do. They are shown below.

(1)
$$CH_4 + 2Cl_2 \rightarrow CCl_4 + H_2$$

Side Products: CHCl₃, CH₂Cl₂, CH₃Cl

Side Products: C₂HCl₅, C₂H₂Cl₄, ..., CH₅Cl

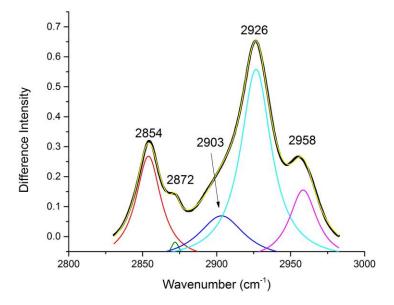


Figure 5.1 – Experimental FTIR spectrum of typical CCl_4 hydrocarbon impurities resonances (black line). Lorentzian fits deconvolution shows at least 5 resonances combined to form an excellent cumulative spectrum (yellow).

Although the second reaction does not produce hydrocarbon side products directly, the perchloroethane that was used for the reaction could have contained some side products. The first reaction produces the side products chloroform (CHCl₃), dichloromethane (DCM, CH₂Cl₂), and Chloromethane (CH₃Cl). These side products are easy to investigate: chloromethane is a gas at room temperature (boiling point -24.2 °C), therefore is not expected to be in solution. Chloroform and DCM are common solvents, thus were injected to CCl₄ solution with the same procedure as described above. Their peaks are shown in Figure 5.2. One C—H stretch peak is observed for chloroform at 3017 cm⁻¹ and two for DCM (symmetric, 2982 cm⁻¹, and asymmetric,

3049 cm⁻¹), as expected. These peaks, although are within the region, are not the exact peaks of the impurities. Therefore, they are ruled out.

The side products of the second reaction are more complicated, as they are more numerous. A literature search¹⁰⁵ or their resonances in CCl₄ as well as some pilot theoretical calculations show that indeed they are within the region, but no matches were found. Although the search for the identity(ies) of the hydrocarbon impurity(ies) was unsuccessful, it is confirmed that they are indeed hydrocarbon impurities, since they are within the hydrocarbon impurities region. Since the motivation for this project was to observed pTSA's role in promoting gas hydrate formation, it is important to observe how the impurity peaks are affected by the addition of pTSA and water and what those effects imply.

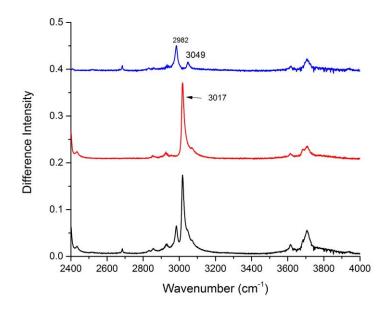


Figure 5.2 – FTIR spectra of DCM and chloroform against room temperature CCl₄ (black line), of only chloroform (red line), and of only DCM (blue line). Spectra were taken at room temperature

5.2 Implications of the Hydrocarbon Impurities in the Presence of pTSA

Figure 5.3a and 5.3b show the deconvoluted experimental spectra of the room temperature and cooled 1:6 pTSA/water from 2800-3200 cm⁻¹. These features were covered in Figure 3.1 for clarity. The features in Figure 5.3a are the same as the impurities features in Figure 5.1. Comparing Figure 5.3b to Figure 5.1, one could see that there are some subtle differences. The most obvious one is the 2835 cm⁻¹, as discussed above. A small shoulder (cyan line) also seem to occur at 2894 cm⁻¹. Perhaps the most pronounced differences are observed in the deconvolution of the peaks at the hydrocarbon stretches in the 1:26 experiment, shown in Figure 5.3c. These peaks are intense and sharp as well as being at completely different frequency than the impurities peaks, save the 2953 cm⁻¹ resonance. These resonances did not show up in previous experiments with only water, but shows up here after pTSA is added. Thus, there are reasons to believe that perhaps these are either pTSA-water cluster peaks or pTSA-waterhydrocarbon related peaks. The 2835 cm⁻¹, although are closed in frequency to these resonances, can be classified as not an impurity because it changes with the addition of water, whereas the impurities resonances only change with the initial addition of pTSA-water cluster. A method of cleaning the CCl₄ solution using TiO₂ is currently underway. Removing the hydrocarbon impurities would elucidate the spectrum more, thus provide more insight into the solvation mechanism of pTSA and water.

The increase in intensity of these hydrocarbon impurities resonances in the presence of pTSA is consistent with Gnanendran and Amin works^{16,17} on pTSA as a hydrate promoter as mentioned in the earlier chapters. They claim that pTSA improves the efficiency of gas hydrate formation by solubilizing more nonpolar gases (CH₄, for example) into the solution, thereby increasing the hydrate storage capacity. They also claim that the anionic species is more efficient at this task.¹⁶ This is also observed in the experimental data: the impurities peak for the salt

experiment at 2854 cm⁻¹ (Figure 3.3) is sharper and more intense than the ones in the other two experiments where pTSA and not Na⁺pTS⁻ was used.

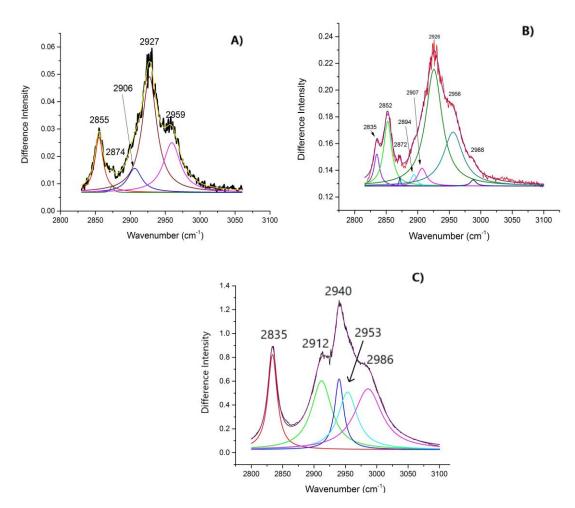


Figure 5.3 – A) Experimental data of the black line (room temperature 1:6 pTSA:water against CCl₄ background) covered by the gray bar in Figure 3.1. B) Experimental data of the red line (-20 °C 1:6 pTSA:water against CCl₄ background) covered by the gray bar in Figure 3.1. C) Experimental data of the black line (room temperature 1:26 pTSA:water against CCl₄ background) covered by the gray bar in Figure 4.1. All three spectra are deconvoluted by Lorentzian fits.

Consider a system where the hydrocarbon impurities concentration is high enough so that they phase separated to another layer not in the CCl₄ solution and only a small, saturated amount is left. This small, saturated amount gave the original peaks, which would be canceled out from the background subtraction. The addition of pTSA, then, solubilizes more hydrocarbon impurities from the phase separated layer, bringing more of them into the CCl₄ solution, thus increase the intensity of the impurities resonances. The anionic species is more efficient at this, therefore more impurities are solubilized and thus the peaks are sharper. To test this hypothesis, SFG could be use to determine if an interface between some hydrocarbon and CCl₄. This interpretation is a promising insight, since it shows pTSA potential as a gas hydrate promoter. Eventually, perhaps gas hydrate formation mechanism could be studied using this system by employing a known hydrocarbon, such as methane.

6: Summary and Future Works

Natural gas hydrate is a potential new technology that could be used to remove CO₂ from the atmosphere or as a medium for gas storage and transport. Current manufacturing methods of natural gas hydrates are not cost efficient, since gas hydrates require a high pressure and low temperature environment to form. Even then, the production rate is too slow. Therefore, gas hydrate promoters, which increase the production rate thereby improve cost-effectiveness, are of great interests. Among the promoters, pTSA have been studied on a macroscopic level and concluded to be more efficient than surfactants.^{16,17} This project studies pTSA's solvation mechanism in water on a molecular level using a CCl₄ matrix and FTIR spectroscopy in an attempt to give some insights into how pTSA works as a gas hydrate promoter.

Since water exists as monomers in CCl₄,¹ only the symmetric and asymmetric stretches as well as the rotational wing about the symmetry axis can be observed at the free-OH stretching region. Monitoring this picture after adding a molecule of interest could give insight into the molecule's interaction with water. Experimental results for pTSA small clusters ($n \le 10$) found a few new features arising at -20 °C: 2835 cm⁻¹ resonance, 3642 cm⁻¹ resonance, and a small but structured H-bonded region. Through complex computational methods, the 2835 cm⁻¹ is assigned the SO—H---OH₂ of the dihydrate species where the acidic proton is not yet ionized. The 3642 cm⁻¹ is assigned the OH--- π . The cluster that is responsible for the 3642 cm⁻¹ is expected to be large ($n \ge 12$), although the specific size and geometry of this cluster is unknown. This cluster could also be a multi-acid multi-water cluster. The existence of both the dihydrate and larger cluster give rise to the possible asymmetric fission model proposed by Donald, Tuck, and Vaida.^{92,93}

A study of larger cluster ($n \ge 20$) also contains the above features with some new features: a small "bump" at 3420 cm⁻¹ and a sharp peak at 1717 cm⁻¹. The region between 3400 and 3450 cm⁻¹ has been observed in many SFG and ice studies^{99,100} as well as in aqueous solutions^{61,101,102}. It is assigned in these studies to be the feature of a 3-coordinate water on the top layer at the air-liquid interface donating to the second layer of the water H-bonded system with one weak H-bond. The rise of this resonance is caused by an increased in ordering of the water H-bonded network caused by the local electric field of the anion. The increase in ordering could also be caused by having the hydronium ion at the interface.⁶⁶ The hydronium ion is amphiphilic, thus have an asymmetric solvation mechanism. Therefore, the water around the hydronium can only accepts H-bond donated by it. The water in its vicinity then becomes a 3-coordinate water (DDA) and donate to the bottom half layer at the interface. Thus, the existence of this peak is evidence that the hydronium is in the solution, meaning the acid is ionized.

The 1717 cm⁻¹ is also another piece of potential evidence for the existence of the hydronium in solution. The bending mode of the hydronium ion has been reported to be from 1700 to 1800 cm^{-1} ,¹⁰³ although not at ambient energy condition. If the 1717 cm⁻¹ is indeed the

bending mode of the hydronium ion, then this study would be the first to report a hydronium feature at ambient energy condition. Unfortunately, the bending mode of the hydronium ion is difficult to model computationally. Furthermore, the structure, size, and geometry of the big cluster, where the hydronium ion supposedly exist, are still unknown. Therefore, more experimental and computational results are needed for definitive assignments. Nonetheless, the 3420 cm⁻¹ and 1717 cm⁻¹ give rise to a little glimpse into what the large cluster looks like: water is sandwiched between the anionic pTS⁻ species and the cationic hydronium ion. This picture is consistent with the works done on ionic salts reported by Bisson and Shultz,⁶¹ where they observed a similar feature at 3440 cm⁻¹.

Some efforts have been given to identify the hydrocarbon impurities that give rise to the features from 2850-3000 cm⁻¹. Although the conclusion was not definitive, the behaviors of these features show pTSA's potential as a promoter. Gnanendran and Amin reported that pTSA, a hydrotrope, is more efficient than surfactants as a gas hydrate promoter because it can solubilize more hydrophobic species, thus increase the storage capacity of gas hydrate.^{16,17} They also reported that the anionic species is more efficient at this task. Since the experiments were done with background subtractions, an increase in intensity of the hydrocarbon impurities peaks after the addition of pTSA means that more of those molecules are brought into solution as a result of adding pTSA. This increase is especially more intense in the salt experiment. Thus, there is reason to believe that there is another layer of hydrocarbon impurities on or below CCl₄, and pTSA solubilize them into solutions. This result is consistent with Gnanendran and Amin's works and is promising for future works with hydrocarbon-pTSA-water systems.

A project is currently being developed to clean CCl₄ of the hydrocarbon impurities with TiO₂. This work takes precedent, since more information could be extracted at that region. After

CCl₄ is cleaned, an experiment could be conducted where a hydrocarbon of interest is introduced to the solution so that its molecular interactions with pTSA-water could be elucidated. This experiment has the potential to impact the petroleum industry, since the promoter-hydrate formation mechanism is still lacking. An experiment where CsOH is added to a system where the 1717 cm⁻¹ appears could be performed. CsOH is a base, and Cs⁺ is polarizable since it is a big ion therefore is not expected to interact with the system. If the 1717 cm⁻¹ is quenched after the addition of CsOH, then it is indeed an acidic feature. Another interesting experiment could be to study pTSA-water in CCl₄ at the interface using SFG. This could potentially give more insight to the orientation of pTSA-water clusters at the interface with CCl₄, and perhaps give more information on the hydronium features at the interface.

Appendix

- 1. Spectra of CCl₄ at room temperature and at -20 °C Figure S 1
- 2. Spectra of CCl₄ with no acid-water: both 99.5% and 99.9% pure Figure S 2
- 3. Selected clusters from Table I and illustrated in Figure 4 coordinates Table S 1
- 4. Predicted stretch spectrum of the dihydrate Figure S 3
- 5. A preliminary look at the case of n=7 water finds frequent interchange among the top four geometries (see Figures 4(e), 4(f)) but no proton transfers that affect the embedded H₃O⁺.

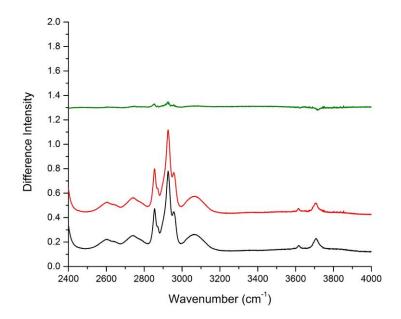
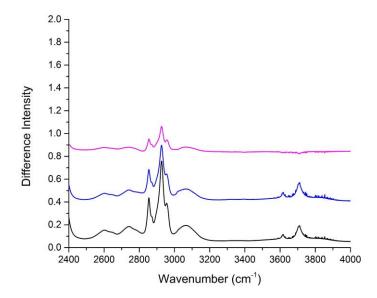


Figure S 2. 99.5% pure CCl₄ at room temperature (black line) and -20 °C (red line) along with the difference spectrum (green line).



*Figure S 1. 99.5% pure CCl*⁴ (black line), 99.9% pure CCl⁴ (blue line), and the difference between the two (magenta line) at room temperature.

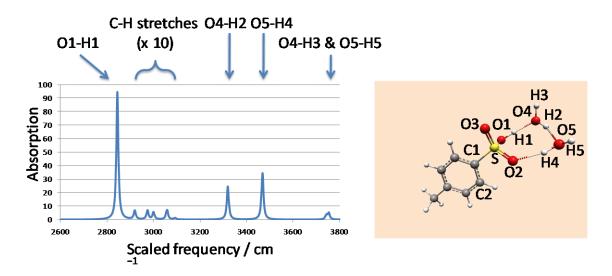


Figure S 3: Stretch region of predicted IR spectrum of $(pTSA)(H_2O)_2$. Computed frequencies are scaled by 0.965. Though plotted here as if all peaks have width 10 cm⁻¹ at half-height, we propose that the water-O stretches at 3320 and 3469 cm⁻¹ are not seen experimentally because they are substantially broader than the (S-)O1-H1 stretch at 2844 cm⁻¹. Figure 4(a), containing the labeling of O's and H's, is reproduced here for the reader's convenience.

pTSA.W1

с	2.299258	-0.165678	-0.362763
с	1.565039	-0.231573	0.826572
h	2.085338	-0.217096	1.778121
с	0.177151	-0.319187	0.812870
h	-0.383270	-0.382409	1.736834
с	-0.484035	-0.338709	-0.412119
с	0.217805	-0.278606	-1.613842
h	-0.316102	-0.308778	-2.554976
с	1.604384	-0.190416	-1.578027
h	2.155913	-0.142747	-2.510860
с	3.805872	-0.101591	-0.339756
h	4.237374	-1.101772	-0.454423
h	4.190303	0.514779	-1.155441
h	4.173869	0.309418	0.602229
S	-2.269551	-0.428296	-0.455030
0	-2.722930	1.127561	-0.424035
0	-2.676026	-0.963732	-1.738764
0	-2.736518	-1.042721	0.796065
h	-3.069109	1.353925	0.479271
о	-3.668971	1.218118	2.113840
h	-3.610266	0.248375	2.106879

h -4.575529 1.441136 2.347354

25

pTSA.W2A

- c 3.127038 -0.176552 -0.212190
- c 2.363780 -0.022151 0.952158
- h 2.863448 0.120776 1.904230
- c 0.975401 -0.049665 0.911681
- h 0.389487 0.060527 1.815223
- c 0.343188 -0.231245 -0.316204
- c 1.071106 -0.389127 -1.491356
- h 0.555177 -0.537121 -2.431367
- c 2.460498 -0.359286 -1.428434
- h 3.034666 -0.480316 -2.340595
- c 4.633925 -0.176099 -0.149602
- h 5.012470 -1.182359 0.060115
- h 5.072320 0.148958 -1.095173
- h 4.999627 0.481997 0.641500
- s -1.437893 -0.264915 -0.393427
- o -1.743928 1.308565 -0.382285
- o -1.953951 -0.850181 0.856974
- o -1.865026 -0.855006 -1.655657
- h -2.731584 1.487362 -0.253289
- o -4.305037 1.687532 0.048273
- h -4.602059 0.977046 0.662001
- h -4.910702 1.686067 -0.699060
- o -4.558521 -0.429651 1.727716
- h -3.658423 -0.756304 1.533697
- h -5.135007 -1.198853 1.735072

pTSA.W3A

c 3.713522 -0.361455 -0.279360

- c 2.831550 -0.069020 -1.326465
- h 3.211457 0.011524 -2.339205
- c 1.475581 0.128383 -1.092352
- h 0.798793 0.369232 -1.902007
- c 0.998089 0.029294 0.211688
- c 1.849133 -0.252185 1.276594
- h 1.456749 -0.304605 2.284159
- c 3.202394 -0.443974 1.020978
- h 3.872694 -0.655969 1.846981
- c 5.177934 -0.602159 -0.548427
- h 5.795159 -0.291969 0.297344
- h 5.368838 -1.666918 -0.720758
- h 5.515180 -0.061161 -1.434853
- s -0.738260 0.277786 0.537702
- o -1.295662 -1.186922 0.243627
- o -1.262500 1.206090 -0.479070
- o -0.940128 0.616668 1.943447
- h -2.258107 -1.299244 0.573006
- o -3.719122 -1.488093 1.057049
- h -4.389280 -0.917213 0.602603
- h -3.845542 -1.360712 2.002728
- o -5.420673 0.186913 -0.168671
- h -5.962000 -0.088073 -0.914171
- h -4.910170 0.975766 -0.466950
- o -3.800501 2.230837 -0.921585
- h -3.842419 3.099294 -0.510654
- h -2.882707 1.917885 -0.793378

pTSA.W3B

с	3.553727	-0.156419	0.000000
с	2.838353	-0.178309	1.202984
h	3.375423	-0.162883	2.145174
с	1.448930	-0.220932	1.214211
h	0.898556	-0.247635	2.145800
с	0.768075	-0.239468	0.000000
с	1.448930	-0.220932	-1.214211
h	0.898556	-0.247635	-2.145800
с	2.838353	-0.178309	-1.202984
h	3.375423	-0.162883	-2.145174
с	5.061769	-0.141235	0.000000
h	5.458565	-1.162191	0.000000
h	5.455341	0.363073	-0.884884
h	5.455341	0.363073	0.884884
s	-1.013666	-0.270982	0.000000
0	-1.352509	1.273739	0.000000
0	-1.479065	-0.880726	1.255895
0	-1.479065	-0.880726	-1.255895
h	-2.397153	1.434814	0.000000
0	-4.090446	-0.197973	2.133167
h	-3.208945	-0.590714	1.997912
h	-4.186010	-0.054569	3.078825
0	-3.854247	1.532238	0.000000
h	-4.164417	1.035420	0.786124
h	-4.164417	1.035420	-0.786124
0	-4.090446	-0.197973	-2.133167
h	-3.208945	-0.590714	-1.997912
h	-4 186010	-0 054569	-3 078825

h -4.186010 -0.054569 -3.078825

pTSA.W3C

С	3.597551	-0.136939	-0.033033
с	2.859308	-0.836272	0.928622
h	3.377809	-1.427421	1.675604
с	1.470010	-0.782999	0.949658
h	0.901349	-1.314227	1.702004
с	0.814572	-0.015605	-0.008965
с	1.518315	0.699089	-0.974958
h	0.986694	1.304355	-1.698053
С	2.906514	0.632427	-0.977021
h	3.462258	1.191895	-1.721669
С	5.102678	-0.225323	-0.065572
h	5.427167	-1.028639	-0.735767
h	5.548907	0.703032	-0.428269
h	5.511755	-0.438280	0.923989
S	-0.965388	0.060880	-0.003027
0	-1.408263	1.313510	-0.625309
0	-1.444929	-0.219221	1.360040
0	-1.312951	-1.165626	-0.954294
h	-2.305745	-1.175045	-1.221572
0	-4.070134	0.544425	2.086919
h	-3.144389	0.249714	2.006790
h	-4.123718	1.052594	2.901220
0	-4.391995	1.290074	-0.587843
h	-3.523896	1.699745	-0.719617
h	-4.450050	1.162409	0.380079
0	-3.798639	-1.107053	-1.603308

- h -4.159298 -0.236017 -1.280720
- h -4.385495 -1.795482 -1.275928

pTSA.W3D

С	2.789409	2.333818	0.543405
С	1.735798	1.982123	1.395413
h	1.526225	2.593042	2.266956
с	0.953060	0.860234	1.147513
h	0.146892	0.584267	1.815289
с	1.229570	0.080891	0.027628
с	2.271753	0.401416	-0.837400
h	2.474160	-0.227722	-1.694748
с	3.042408	1.528434	-0.572699
h	3.855299	1.784213	-1.243946
с	3.653767	3.534378	0.838526
h	4.006896	4.006590	-0.080827
h	3.112403	4.282553	1.421082
h	4.537092	3.242969	1.417040
S	0.235392	-1.363941	-0.312811
0	-0.989998	-0.693728	-1.078463
0	-0.249722	-1.906680	0.970779
0	0.974082	-2.254026	-1.198532
h	-1.823101	-1.303953	-1.111590
0	-2.476428	-3.564545	1.178440
h	-2.254410	-4.500081	1.199536
h	-1.630253	-3.084479	1.267451
0	-3.060965	-2.189398	-1.113875
h	-3.050969	-2.769358	-0.321736

- h -3.925974 -1.745824 -1.156365
- o -5.529657 -0.832023 -1.401576
- h -5.930361 -0.852558 -2.276791
- h -5.622366 0.073386 -1.087699

pTS-.H+W3A

С	3.560577	-0.127283	-0.040154
с	2.853870	-0.189736	1.165682
h	3.397972	-0.212230	2.103980
с	1.463438	-0.225973	1.185304
h	0.918916	-0.285414	2.118784
с	0.770778	-0.198314	-0.021299
с	1.445259	-0.137829	-1.237976
h	0.888504	-0.129349	-2.166275
с	2.834971	-0.101547	-1.236940
h	3.364582	-0.054412	-2.182567
с	5.069147	-0.120568	-0.052193
h	5.461581	-1.142023	-0.103433
h	5.458420	0.422958	-0.915748
h	5.473036	0.340303	0.851717
S	-1.016819	-0.199207	-0.011852
0	-1.417866	1.274986	-0.066362
0	-1.457946	-0.814793	1.266869
0	-1.470853	-0.909378	-1.234351
0	-3.982434	-0.080109	2.022213
h	-3.082468	-0.465821	1.908936
h	-4.079303	0.149317	2.951013
0	-3.852310	1.499703	-0.057906

- h -4.077306 0.997374 0.782529
- h -2.746973 1.493679 -0.072397
- h -4.102851 0.909055 -0.833719
- o -3.986016 -0.190681 -2.043665
- h -3.091608 -0.580111 -1.907243
- h -4.596297 -0.922595 -2.172948

pTS-.H+W4A

С	4.029641	0.050271	-0.001018
С	3.313914	-0.042414	1.197695
h	3.851432	-0.086316	2.139266
с	1.923495	-0.082349	1.206157
h	1.374480	-0.166240	2.135250
с	1.236028	-0.027570	-0.002988
с	1.921002	0.064488	-1.211300
h	1.369508	0.093744	-2.142293
с	3.311553	0.103178	-1.200678
h	3.847177	0.173607	-2.141724
с	5.538663	0.059742	-0.001934
h	5.934900	-0.959528	-0.067268
h	5.933277	0.618952	-0.853120
h	5.935081	0.506543	0.912477
S	-0.558230	-0.014268	-0.000539
0	-0.976584	1.417616	0.104867
0	-0.976519	-0.812619	1.191497
0	-0.977233	-0.630354	-1.296137
0	-3.479057	-1.104882	1.987134
h	-2.509852	-1.035888	1.765308

- h -3.586769 -0.806537 2.895135
- o -3.475774 2.240100 -0.044807
- h -2.504653 2.016223 0.008553
- h -3.582403 2.889224 -0.746463
- o -3.473349 -1.204550 -1.933968
- h -2.504991 -1.036547 -1.767873
- h -3.572664 -2.143212 -2.119068
- o -4.620127 -0.019856 -0.004384
- h -4.244363 -0.446200 0.853844
- h -4.238543 0.936604 -0.061403
- h -4.239039 -0.550962 -0.800217

pTS-.H+W4B

С	3.931820	-0.731404	-0.421498
с	2.998535	-1.588747	-1.012622
h	3.343146	-2.399061	-1.646475
с	1.632799	-1.424638	-0.800627
h	0.913114	-2.096848	-1.249531
с	1.194566	-0.386583	0.015004
с	2.100362	0.482251	0.619016
h	1.742467	1.276349	1.261861
с	3.460385	0.304103	0.394523
h	4.168098	0.979415	0.863920
с	5.412205	-0.935290	-0.629842
h	5.615433	-1.450106	-1.571127
h	5.948013	0.016610	-0.639116
h	5.838469	-1.542562	0.176099
S	-0.561402	-0.134567	0.265410

- o -0.728957 0.327683 1.664887
- o -1.234103 -1.426817 -0.045057
- o -0.961720 0.940623 -0.727131
- o -3.690978 -1.402882 -1.159540
- h -2.816741 -1.583518 -0.735421
- h -3.790734 -2.022552 -1.887966
- o -3.363047 1.155753 -1.317404
- h -2.328086 1.120384 -1.094375
- h -3.645658 0.185828 -1.352194
- h -3.838045 1.624950 -0.543994
- o -4.452913 2.257984 0.717759
- h -3.952038 1.975106 1.530348
- h -4.557899 3.212833 0.766592
- o -2.962172 1.425303 2.751230
- h -2.124350 1.040093 2.403054
- h -3.290570 0.800308 3.404217

pTSA.W4A

С	4.042125	0.038675	-0.001049
С	3.318806	0.838430	0.891833
h	3.849981	1.479744	1.586763
С	1.928790	0.832462	0.900897
h	1.372297	1.462119	1.583217
С	1.255719	0.012226	-0.000651
С	1.943824	-0.789207	-0.907205
h	1.398174	-1.404181	-1.611165
с	3.333966	-0.767661	-0.899578

h 3.876798 -1.384262 -1.607734

- c 5.550072 0.030877 0.018783
- h 5.922601 -0.729415 0.713750
- h 5.960942 -0.196919 -0.966848
- h 5.949931 0.994062 0.342286
- s -0.526903 -0.010802 0.005374
- o -0.803125 -1.150505 1.061798
- o -0.997465 -0.427259 -1.324084
- o -1.015371 1.279522 0.514969
- h -1.818026 -1.333980 1.248027
- o -3.222743 -1.657600 1.523566
- h -3.880439 -1.107030 0.991509
- h -3.502771 -1.629808 2.443390
- o -4.791175 -0.156987 0.079720
- h -4.547349 -0.295193 -0.857161
- h -4.560250 0.773888 0.270923
- o -3.603482 -0.485685 -2.426194
- h -2.670302 -0.471183 -2.146819
- h -3.692589 -1.235037 -3.021833
- o -3.641959 2.307733 0.714702
- h -2.703568 2.052832 0.654928
- h -3.726510 3.130141 0.223887

pTSA.W4B

С	4.230237	-0.454267	-0.133676
С	3.513260	0.124246	-1.186693
h	4.026079	0.375139	-2.108931
с	2.153525	0.393162	-1.071165
h	1.605139	0.855451	-1.881558

- c 1.503859 0.076106 0.118019
- c 2.190562 -0.492270 1.188191
- h 1.671745 -0.709733 2.113140
- c 3.549154 -0.751784 1.052991
- h 4.090807 -1.186472 1.886203
- c 5.698407 -0.767884 -0.277876
- h 6.224641 -0.658262 0.672828
- h 5.841707 -1.800396 -0.614334
- h 6.176165 -0.114534 -1.010549
- s -0.244880 0.400044 0.286214
- o -0.904035 -0.955770 -0.221590
- o -0.586135 1.464015 -0.667704
- o -0.573124 0.587493 1.700131
- h -1.626053 -1.282269 0.446207
- o -2.643030 -1.665654 1.488453
- h -3.580912 -1.454831 1.247752
- h -2.435274 -1.125511 2.260702
- o -5.153108 -0.979702 0.841435
- h -5.764576 -1.589528 0.419458
- h -5.237586 -0.120283 0.364112
- o -3.051977 2.313451 -1.653913
- h -2.193686 2.050470 -1.267237
- h -2.945423 2.225855 -2.605508
- o -5.326990 1.386471 -0.466457
- h -5.700943 2.147319 -0.012787
- h -4.507554 1.709926 -0.905857

pTSA.W4C

- c 3.925307 -0.824580 -0.458835
- c 3.240246 0.084554 -1.272498
- h 3.765519 0.579496 -2.082070
- c 1.894250 0.365989 -1.064247
- h 1.363880 1.062765 -1.700683
- c 1.228234 -0.275596 -0.024292
- c 1.879067 -1.188151 0.802389
- h 1.337855 -1.682767 1.598722
- c 3.224888 -1.452500 0.578580
- h 3.739324 -2.160089 1.219847
- c 5.377042 -1.146132 -0.710859
- h 5.883615 -0.327579 -1.225681
- h 5.908548 -1.344753 0.222368
- h 5.472044 -2.039092 -1.338032
- s -0.496158 0.082294 0.254136
- o -1.126002 -1.106498 0.847541
- o -1.088843 0.611019 -0.980996
- o -0.386345 1.230169 1.338773
- h -1.249185 1.786867 1.465280
- o -2.418584 2.701596 1.739452
- h -2.251626 3.632953 1.565027
- h -3.294271 2.487741 1.312488
- o -4.686369 2.156350 0.491040
- h -5.474353 1.867263 0.959912
- h -4.516047 1.483744 -0.220337
- o -3.993176 0.323304 -1.343567
- h -4.029016 -0.566486 -0.940847
- h -3.045549 0.479334 -1.480214
- o -3.703482 -2.039208 0.139214

- h -2.814834 -1.834598 0.482359
- h -3.665977 -2.947406 -0.173970

pTS-.H+W5A

С	4.468412	-0.130471	0.082669
с	3.750525	-0.043204	1.279450
h	4.285507	-0.013233	2.222990
с	2.359108	-0.001529	1.285199
h	1.805865	0.051239	2.214134
с	1.673786	-0.050774	0.075395
с	2.362248	-0.144770	-1.131633
h	1.813049	-0.203729	-2.062694
с	3.751981	-0.184992	-1.118882
h	4.289218	-0.266972	-2.058207
с	5.977481	-0.146367	0.081806
h	6.369200	-0.779506	-0.717829
h	6.377693	0.861433	-0.074042
h	6.373340	-0.513196	1.031109
S	-0.119273	0.052391	0.068644
0	-0.567458	-0.473245	1.387061
0	-0.571966	-0.783374	-1.086976
0	-0.454971	1.497355	-0.119773
0	-2.809022	-0.575409	-2.448956
h	-1.934890	-0.664749	-1.974522
h	-3.032676	-1.440941	-2.802871
0	-2.804239	2.651313	-0.213899
h	-1.891745	2.252827	-0.123629
h	-2.704955	3.472059	-0.705570

- o -2.849189 -1.653924 2.143255
- h -2.005046 -1.210602 1.880987
- h -2.672468 -2.598420 2.112190
- o -4.988124 -0.598555 1.031665
- h -4.200658 -1.025982 1.473700
- h -5.544288 -0.238449 1.728812
- o -4.343370 0.811370 -0.988647
- h -3.783864 0.209872 -1.610020
- h -4.628365 0.274092 -0.163772
- h -3.750846 1.608445 -0.697144

pTS-.H+W5B

С	4.449152	-0.253818	0.093546
с	3.728491	-0.571157	1.248205
h	4.260936	-0.847203	2.152286
с	2.336113	-0.540032	1.261852
h	1.777307	-0.791412	2.154049
с	1.657602	-0.184687	0.101624
с	2.348178	0.136952	-1.065300
h	1.805179	0.405650	-1.962693
с	3.736770	0.100814	-1.059554
h	4.277729	0.352048	-1.965973
с	5.956568	-0.318382	0.076802
h	6.300795	-1.279982	-0.319245
h	6.381728	0.464973	-0.555047
h	6.371658	-0.209021	1.080727
S	-0.132324	-0.113340	0.112209
0	-0.600403	-0.789062	-1.134246

- o -0.465837 1.362253 0.084722
- o -0.593483 -0.773668 1.357940
- o -3.338299 -1.274163 1.449266
- h -2.363971 -1.228729 1.540833
- h -3.478539 -1.554987 0.530450
- o -4.150590 1.217632 1.804586
- h -3.912746 0.247454 1.761225
- h -4.121616 1.484165 2.728134
- o -2.774616 2.244589 -0.087372
- h -3.298498 1.920940 0.727831
- h -1.783780 1.874503 -0.026877
- h -3.228128 1.851136 -0.896525
- o -3.130110 -1.513875 -1.493519
- h -3.209982 -2.301254 -2.041183
- h -2.157319 -1.325467 -1.410498
- o -3.964646 1.026460 -2.069856
- h -3.769386 0.065045 -1.982195
- h -4.913935 1.115624 -2.196203

pTS-.H+W5C

- c 4.453865 0.466203 -0.226856
- c 3.819652 0.262155 1.004226
- h 4.413434 0.223040 1.911525
- c 2.440504 0.106252 1.087491
- h 1.955647 -0.062385 2.040603
- c 1.681883 0.157697 -0.079135
- c 2.283668 0.359109 -1.317472
- h 1.677207 0.384312 -2.213773

- c 3.665247 0.511956 -1.381242
- h 4.137238 0.667822 -2.345660
- c 5.954480 0.601056 -0.309010
- h 6.428827 -0.381330 -0.409294
- h 6.254359 1.199010 -1.172206
- h 6.362176 1.070374 0.589079
- s -0.101027 0.006275 0.021239
- o -0.608183 1.402466 0.310959
- o -0.386162 -0.921310 1.148301
- o -0.574927 -0.483078 -1.302186
- o -2.418206 -2.689267 1.257897
- h -1.655154 -2.061728 1.249471
- h -2.064841 -3.531047 0.955447
- o -4.561694 1.016189 1.410558
- h -4.355326 0.975948 2.349677
- h -4.634368 0.089573 1.092715
- o -2.919965 2.021688 -0.379699
- h -3.533589 1.779715 0.376358
- h -1.933331 1.823733 -0.067723
- h -3.099021 1.303868 -1.106318
- o -3.193883 0.042366 -1.949446
- h -2.266318 -0.280587 -1.906542
- h -3.716966 -0.599653 -1.436226
- o -4.489196 -1.447063 0.155684
- h -3.754681 -1.995044 0.549196
- h -5.261349 -2.018315 0.092360

pTS-.H+W5D

- c 4.204375 0.272620 -0.006562
- c 3.453331 0.274374 1.175117
- h 3.963305 0.280443 2.132764
- c 2.063557 0.267130 1.145728
- h 1.489605 0.259440 2.063717
- c 1.411772 0.257975 -0.085555
- c 2.131103 0.257500 -1.276806
- h 1.605598 0.241895 -2.223046
- c 3.522020 0.264659 -1.227399
- h 4.085061 0.263229 -2.154805
- c 5.712288 0.248523 0.038199
- h 6.145006 0.700134 -0.856893
- h 6.093582 0.785489 0.909573
- h 6.083563 -0.780292 0.099575
- s -0.379512 0.310388 -0.136176
- o -0.789261 1.730729 0.053603
- o -0.855985 -0.545469 1.014252
- o -0.788676 -0.249856 -1.454921
- o -3.517007 -0.057926 -1.841027
- h -2.537585 -0.133133 -1.906695
- h -3.665249 0.844443 -1.501409
- o -4.121755 -1.575983 0.086294
- h -3.929259 -1.008049 -0.769986
- h -3.362497 -2.243532 0.181550
- h -3.977740 -0.902927 0.860151
- o -3.484512 0.171373 1.823766
- h -2.514795 0.035484 1.810025
- h -3.617229 1.033398 1.386485
- o -3.412292 2.244335 -0.132606

- h -3.680196 3.166835 -0.186679
- h -2.423187 2.229096 -0.096704
- o -1.982909 -2.954635 0.513534
- h -1.530336 -3.504910 -0.133044
- h -1.394181 -2.182780 0.695980

pTSA.W5A

с	4.460746	0.506360	-0.191526
с	3.863270	0.028530	0.980280
h	4.478956	-0.178608	1.848937
с	2.491480	-0.187289	1.054009
h	2.032220	-0.564676	1.958495
с	1.707982	0.083605	-0.064148
с	2.270859	0.559418	-1.245393
h	1.645133	0.754780	-2.106834
с	3.644245	0.766884	-1.298463
h	4.088958	1.137135	-2.215857
с	5.953525	0.707016	-0.269258
h	6.452881	-0.217911	-0.577465
h	6.214892	1.478699	-0.996079
h	6.367596	0.995032	0.699302
S	-0.057377	-0.160636	0.027213
0	-0.573949	1.256829	0.421445
0	-0.324755	-1.131286	1.099691
0	-0.551645	-0.518347	-1.320416
h	-1.610845	1.505589	0.122578
0	-2.614583	-2.661459	1.630530
h	-1.791329	-2.156795	1.491845

- h -2.369750 -3.589825 1.583798
- o -4.813862 1.101282 1.362378
- h -4.844421 1.106005 2.323021
- h -4.868627 0.155089 1.073055
- o -2.844064 1.852524 -0.259845
- h -3.541452 1.690983 0.431888
- h -3.104763 1.301194 -1.023509
- o -3.185629 -0.289431 -2.044702
- h -2.238356 -0.514348 -1.926309
- h -3.378657 -0.367333 -2.984096
- o -4.667489 -1.359749 0.274468
- h -4.007354 -1.913468 0.738175
- h -4.283836 -1.175794 -0.597210

pTSA.W5B

С	4.510079	-0.197939	-0.032567
С	3.784204	-0.954543	0.894102
h	4.312719	-1.574900	1.609822
С	2.393852	-0.926968	0.916030
h	1.834209	-1.507375	1.638171
с	1.723998	-0.129010	-0.006620
с	2.415789	0.637091	-0.941505
h	1.874246	1.257048	-1.644539
С	3.804884	0.594474	-0.946619
h	4.350168	1.187375	-1.673007
с	6.018194	-0.211486	-0.028182
h	6.420938	-0.055393	-1.031135
h	6.410364	0.587173	0.610709

- h 6.407364 -1.157507 0.353494
- s -0.059555 -0.097752 -0.003270
- o -0.505516 1.220397 -0.480314
- o -0.529403 -0.519495 1.325413
- o -0.379054 -1.200934 -1.075358
- h -1.404715 -1.333937 -1.319031
- o -3.044783 -1.233630 2.375492
- h -2.149060 -0.984322 2.083837
- h -3.190890 -0.779483 3.210046
- o -2.748782 -1.582336 -1.722709
- h -3.469397 -1.341296 -1.048029
- h -2.926476 -2.478238 -2.024667
- o -2.715276 2.945052 -0.392203
- h -1.970793 2.318459 -0.459255
- h -2.439831 3.603409 0.252228
- o -5.169393 1.719718 -0.190165
- h -4.320011 2.205178 -0.272407
- h -5.730134 2.038664 -0.902914
- o -4.526410 -0.959134 0.033138
- h -4.148482 -1.058561 0.929828
- h -4.833048 -0.029798 -0.037815

pTS-.H+W6A

С	4.575628	-0.431328	0.000000
С	3.858610	-0.444013	1.201523
h	4.395128	-0.436518	2.144506
с	2.467967	-0.469187	1.211295
h	1.918860	-0.491667	2.143870

- c 1.781720 -0.480301 0.000000
- c 2.467967 -0.469187 -1.211295
- h 1.918860 -0.491667 -2.143870
- c 3.858610 -0.444013 -1.201523
- h 4.395128 -0.436518 -2.144506
- c 6.084368 -0.437072 0.000000
- h 6.469335 -1.462667 0.000000
- h 6.484917 0.062614 -0.884630
- h 6.484917 0.062614 0.884630
- s -0.011213 -0.444759 0.000000
- o -0.426295 1.006102 0.000000
- o -0.444619 -1.126800 1.251536
- o -0.444619 -1.126800 -1.251536
- o -2.912911 -0.883063 -2.246942
- h -2.004568 -1.096465 -1.913810
- h -3.085169 -1.476647 -2.984511
- o -2.244881 1.902695 -1.909713
- h -1.437605 1.679006 -1.397382
- h -2.486472 1.067932 -2.349199
- o -2.912911 -0.883063 2.246942
- h -2.004568 -1.096465 1.913810
- h -3.085169 -1.476647 2.984511
- o -4.536405 -0.374051 0.000000
- h -4.075289 -0.734404 -0.784165
- h -4.075289 -0.734404 0.784165
- o -3.924092 2.105712 0.000000
- h -3.295650 2.135277 0.820221
- h -4.289685 1.140840 0.000000
- h -3.295650 2.135277 -0.820221

- o -2.244881 1.902695 1.909713
- h -2.486472 1.067932 2.349199
- h -1.437605 1.679006 1.397382

pTS-.H+W6B

С	4.844910	-0.058942	-0.210842
С	4.148573	-0.523193	0.907278
h	4.700151	-0.893911	1.764952
С	2.755067	-0.524067	0.941961
h	2.215008	-0.890501	1.805388
С	2.049732	-0.054978	-0.159483
С	2.718282	0.409251	-1.291743
h	2.156489	0.760426	-2.148276
С	4.106747	0.403398	-1.309048
h	4.628292	0.760303	-2.191375
с	6.353695	-0.036641	-0.236438
h	6.740533	-0.385908	-1.197339
h	6.732399	0.979715	-0.083832
h	6.775281	-0.667498	0.548455
S	0.254137	-0.028279	-0.123234
0	-0.136732	1.418878	-0.123403
0	-0.158290	-0.736147	1.118521
0	-0.186103	-0.715654	-1.375195
0	-4.376802	-1.491675	1.566730
h	-3.645605	-0.959099	2.011089
h	-5.129185	-1.535939	2.162990
0	-2.442058	-1.866389	-2.001484
h	-1.574535	-1.431376	-1.755709

- h -2.294449 -2.816540 -1.975490
- o -2.348029 2.377161 0.881083
- h -1.474391 2.061290 0.505798
- h -2.200322 3.265741 1.219953
- o -4.210844 1.704616 -0.954071
- h -3.553237 2.041478 -0.295515
- h -4.000908 2.123642 -1.794578
- o -4.446697 -0.866300 -0.870530
- h -4.482494 -1.131882 0.127071
- h -4.369650 0.143170 -0.941524
- h -3.624329 -1.311418 -1.328053
- o -2.434631 -0.038274 2.504305
- h -1.585113 -0.357764 2.127360
- h -2.504890 0.873307 2.174226

pTS-.H+W6C

- c 4.276233 0.460444 0.168700
- c 3.520694 0.051900 1.270970
- h 4.025209 -0.270289 2.175622
- c 2.128194 0.051297 1.232619
- h 1.545778 -0.268551 2.086903
- c 1.483717 0.464897 0.071901
- c 2.210720 0.880244 -1.043418
- h 1.696087 1.199726 -1.941137
- c 3.598471 0.874576 -0.986042
- h 4.166356 1.198854 -1.851881
- c 5.784193 0.436266 0.206520
- h 6.172990 -0.446758 -0.311872

- h 6.208101 1.314875 -0.285839
- h 6.157114 0.407639 1.231958
- s -0.307861 0.520307 0.017341
- o -0.712626 1.947420 0.119679
- o -0.794409 -0.295725 1.179248
- o -0.710969 -0.078523 -1.301554
- o -3.525511 0.229095 -1.622518
- h -2.557964 0.136953 -1.735546
- h -3.642918 1.126654 -1.260566
- o -4.103912 -1.343858 0.329017
- h -3.945709 -0.786355 -0.518353
- h -3.305097 -2.019125 0.406950
- h -3.957659 -0.671507 1.086636
- o -3.425056 0.451202 2.050143
- h -2.458849 0.310840 2.023578
- h -3.557607 1.307426 1.602898
- o -3.339785 2.525365 0.086920
- h -3.590314 3.452889 0.035835
- h -2.352908 2.488706 0.078256
- o -2.040156 -2.756056 0.604671
- h -1.638067 -3.022157 -0.250885
- h -1.443902 -2.054138 0.934873
- o -0.841137 -2.782979 -1.846146
- h -0.637266 -1.827213 -1.794030
- h -0.084177 -3.214343 -2.251453

pTS-.H+W7A

c 4.648204 0.413359 0.604882

- c 3.869973 1.491455 0.176998
- h 4.311796 2.480393 0.118953
- c 2.532730 1.322573 -0.177165
- h 1.931664 2.161405 -0.503389
- c 1.967328 0.055156 -0.097622
- c 2.719561 -1.040695 0.326009
- h 2.264504 -2.021409 0.387879
- c 4.050598 -0.853078 0.673684
- h 4.637130 -1.703836 1.005065
- c 6.095275 0.597284 0.990138
- h 6.260146 0.331481 2.038835
- h 6.747206 -0.042286 0.387839
- h 6.416809 1.630829 0.851757
- s 0.262796 -0.195696 -0.596074
- o -0.346052 -1.112918 0.426069
- o 0.278531 -0.810174 -1.948955
- o -0.380094 1.161345 -0.595941
- o -2.736249 -2.422295 -0.493871
- h -1.892166 -2.105879 -0.120113
- h -2.650167 -2.270946 -1.450963
- o -1.461924 -0.326957 2.821416
- h -0.973526 -0.699061 2.056857
- h -0.905527 -0.457779 3.594389
- o -2.144919 1.933463 1.435369
- h -1.971583 1.246500 2.115875
- h -1.404032 1.823733 0.804175
- o -4.113563 1.361970 -0.029244
- h -3.352501 1.621531 0.640488
- h -3.660766 1.321049 -0.950959

- h -4.440508 0.421920 0.202198
- o -2.735958 1.266944 -2.184732
- h -2.707247 0.415458 -2.662001
- h -1.833769 1.369139 -1.821367
- o -2.100367 -1.346451 -3.099829
- h -1.159407 -1.224818 -2.828301
- h -2.093527 -1.725244 -3.984251
- o -4.801730 -1.056951 0.441678
- h -5.076381 -1.369098 1.308784
- h -4.052286 -1.641419 0.135436

pTS-.H+W7B

с	4.454441	-0.471129	1.201147
с	3.682169	0.648557	1.515826
h	4.113596	1.446004	2.111261
С	2.363982	0.765227	1.076407
h	1.770002	1.637287	1.316727
С	1.813427	-0.256119	0.312122
С	2.562132	-1.385731	-0.020820
h	2.125237	-2.171590	-0.624446
С	3.872329	-1.484454	0.425537
h	4.456406	-2.361620	0.166625
С	5.881972	-0.595599	1.673091
h	6.174721	0.256841	2.288224
h	6.023129	-1.503992	2.266285
h	6.572419	-0.651727	0.825960
S	0.114581	-0.140588	-0.253922
0	-0.643169	-1.233415	0.444291

- o 0.136343 -0.339738 -1.727205
- o -0.369763 1.221078 0.146397
- o -2.219650 -1.006471 2.743503
- h -1.810261 -1.248219 3.578772
- h -1.569783 -1.207204 2.042313
- o -2.719880 1.463516 1.663665
- h -2.763438 0.640965 2.195413
- h -1.810076 1.454076 1.304921
- o -3.952603 1.817250 -0.511640
- h -3.567788 1.613196 0.430760
- h -3.210319 2.333316 -0.984639
- h -4.039401 0.937699 -1.057202
- o -1.864946 2.816397 -1.672783
- h -1.774963 2.303477 -2.496651
- h -1.169317 2.450935 -1.092178
- o -1.689281 0.616429 -3.467066
- h -0.977668 0.207586 -2.917568
- h -1.432579 0.502321 -4.387266
- o -3.976282 -0.236026 -1.997615
- h -3.301445 -0.036959 -2.673671
- h -3.678872 -1.075894 -1.577058
- o -2.865236 -2.393201 -0.766006
- h -2.034411 -2.077329 -0.355291
- h -2.675755 -3.255151 -1.146223

pTS-.H+W7C

- c 4.534610 -0.427781 1.202265
- c 3.784771 0.733253 1.399583

- h 4.237942 1.585197 1.894982
- c 2.460860 0.821145 0.970933
- h 1.881752 1.721997 1.126137
- c 1.881974 -0.271678 0.337757
- c 2.606406 -1.445636 0.126518
- h 2.144108 -2.291181 -0.367668
- c 3.922911 -1.514222 0.559791
- h 4.487920 -2.426282 0.397076
- c 5.967832 -0.522165 1.663884
- h 6.292542 0.399865 2.148840
- h 6.098989 -1.342014 2.376519
- h 6.639972 -0.714509 0.822156
- s 0.187084 -0.181954 -0.242427
- o -0.541490 -1.328769 0.392294
- o 0.227259 -0.304482 -1.723511
- o -0.346166 1.146313 0.212766
- o -2.527507 -1.420931 2.265113
- h -2.371569 -1.826803 3.123206
- h -1.676570 -1.444805 1.777397
- o -2.770327 1.294522 1.609860
- h -2.896473 0.405874 1.995055
- h -1.840576 1.290590 1.302740
- o -3.882351 1.909231 -0.603264
- h -3.553990 1.639142 0.333685
- h -3.085948 2.389352 -1.028887
- h -3.998092 1.039044 -1.177325
- o -1.700853 2.806821 -1.649133
- h -1.611658 2.298863 -2.475914
- h -1.046655 2.400124 -1.046882

- o -1.622854 0.597699 -3.458217
- h -0.909798 0.192157 -2.907961
- h -1.375057 0.462131 -4.378141
- o -3.977650 -0.142723 -2.056821
- h -3.264922 -0.008333 -2.710811
- h -3.736768 -0.961920 -1.559060
- o -3.081726 -2.237270 -0.559384
- h -2.129033 -2.066464 -0.443529
- h -3.408896 -2.253398 0.350376

pTS-.H+W7D

с	4.584269	0.859585	0.088992
с	3.829774	0.981262	1.263735
h	4.335968	1.140425	2.210214
с	2.443288	0.902878	1.240385
h	1.868618	0.993297	2.153691
с	1.795652	0.698643	0.022379
с	2.517177	0.577375	-1.160208
h	1.997015	0.416798	-2.095647
с	3.907159	0.657570	-1.116588
h	4.471822	0.561456	-2.037751
с	6.090732	0.924674	0.135487
h	6.514445	1.077314	-0.858661
h	6.432823	1.738522	0.779879
h	6.509317	-0.004771	0.535777
S	0.003640	0.658531	-0.018998
0	-0.478334	2.061336	0.061004

o -0.429877 -0.143901 1.176825

- o -0.375329 -0.000958 -1.312763
- o -3.155300 0.488913 2.005555
- h -2.190548 0.396277 1.912610
- h -3.374063 1.264958 1.463063
- o -3.136313 2.549393 -0.037711
- h -3.421085 3.468461 -0.044152
- h -2.151622 2.551091 0.012943
- o -3.046925 0.617551 -2.097494
- h -3.293656 1.347955 -1.497571
- h -2.074488 0.557252 -2.018722
- o -3.674251 -1.780694 -1.465923
- h -3.542751 -0.795501 -1.725840
- h -2.754962 -2.211748 -1.600483
- h -3.840616 -1.831079 -0.432027
- o -1.249394 -2.649668 -1.568810
- h -1.086592 -2.988563 -0.669823
- h -0.762782 -1.800335 -1.588694
- o -1.133753 -2.765521 1.267928
- h -0.775382 -1.846975 1.305611
- h -0.650501 -3.280216 1.921752
- o -3.874041 -1.935079 1.040330
- h -3.070218 -2.418636 1.304278
- h -3.758916 -1.037871 1.445424

pTS-.H+W7E

С	4.555538	0.847992	0.066776
с	3.797158	0.984580	1.236229
h	4.301152	1.125893	2.186565

- c 2.408300 0.946288 1.202774
- h 1.831594 1.052040 2.113116
- c 1.762810 0.766606 -0.019284
- c 2.489878 0.632954 -1.198292
- h 1.971685 0.496038 -2.138532
- c 3.880368 0.673680 -1.145209
- h 4.447977 0.570083 -2.063891
- c 6.063162 0.862297 0.118419
- h 6.456351 -0.146629 0.284264
- h 6.490257 1.230282 -0.816708
- h 6.428120 1.493208 0.931736
- s -0.030476 0.788433 -0.079607
- o -0.485000 2.187285 0.047235
- o -0.489809 -0.049264 1.099075
- o -0.412332 0.154567 -1.383491
- o -2.978214 0.507330 2.126843
- h -2.033838 0.397501 1.866955
- h -2.983162 0.772694 3.051981
- o -3.337378 2.447900 -0.044750
- h -3.476874 1.926267 0.761923
- h -2.386940 2.658501 -0.010960
- o -3.123297 0.725794 -2.126780
- h -3.361394 1.398144 -1.437964
- h -2.147615 0.697973 -2.096138
- o -3.682497 -1.639125 -1.582879
- h -3.538302 -0.611505 -1.814198
- h -2.769647 -2.070242 -1.740747
- h -3.838117 -1.729827 -0.579825
- o -1.248107 -2.491525 -1.750588

- h -1.060263 -2.864454 -0.870791
- h -0.777908 -1.632790 -1.740821
- o -1.040934 -2.700168 1.080356
- h -0.713850 -1.770009 1.140698
- h -0.505378 -3.226894 1.681608
- o -3.793024 -1.930476 0.989038
- h -2.957548 -2.395162 1.180346
- h -3.712444 -1.086260 1.475519

pTS-.H+W8A

с	4.905338	-0.136598	0.172095
с	4.240151	0.805126	0.970441
h	4.809961	1.418729	1.660586
С	2.864397	0.971083	0.892640
h	2.362388	1.707699	1.507575
С	2.133998	0.182872	0.002597
С	2.765697	-0.757118	-0.802892
h	2.183486	-1.352629	-1.493969
С	4.148444	-0.909313	-0.711021
h	4.643108	-1.640789	-1.340993
С	6.402407	-0.297029	0.267279
h	6.760286	-1.097346	-0.382464
h	6.915261	0.625372	-0.022096
h	6.709441	-0.530418	1.291002
S	0.351951	0.377861	-0.069838
0	0.084388	1.834151	-0.192048
0	-0.199398	-0.184802	1.208743

o -0.104504 -0.405381 -1.265423

- o -1.617364 0.715448 -3.308025
- h -1.149118 1.252086 -3.953263
- h -0.948677 0.385503 -2.674940
- o -3.749398 1.271957 -1.685698
- h -3.059803 1.164084 -2.378200
- h -3.385991 1.931400 -1.061952
- o -2.368403 2.940523 0.121211
- h -1.441201 2.641443 -0.033997
- h -2.360756 3.902183 0.086934
- o -2.444486 1.194276 2.425046
- h -2.579142 1.936159 1.812046
- h -1.578098 0.832663 2.167450
- o -1.503020 -2.523696 1.703058
- h -0.919095 -1.738267 1.592674
- h -1.052420 -3.124602 2.304409
- o -3.947763 -0.995312 1.853473
- h -3.266847 -1.670861 2.020153
- h -3.523417 -0.150773 2.146995
- o -4.194921 -1.005174 -0.672892
- h -4.174597 -0.959809 0.363628
- h -3.391996 -1.584260 -0.926982
- h -4.052413 -0.057935 -1.076266
- o -2.068799 -2.411348 -1.074240
- h -1.865144 -2.766401 -0.190295
- h -1.333933 -1.790754 -1.252543

pTS-.H+W8B

c 4.304262 -1.050479 1.662325

- c 3.607590 -2.020640 0.930545
- h 4.130556 -2.905015 0.581799
- c 2.256279 -1.872920 0.642638
- h 1.726608 -2.626427 0.073279
- c 1.586742 -0.736295 1.092671
- c 2.250963 0.241936 1.825515
- h 1.715045 1.118667 2.165396
- c 3.605270 0.076994 2.103458
- h 4.125080 0.839662 2.673499
- c 5.778375 -1.208206 1.941646
- h 6.043229 -2.255668 2.102992
- h 6.375031 -0.846404 1.097247
- h 6.080051 -0.640407 2.823993
- s -0.170825 -0.571874 0.775545
- o -0.885701 -1.441777 1.745278
- o -0.389745 -1.037263 -0.638168
- o -0.502270 0.877199 0.961745
- o -0.478480 0.025229 -3.117997
- h -0.346459 -0.325464 -2.202915
- h -0.175016 -0.669788 -3.711310
- o 0.476882 2.642582 -3.073046
- h 0.401668 3.131059 -3.897530
- h 0.216650 1.718759 -3.259901
- o -0.904789 3.058456 -0.733862
- h -0.371813 3.049290 -1.557743
- h -0.536838 2.362083 -0.155089
- o -3.319258 2.326120 -0.908984
- h -2.350083 2.701051 -0.901448
- h -3.444388 1.913197 0.018137

- h -3.349034 1.562267 -1.610086
- o -3.289995 1.301698 1.446504
- h -3.632951 0.401388 1.603812
- h -2.322032 1.210310 1.538900
- o -3.595957 -1.500702 1.710576
- h -2.625334 -1.597003 1.853845
- h -4.034506 -2.025369 2.387368
- o -3.134671 -1.865818 -1.125484
- h -2.179248 -1.805344 -0.947845
- h -3.537690 -1.887312 -0.241573
- o -3.295124 0.418548 -2.587751
- h -3.397941 -0.435790 -2.099170
- h -2.410133 0.349471 -2.988811

pTS-.H+W9A

С	4.900088	0.206307	1.265017
с	4.010458	1.281182	1.317369
h	4.385142	2.276465	1.530485
С	2.644258	1.100891	1.105364
h	1.957633	1.935741	1.156757
с	2.165309	-0.175432	0.835106
с	3.029708	-1.269727	0.780436
h	2.642950	-2.261480	0.581210
с	4.386383	-1.070486	0.994023
h	5.060065	-1.920249	0.954354
с	6.377221	0.400599	1.501708
h	6.963833	0.036512	0.653112
h	6.622117	1.452969	1.653817

- h 6.708909 -0.152716 2.385650
- s 0.413984 -0.432461 0.542414
- o -0.019848 -1.540852 1.429639
- o 0.281425 -0.818583 -0.911312
- o -0.259948 0.868984 0.832796
- o -4.397737 0.729738 0.210810
- h -4.310330 -0.272154 0.080570
- h -3.794742 0.931081 1.000888
- o -3.950649 -1.828142 0.067812
- h -3.286657 -2.051999 -0.613170
- h -3.531768 -2.035496 0.924268
- o -2.476239 -1.885633 2.499384
- h -2.490731 -2.497835 3.241499
- h -1.545527 -1.847447 2.176903
- o -2.738222 0.958033 2.203090
- h -2.746352 0.067605 2.596185
- h -1.836526 1.045988 1.837671
- o -3.524553 1.884017 -1.773794
- h -2.669073 2.330496 -1.552373
- h -3.282931 1.242061 -2.485984
- h -4.029151 1.250470 -0.675974
- o -1.002462 2.754355 -1.212661
- h -0.493262 2.392783 -1.956798
- h -0.706512 2.223051 -0.452471
- o 0.235866 0.895682 -3.052164
- h 0.400644 0.270998 -2.311568
- h 1.031036 0.904034 -3.593518
- o -2.494965 0.079793 -3.554054
- h -2.422890 -0.789396 -3.124918

- h -1.573647 0.381772 -3.631479
- o -1.898000 -2.190977 -1.844326
- h -1.080947 -1.788204 -1.467495
- h -1.653362 -3.072336 -2.144278

pTS-.H+W9B

С	4.914966	0.190526	1.244988
С	3.975907	0.852726	2.046740
h	4.318248	1.509635	2.839445
С	2.611513	0.689451	1.842549
h	1.894584	1.214619	2.461245
с	2.174004	-0.148272	0.817488
с	3.082654	-0.816527	0.003514
h	2.726207	-1.452545	-0.796129
с	4.446420	-0.639268	0.222578
h	5.155718	-1.152350	-0.417902
с	6.393241	0.356830	1.495080
h	6.724981	-0.297869	2.308120
h	6.978827	0.104388	0.609161
h	6.635563	1.382298	1.783667
S	0.413357	-0.400419	0.585893
0	-0.046628	-1.336775	1.642882
0	0.251481	-0.959294	-0.794244
0	-0.227551	0.955182	0.726467
0	-4.338753	0.727548	0.024761
h	-4.267270	-0.291117	-0.002218
h	-3.748365	1.000882	0.806878
0	-3.960708	-1.817327	0.146048

- h -3.298089 -2.129010 -0.503219
- h -3.555984 -1.947877 1.024770
- o -2.546135 -1.643826 2.598209
- h -2.587523 -2.193938 3.386585
- h -1.602650 -1.626008 2.309981
- o -2.728801 1.155074 2.009891
- h -2.767810 0.316055 2.502358
- h -1.817314 1.180855 1.654172
- o -3.357705 1.668884 -2.048680
- h -2.522264 2.136785 -1.851347
- h -3.112175 0.958252 -2.701707
- h -3.938762 1.139922 -0.877082
- o -0.794787 2.635723 -1.362279
- h -0.539629 3.544299 -1.174322
- h -0.524216 2.097981 -0.583810
- o 0.189015 0.600724 -3.240997
- h 0.395499 0.041311 -2.473552
- h 0.013478 1.467458 -2.843051
- o -2.440658 -0.287671 -3.650516
- h -2.406954 -1.116292 -3.145980
- h -1.501190 -0.015240 -3.719540
- o -1.909874 -2.398937 -1.676481
- h -1.101802 -1.959089 -1.325188
- h -1.652068 -3.297757 -1.905227

pTS-.H+W9C

- c 5.008559 0.058962 1.322817
- c 4.215140 1.208643 1.263279

- h 4.673197 2.182636 1.398928
- c 2.843316 1.128995 1.038459
- h 2.231821 2.021466 1.007874
- c 2.257046 -0.121034 0.867221
- c 3.022030 -1.284663 0.923692
- h 2.551025 -2.252331 0.804160
- c 4.389605 -1.185430 1.148954
- h 4.986460 -2.090267 1.194558
- c 6.488201 0.152271 1.601428
- h 6.685464 0.083203 2.676674
- h 7.037318 -0.657477 1.116044
- h 6.900686 1.101438 1.253274
- s 0.498423 -0.247309 0.534765
- o -0.009971 -1.388159 1.336542
- o 0.358889 -0.516191 -0.944436
- o -0.099835 1.069581 0.912242
- o -4.462538 0.250391 -1.479996
- h -3.903754 0.165541 -2.292138
- h -4.338539 -0.611719 -1.005650
- o -3.777693 -2.089611 -0.303461
- h -2.994273 -2.342544 -0.823049
- h -3.472949 -2.050242 0.619990
- o -2.545106 -1.702264 2.255378
- h -2.605900 -2.279411 3.022979
- h -1.596639 -1.677668 1.991047
- o -2.690018 1.132332 2.135081
- h -2.845646 0.190170 2.331337
- h -1.738389 1.173807 1.926146
- o -3.708104 2.099031 -0.065250

- h -3.431268 1.723482 0.832423
- h -2.857802 2.481172 -0.455589
- h -4.061271 1.273514 -0.708616
- o -1.346715 2.883074 -0.865349
- h -1.046810 2.556588 -1.732559
- h -0.792540 2.387770 -0.232156
- o -0.264558 1.265900 -2.926288
- h 0.089576 0.643349 -2.251036
- h 0.455655 1.457147 -3.535035
- o -2.676319 -0.216854 -3.488739
- h -2.305933 -1.063940 -3.186886
- h -1.912580 0.384883 -3.488420
- o -1.441844 -2.340232 -1.965203
- h -0.742785 -1.789492 -1.547193
- h -1.022633 -3.168510 -2.218247

pTS-.H+W9D

- С 5.009419 0.026256 1.302984 4.089843 0.554553 2.219982 С h 4.451115 1.023230 3.129428 2.722849 0.489141 1.987304 С h 2.023011 0.903856 2.702259 2.260335 -0.115947 0.818415 С 3.148104 -0.650180 -0.108003 С 2.771262 -1.112506 -1.010778 h 4.517028 -0.574152 0.142088 С h 5.210519 -0.990457 -0.580714
- c 6.492498 0.128223 1.559389

- h 7.054957 -0.563636 0.929906
- h 6.855624 1.139461 1.347441
- h 6.731026 -0.090114 2.603310
- s 0.491025 -0.225490 0.541007
- o -0.058516 -1.153021 1.563605
- o 0.319504 -0.743936 -0.857080
- o -0.048263 1.166258 0.697460
- o -4.339388 -0.009074 -1.620649
- h -3.701113 -0.228379 -2.406168
- h -4.201254 -0.761589 -0.945512
- o -3.775020 -1.990830 -0.074040
- h -3.004565 -2.390981 -0.520834
- h -3.486845 -1.812375 0.843536
- o -2.623309 -1.330302 2.394491
- h -2.714706 -1.832928 3.209998
- h -1.663699 -1.330586 2.164804
- o -2.626850 1.531010 1.951027
- h -2.836436 0.642727 2.283464
- h -1.688284 1.469783 1.699318
- o -3.630703 2.139855 -0.507734
- h -3.399395 1.964511 0.437736
- h -2.819067 2.532607 -0.881043
- h -4.059218 0.894859 -1.180366
- o -0.977342 2.877013 -1.220439
- h -0.611692 3.764341 -1.148260
- h -0.545374 2.335245 -0.521212
- o -0.437261 0.819326 -3.149088
- h -0.008339 0.311137 -2.437894
- h -0.613780 1.681041 -2.735944

- o -2.678525 -0.699807 -3.417011
- h -2.310207 -1.528797 -3.064386
- h -1.899993 -0.090389 -3.457640
- o -1.500530 -2.632200 -1.632413
- h -0.788635 -2.037970 -1.298319
- h -1.099812 -3.498789 -1.753423

pTS-.H+W9E

с	4.889789	-0.437138	0.351762
с	4.188607	0.240215	1.357399
h	4.727349	0.632187	2.213457
с	2.812005	0.416019	1.283080
h	2.280440	0.935195	2.071065
с	2.121677	-0.095175	0.185344
с	2.790034	-0.776694	-0.827471
h	2.233685	-1.176534	-1.665246
с	4.169122	-0.942974	-0.734827
h	4.690872	-1.479977	-1.519939
с	6.387742	-0.598673	0.426693
h	6.707220	-1.552582	0.001243
h	6.891927	0.194874	-0.135342
h	6.744011	-0.547960	1.457387
S	0.347690	0.148074	0.077834
0	0.127204	1.617924	-0.123720
0	-0.232260	-0.303278	1.382524
0	-0.121533	-0.663375	-1.083554
0	0.021408	3.002585	-2.547087
h	0 001000	3.374335	2 020002

- 0.282030 2.582064 -1.706657 h
- -1.583213 0.749166 -3.181093 0
- -1.092378 1.589892 -3.226020 h
- -1.020371 0.185914 -2.624142 h
- -3.765915 1.001736 -1.595084 0
- h -3.050530 0.928598 -2.278218
- -3.429461 1.662703 -0.959459 h
- -2.396549 2.720196 0.211243 0
- h -1.475276 2.412356 0.066056
- -2.394836 3.666529 0.033311 h
- -2.511674 1.077840 2.564139 0
- -2.647775 1.815699 1.947006 h
- h -1.642572 0.721921 2.311586
- -4.013130 -1.122931 1.995302 0
- -3.338690 -1.793935 2.198489 h
- -3.595112 -0.272641 2.277878 h
- 0 -4.227282 -1.227654 -0.545686
- -4.225597 -1.142790 0.482834 h
- h -3.428277 -1.821186 -0.759408
- -4.065568 -0.285538 -0.985244 h
- -2.090827 -2.664678 -0.845110 0
- -1.902108 -2.980290 0.056396 h
- h -1.350947 -2.056648 -1.039150
- -1.551264 -2.645554 1.950806 ο
- -0.968443 -1.863876 1.827454 h
- -1.112247 -3.223939 2.582174 h
- 46

pTS-.H+W9X

- c 4.370311 -0.533749 0.962283
- c 3.603593 0.182144 1.892004
- h 4.101994 0.790873 2.639097
- c 2.215411 0.130092 1.874988
- h 1.633659 0.693646 2.593771
- c 1.573248 -0.648143 0.912265
- c 2.307002 -1.370600 -0.024020
- h 1.791786 -1.973736 -0.760891
- c 3.700916 -1.303997 0.005428
- h 4.273453 -1.872517 -0.720150
- c 5.877245 -0.489786 1.014543
- h 6.252897 -1.088389 1.850792
- h 6.321696 -0.885470 0.099573
- h 6.240890 0.530917 1.157493
- s -0.222999 -0.743117 0.910780
- o -0.626969 -1.598744 2.054506
- o -0.599500 -1.337457 -0.409984
- o -0.707070 0.667419 1.079137
- o -3.285322 -2.344255 -0.476703
- h -2.314157 -2.253086 -0.461893
- h -3.535591 -2.351791 0.465679
- o -3.311915 -2.041300 2.355364
- h -3.586374 -2.645287 3.052161
- h -2.329846 -1.988318 2.392292
- o -3.457654 0.838684 2.015564
- h -3.667307 -0.074279 2.272387
- h -2.490974 0.833474 1.904123
- o -4.032148 1.709859 -0.510418
- h -3.991042 1.413730 0.430945

- h -3.240684 2.270046 -0.613638
- o -1.380794 2.811960 -0.472899
- h -1.189696 3.614382 0.023586
- h -1.046230 2.064218 0.072632
- o -1.446404 -0.272150 -2.863335
- h -0.983113 -0.644046 -2.088672
- h -1.003572 0.597460 -3.031020
- o -3.813175 -0.321565 -2.013695
- h -3.946323 0.518437 -1.417299
- h -2.867710 -0.246925 -2.455696
- h -3.749246 -1.129330 -1.401646
- o -0.160529 2.105014 -2.954458
- h -0.535624 2.545314 -2.171846
- h 0.780940 1.971878 -2.749845
- o 2.537590 1.548314 -2.146503
- h 3.282814 1.508146 -2.753950
- h 2.656523 0.812030 -1.532834

pTS-.H+W10A

- c 5.222075 -0.281363 1.053335
- c 4.672279 0.350542 -0.071538
- h 5.329899 0.814580 -0.799257
- c 3.300056 0.391392 -0.275323
- h 2.890280 0.877326 -1.152221
- c 2.455830 -0.209403 0.659385
- c 2.971600 -0.844496 1.782591
- h 2.300195 -1.312666 2.490552
- c 4.352736 -0.875665 1.970271

- h 4.756832 -1.374043 2.844877
- c 6.716547 -0.312913 1.256557
- h 6.985182 -0.861790 2.160577
- h 7.219002 -0.790306 0.410104
- h 7.122325 0.699525 1.344373
- s 0.679494 -0.134607 0.411882
- o 0.075721 -1.029606 1.430759
- o 0.426955 -0.583853 -0.996067
- o 0.283718 1.303581 0.590112
- o -4.268157 -0.388192 -1.903004
- h -3.589711 -0.466182 -2.672259
- h -4.052898 -1.163836 -1.279454
- h -4.110792 0.514431 -1.376742
- o -3.866801 1.715527 -0.620921
- h -3.847422 1.562120 0.360278
- h -3.020277 2.155212 -0.834091
- o -1.294550 2.787560 -1.106149
- h -1.166077 3.723758 -0.921514
- h -0.710100 2.302536 -0.481460
- o -0.431407 1.033286 -3.214336
- h 0.027340 0.522089 -2.524658
- h -0.743478 1.821225 -2.739406
- o -2.458047 -0.744437 -3.676942
- h -1.998677 -1.537508 -3.350763
- h -1.765995 -0.040032 -3.652629
- o -1.102812 -2.633389 -1.946598
- h -0.481768 -1.978452 -1.552518
- h -0.587456 -3.426272 -2.125151
- o -3.483639 -2.367957 -0.438143

- h -2.658518 -2.645695 -0.879487
- h -3.231556 -2.139920 0.478862
- o -2.526459 -1.494268 2.052185
- h -2.560774 -2.047622 2.839020
- h -1.572058 -1.328806 1.869310
- o -3.599710 1.200801 2.013696
- h -3.334559 0.273025 2.131596
- h -2.844495 1.721182 2.349121
- o -1.189214 2.516187 2.613154
- h -0.552770 2.058871 2.030131
- h -0.737889 2.668527 3.447899

pTS-.H+W10B

С	5.008825	-0.002354	1.679372
с	4.111967	1.017452	2.015145
h	4.474781	1.900765	2.530184
С	2.759176	0.921048	1.703440
h	2.069344	1.713245	1.963824
С	2.294146	-0.213269	1.043937
С	3.163284	-1.246820	0.700717
h	2.790612	-2.125555	0.189233
С	4.511936	-1.132824	1.019847
h	5.189338	-1.937426	0.754200
С	6.477575	0.126069	1.998853
h	6.962457	-0.850825	2.047070
h	6.993165	0.712666	1.230851
h	6.633796	0.631889	2.954367
S	0.542550	-0.384170	0.688904

- o -0.021858 -1.323568 1.689741
- o 0.456040 -0.937402 -0.705284
- o -0.043037 0.991411 0.794152
- o -1.652291 -2.417456 -1.647763
- h -0.871847 -1.936576 -1.286921
- h -1.321776 -3.260659 -1.973964
- o -3.086345 -0.453804 -3.182113
- h -2.546764 -1.199493 -2.863191
- h -2.438144 0.257376 -3.429307
- o -4.589616 0.048789 -1.227578
- h -4.009275 -0.099120 -2.085165
- h -4.379885 -0.742504 -0.623359
- h -4.274787 0.915383 -0.751434
- o -3.829601 -2.016143 0.129513
- h -3.073707 -2.337366 -0.397557
- h -3.491807 -1.896851 1.039295
- o -2.566091 -1.505353 2.585520
- h -2.645319 -2.044505 3.378654
- h -1.614571 -1.514205 2.328707
- o -2.518308 1.371039 2.271975
- h -2.725514 0.471801 2.575546
- h -1.607784 1.299599 1.936697
- o -3.748192 2.127369 -0.046603
- h -3.432654 1.901199 0.861842
- h -2.955636 2.487693 -0.486445
- o -1.142104 2.722561 -1.039295
- h -0.738010 3.587767 -0.917610
- h -0.660438 2.107011 -0.442559
- o -1.215306 1.434507 -3.621177

- h -0.345548 0.992640 -3.654648
- h -1.178196 1.984962 -2.820461
- o 1.154276 -0.022520 -3.239276
- h 2.082865 0.195873 -3.355032
- h 1.048280 -0.319347 -2.314844

pTS-.H+W10C

с	5.210328	-0.004471	0.105569
С	4.485087	-0.093243	1.295515
h	5.013478	-0.156685	2.240597
С	3.091227	-0.106257	1.293973
h	2.534548	-0.185164	2.218407
с	2.415197	-0.030148	0.082387
с	3.113116	0.054747	-1.122973
h	2.578294	0.100379	-2.063644
с	4.500778	0.066939	-1.101919
h	5.044530	0.129164	-2.038820
С	6.718607	0.015287	0.107418
h	7.124074	-0.786500	-0.516452
h	7.098519	0.960463	-0.292472
h	7.118394	-0.106824	1.115304
S	0.620630	-0.007074	0.053092
0	0.207653	1.381924	-0.320650
0	0.171641	-0.379511	1.430377
0	0.201952	-1.005273	-0.980204
0	-1.292436	2.270623	-2.460138
h	-0.657274	2.011927	-1.757601
h	-0.936095	3.059680	-2.880381

- o -1.500573 -0.463567 -3.254018
- h -0.797662 -0.646578 -2.607151
- h -1.534763 0.506734 -3.281152
- o -1.387622 -3.249744 -0.757320
- h -0.718642 -2.537960 -0.852875
- h -1.024597 -4.031055 -1.186335
- o -3.514314 -1.652487 -1.829347
- h -2.969090 -2.361897 -1.444017
- h -2.901339 -1.205218 -2.460626
- o -4.442995 0.005642 -0.131875
- h -4.088120 0.947252 -0.362395
- h -4.084030 -0.678316 -0.816288
- h -4.135222 -0.258301 0.816952
- o -3.529397 2.321838 -0.669720
- h -2.940529 2.355504 -1.444992
- h -2.965348 2.667130 0.063449
- o -1.641210 3.046371 1.174060
- h -1.711087 2.582389 2.024397
- h -0.882998 2.615765 0.743813
- o -1.508875 0.991313 3.152977
- h -0.825708 0.531356 2.619204
- h -1.200052 0.986262 4.064453
- o -3.640336 -0.664893 2.191918
- h -3.100445 0.010319 2.640536
- h -3.043364 -1.450470 2.154671
- o -1.669621 -2.553765 1.997144
- h -0.921779 -1.943839 1.879288
- h -1.682531 -3.059769 1.167865

pTS-.H+W10D

С	5.203390	0.037077	0.101058
с	4.489326	1.212940	-0.155819
h	5.027813	2.134145	-0.351990
с	3.098392	1.226687	-0.159006
h	2.551256	2.142102	-0.344139
С	2.408929	0.044331	0.097401
С	3.092004	-1.141011	0.357990
h	2.545006	-2.051007	0.570775
с	4.482919	-1.135156	0.356549
h	5.016867	-2.056764	0.562028
с	6.711566	0.044044	0.131429
h	7.078913	0.436773	1.085487
h	7.118370	-0.961595	0.010268
h	7.122153	0.675937	-0.659710
S	0.614996	0.037782	0.046001
0	0.175667	1.444067	0.240707
0	0.164642	-0.888149	1.131091
0	0.225431	-0.492443	-1.310991
0	-1.645741	3.000759	-1.405565
h	-0.921216	2.620616	-0.880811
h	-1.641390	2.460968	-2.212166
0	-1.265483	0.814536	-3.221619
h	-0.612379	0.403209	-2.613451
h	-0.863527	0.834414	-4.095622
0	-1.337693	-2.675140	-1.909382
h	-0.666226	-1.983665	-1.717300
h	-0.959096	-3.258597	-2.574232

- o -3.438424 -0.768435 -2.262708
- h -2.889032 -1.568813 -2.178243
- h -2.856143 -0.137948 -2.724458
- o -4.434376 0.122712 -0.005102
- h -4.095721 1.126763 0.165756
- h -4.071029 -0.223229 -0.880157
- h -4.124595 -0.493307 0.756218
- o -3.633666 2.433087 0.394499
- h -3.046754 2.736649 -0.337954
- h -3.049699 2.471349 1.189874
- o -1.676470 2.370089 2.306826
- h -1.692131 1.596689 2.894223
- h -0.926561 2.189895 1.716051
- o -1.418959 -0.289802 3.325618
- h -0.752668 -0.506525 2.639134
- h -1.073733 -0.618166 4.161760
- o -3.617617 -1.419159 1.869391
- h -3.048673 -0.981748 2.527736
- h -3.046314 -2.140335 1.515463
- o -1.676990 -3.102724 0.899476
- h -0.942853 -2.473914 1.007250
- h -1.702481 -3.249987 -0.059635

pTS-.H+W10E

- c 4.791209 0.602407 -0.709590 c 4.178772 1.583312 0.079728
- h 4.792449 2.309895 0.601597
- c 2.794509 1.650358 0.202422

- h 2.326571 2.420843 0.802138
- c 2.008271 0.719079 -0.472668
- c 2.588936 -0.268130 -1.265683
- h 1.963964 -0.976201 -1.794904
- c 3.975100 -0.319112 -1.376153
- h 4.428622 -1.085472 -1.995675
- c 6.290872 0.562158 -0.867323
- h 6.647043 -0.455613 -1.039664
- h 6.605848 1.168729 -1.723230
- h 6.796244 0.956689 0.016578
- s 0.224912 0.759739 -0.268176
- o -0.147162 2.179765 -0.056349
- o -0.081508 -0.087496 0.944398
- o -0.357094 0.164240 -1.509593
- o -2.072727 1.497761 -3.253848
- h -1.682600 2.096076 -3.896521
- h -1.339377 1.119477 -2.731216
- o -4.117774 1.651433 -1.434418
- h -3.480244 1.712080 -2.178274
- h -3.731986 2.196842 -0.721974
- o -2.658642 3.026787 0.569059
- h -1.728360 2.842107 0.305810
- h -2.718743 3.972350 0.737658
- o -2.473580 0.854985 2.467334
- h -2.672770 1.690388 2.012407
- h -1.589902 0.613001 2.145313
- o -3.940469 -1.279909 1.575871
- h -3.202434 -1.921122 1.530662
- h -3.536809 -0.488543 2.005001

- o -4.391603 -0.817169 -0.863465
- h -4.286564 -0.972205 0.164977
- h -3.585874 -1.286987 -1.283403
- h -4.324530 0.190013 -1.078576
- o -2.242719 -1.993610 -1.672226
- h -1.965878 -2.487570 -0.874192
- h -1.550882 -1.314990 -1.774503
- o -1.521163 -2.562674 0.949313
- h -0.994709 -1.741297 0.950939
- h -0.978705 -3.206215 1.454029
- o 0.342393 -3.937573 2.463185
- h 0.933153 -3.167736 2.600631
- h 0.163752 -4.299104 3.336181
- o 1.716595 -1.537484 2.583630
- h 1.179726 -0.925883 2.044751
- h 2.625867 -1.408417 2.298010

pTS-.H+W11A

С	4.119642	2.419013	2.173988
с	3.772658	1.797551	0.966894
h	4.481016	1.789644	0.145144
с	2.536723	1.184912	0.802301
h	2.291726	0.705263	-0.137285
с	1.627306	1.189529	1.859815
с	1.946179	1.794431	3.070477
h	1.229629	1.779316	3.881151
С	3.190678	2.403228	3.217678
h	3.441366	2.870886	4.163826

- c 5.454130 3.105001 2.329460
- h 5.424406 4.113458 1.902997
- h 5.733845 3.200276 3.380286
- h 6.246130 2.556981 1.813749
- s 0.019081 0.414498 1.656098
- o -0.660402 0.496361 2.970465
- o 0.260151 -0.998747 1.216920
- o -0.707995 1.195264 0.594519
- o -2.478440 0.313224 -3.287180
- h -2.002709 -0.329783 -3.847514
- h -1.783641 0.881426 -2.905189
- o -3.969410 -0.699804 -1.404229
- h -3.429784 -0.310044 -2.165151
- h -3.589698 -1.703045 -1.156052
- h -3.855622 -0.095207 -0.599121
- o -3.112979 -2.920877 -0.832390
- h -2.354317 -3.177068 -1.414513
- h -2.769053 -2.921921 0.096347
- o -2.075937 -2.704270 1.661682
- h -2.594387 -2.148292 2.266896
- h -1.219548 -2.248321 1.588708
- o -3.197329 -0.477797 3.153119
- h -3.652578 -0.414917 3.998241
- h -2.303100 -0.087790 3.275155
- o -3.539975 0.837293 0.645762
- h -3.671305 0.430720 1.521921
- h -2.598771 1.097795 0.638684
- o -0.309582 1.689256 -2.022510
- h -0.175646 2.636596 -2.129991

- h -0.406344 1.533635 -1.052004
- o 1.294696 -0.363875 -3.189688
- h 1.441270 -0.969726 -2.439705
- h 0.876028 0.426956 -2.803557
- o 1.296732 -2.174468 -0.993543
- h 2.032701 -2.723363 -0.703682
- h 0.944709 -1.739916 -0.181438
- o -0.713489 -1.504968 -4.543723
- h -0.572352 -1.666510 -5.481596
- h 0.114211 -1.089856 -4.184194
- o -0.942138 -3.397038 -2.384849
- h -0.150700 -3.090329 -1.909140
- h -0.937783 -2.904580 -3.224037

pTS-.H+W11B

с	4.995129	1.722442	-1.712467
с	3.871331	2.541081	-1.557183
h	3.929048	3.587848	-1.836043
с	2.678444	2.040172	-1.044841
h	1.819916	2.686847	-0.915384
с	2.606419	0.697251	-0.684087
с	3.709267	-0.141502	-0.826371
h	3.641495	-1.179806	-0.527575
с	4.893062	0.377051	-1.339674
h	5.753575	-0.274653	-1.448551
с	6.294423	2.282044	-2.236126
h	6.934517	2.613303	-1.411297
h	6.852335	1.532325	-2.801379

- h 6.125070 3.143298 -2.885522
- s 1.059959 0.016818 -0.076973
- o 1.411996 -1.090768 0.839940
- o 0.307894 -0.476945 -1.289140
- o 0.333787 1.149045 0.578867
- o -1.565538 0.867249 -2.768482
- h -0.800931 0.432834 -2.327394
- h -1.319208 1.000150 -3.689119
- o -1.719898 2.777309 -0.620709
- h -0.940155 2.335676 -0.241749
- h -1.825762 2.344735 -1.483591
- o -3.481923 1.859930 1.286894
- h -2.820095 1.753146 1.993424
- h -2.979541 2.291543 0.557746
- o -1.139691 1.392920 2.890868
- h -0.520931 1.272792 2.137722
- h -0.743854 2.057707 3.463607
- o -1.910313 -1.191301 3.856707
- h -1.146685 -1.739652 3.605700
- h -1.643024 -0.279533 3.647292
- o 0.184201 -2.745894 2.636189
- h 0.644921 -2.143980 2.011401
- h 0.873364 -3.233729 3.097794
- o -1.982920 -3.807403 1.064978
- h -1.203778 -3.622693 1.617914
- h -1.723935 -3.535658 0.167661
- o -3.832051 -2.120805 2.159287
- h -3.277852 -2.817387 1.729880
- h -3.244891 -1.753402 2.865277

- o -4.511708 -0.376173 0.553883
- h -4.148399 -0.602719 -0.359196
- h -4.125138 0.525191 0.848247
- h -4.228124 -1.145075 1.259207
- o -3.494369 -0.986974 -1.783156
- h -2.999238 -0.272589 -2.224525
- h -2.855417 -1.721922 -1.735512
- o -1.248474 -2.723162 -1.493455
- h -0.897757 -3.315915 -2.165842
- h -0.588551 -2.001065 -1.385142

pTS-.H+W11C

С	5.177665	-0.050747	1.715967
С	4.464403	0.973944	1.082397
h	4.982949	1.876629	0.777298
с	3.099690	0.859444	0.841832
h	2.558561	1.666055	0.362581
с	2.434327	-0.299228	1.236507
С	3.117433	-1.334518	1.867569
h	2.580705	-2.222027	2.176482
С	4.482885	-1.201857	2.101655
h	5.015563	-2.007544	2.595806
С	6.650950	0.095016	2.005208
h	6.811378	0.466057	3.023164
h	7.169688	-0.862612	1.921862
h	7.124845	0.800670	1.320096
S	0.680615	-0.474596	0.892326
0	0.198238	-1.609840	1.717810

- o 0.560356 -0.774090 -0.579063
- o 0.040808 0.830000 1.243925
- o -3.395364 0.785083 -3.203130
- h -2.862469 0.037799 -3.562132
- h -2.742296 1.519628 -3.111931
- o -4.453952 0.248103 -1.016538
- h -4.017124 0.471312 -1.960628
- h -4.117078 -0.649060 -0.686469
- h -4.187087 0.968144 -0.333341
- o -3.529109 -2.031230 -0.179400
- h -2.755692 -2.344899 -0.686790
- h -3.261474 -2.084236 0.757759
- o -2.395904 -2.050999 2.420465
- h -2.500670 -2.749460 3.074005
- h -1.433991 -1.986276 2.222771
- o -2.457651 0.809991 2.799546
- h -2.625951 -0.144404 2.867785
- h -1.542685 0.860456 2.476527
- o -3.720840 2.011453 0.683284
- h -3.405135 1.595915 1.520348
- h -2.946993 2.513105 0.370183
- o -1.111502 2.982537 -0.014665
- h -0.764801 3.779721 0.398722
- h -0.625899 2.228356 0.386190
- o -1.319484 2.543265 -2.834386
- h -0.549371 1.970739 -2.990407
- h -1.236009 2.822287 -1.906405
- o 0.686722 0.478047 -3.016372
- h 1.582088 0.503240 -3.367469

- h 0.748887 0.096855 -2.113766
- o -1.564408 -1.132499 -3.954762
- h -1.495400 -1.818898 -3.270246
- h -0.741318 -0.624706 -3.860154
- o -1.148348 -2.685068 -1.554443
- h -0.484374 -2.083314 -1.147083
- h -0.776377 -3.571742 -1.510487

pTS-.H+W11X

С	4.777166	-0.053042	1.633005
С	4.000886	1.092721	1.428322
h	4.473264	2.069444	1.442272
с	2.622021	1.005902	1.227310
h	2.022912	1.895619	1.084057
с	2.013582	-0.244302	1.222680
с	2.764102	-1.402590	1.424613
h	2.280065	-2.371366	1.426134
с	4.134593	-1.299331	1.625914
h	4.716551	-2.201087	1.784936
с	6.260221	0.045349	1.891402
h	6.811715	-0.727980	1.350837
h	6.655196	1.018447	1.594071
h	6.476151	-0.087378	2.956588
S	0.244949	-0.390518	0.925534
0	-0.309190	-1.201169	2.038080
0	0.112902	-1.082307	-0.397925
0	-0.287686	1.008144	0.893227
0	-1.963319	-2.606748	-1.288141

- h -1.189206 -2.109624 -0.934980
- h -1.635070 -3.474941 -1.542942
- o -3.241997 -0.694142 -3.011124
- h -2.748614 -1.439333 -2.624189
- h -2.554348 -0.029559 -3.290866
- o -4.796216 -0.003484 -1.167196
- h -4.179558 -0.233094 -1.986942
- h -4.634420 -0.744464 -0.491688
- h -4.488894 0.892707 -0.746958
- o -4.152678 -1.970142 0.393802
- h -3.408767 -2.373649 -0.092148
- h -3.815906 -1.779564 1.291100
- o -2.902337 -1.283666 2.827557
- h -3.026942 -1.746012 3.662256
- h -1.941125 -1.330398 2.617072
- o -2.798621 1.559543 2.287287
- h -3.021602 0.686733 2.650897
- h -1.879402 1.459050 1.986672
- o -3.974471 2.152854 -0.112959
- h -3.682190 1.991837 0.816088
- h -3.167601 2.472521 -0.558584
- o -1.349192 2.625805 -1.085337
- h -0.912702 3.482151 -1.033480
- h -0.887932 2.040802 -0.445322
- o -1.319826 1.096078 -3.549432
- h -0.433811 0.678650 -3.495091
- h -1.327379 1.731832 -2.814997
- o 1.064549 -0.190702 -2.942002
- h 1.928763 0.231024 -2.810695

- h 0.819580 -0.539513 -2.068237
- o 3.638274 0.934826 -2.175828
- h 4.439188 0.598353 -2.590121
- h 3.774372 0.839713 -1.224527

pTS-.H+W12A

с	5.579711	-0.039769	0.065392
с	4.849410	0.180547	1.236793
h	5.373242	0.300369	2.178940
с	3.458562	0.244684	1.219782
h	2.901130	0.404803	2.133985
с	2.787560	0.084058	0.011351
с	3.488953	-0.139487	-1.172171
h	2.953529	-0.276971	-2.102724
с	4.876515	-0.200360	-1.135594
h	5.423022	-0.380129	-2.055540
с	7.086537	-0.107276	0.084219
h	7.442884	-1.061740	-0.314117
h	7.522014	0.684210	-0.533166
h	7.478873	0.002690	1.096406
S	0.999431	0.234235	-0.030735
0	0.680086	1.674775	-0.266270
0	0.508716	-0.236289	1.301388
0	0.532556	-0.631493	-1.160650
0	-0.674191	3.469410	1.342976
h	-0.076798	2.898826	0.814118
h	-0.184273	4.272606	1.545240
0	-2.879656	3.322597	-0.497661

- h -2.270435 3.515853 0.236724
- h -2.283843 3.209740 -1.274272
- o -0.954477 2.785008 -2.395203
- h -0.253959 2.493148 -1.788058
- h -1.147235 1.986760 -2.912938
- o -1.215650 0.025732 -3.177698
- h -0.549205 -0.224706 -2.500077
- h -0.944595 -0.392550 -4.000966
- o -3.729834 -0.488841 -1.969441
- h -2.912060 -0.300585 -2.466831
- h -3.645999 -1.415550 -1.668178
- o -4.338617 1.239140 -0.133186
- h -3.777285 2.083769 -0.271292
- h -4.115883 0.568863 -0.860961
- h -4.134010 0.807932 0.824752
- o -3.848596 0.236488 2.078624
- h -3.011406 0.593336 2.459809
- h -3.731677 -0.742936 2.033683
- o -3.324578 -2.431673 1.796522
- h -3.452189 -2.729319 0.879303
- h -2.383286 -2.591156 1.980566
- o -3.232909 -3.084594 -1.021508
- h -2.246616 -3.210224 -1.033384
- h -3.620932 -3.866298 -1.427090
- o -0.567920 -3.268863 -0.807053
- h -0.127614 -2.426794 -1.020373
- h -0.432313 -3.359493 0.152513
- o -0.434379 -2.732439 1.982650
- h 0.025274 -3.102391 2.742812

- h -0.040647 -1.848403 1.817888
- o -1.420997 1.187777 2.913644
- h -0.742542 0.680017 2.436544
- h -1.261402 2.102954 2.630514

pTS-.H+W12B

с	5.570225	-0.084595	0.056668
с	4.795411	-0.647198	1.073272
h	5.281340	-1.171060	1.889210
с	3.405303	-0.543967	1.064991
h	2.811486	-0.971731	1.862269
С	2.784218	0.132628	0.021678
С	3.532152	0.708259	-1.006221
h	3.039173	1.245486	-1.807024
с	4.915273	0.594641	-0.980882
h	5.498746	1.045105	-1.777133
с	7.074885	-0.190490	0.069548
h	7.443927	-0.666608	-0.843625
h	7.537320	0.799507	0.127925
h	7.426753	-0.775704	0.920556
S	0.993698	0.249213	-0.026966
0	0.667326	1.666283	-0.369704
0	0.502351	-0.162785	1.315500
0	0.534218	-0.687928	-1.114100
0	-0.710523	3.530093	1.145904
h	-0.106853	2.937410	0.649777
h	-0.218546	4.336658	1.329237
0	-2.892474	3.298022	-0.705867

- h -2.291836 3.528276 0.025294
- h -2.287942 3.154428 -1.470805
- o -0.943119 2.675340 -2.549678
- h -0.263721 2.400431 -1.910997
- h -1.136645 1.854839 -3.031458
- o -1.191096 -0.112150 -3.186986
- h -0.532136 -0.332671 -2.492133
- h -0.917635 -0.573452 -3.986011
- o -3.727914 -0.569769 -1.974159
- h -2.907344 -0.408801 -2.474934
- h -3.618625 -1.460000 -1.560161
- o -4.326813 1.218643 -0.271273
- h -3.771643 2.065234 -0.436685
- h -4.090722 0.503317 -0.977614
- h -4.126797 0.843788 0.677626
- o -3.810017 0.318899 2.029976
- h -2.977761 0.698899 2.401943
- h -3.723781 -0.651728 2.105262
- o -3.313278 -2.456661 2.121308
- h -3.721628 -2.998225 2.804189
- h -2.330135 -2.556438 2.224276
- o -3.254842 -2.947351 -0.718938
- h -2.310526 -3.167003 -0.785018
- h -3.430660 -2.875851 0.236023
- o -0.353834 -3.262146 -0.704969
- h 0.035394 -2.379316 -0.892073
- h 0.179884 -3.908436 -1.177319
- o -0.639395 -2.689216 2.074619
- h -0.472762 -3.127519 1.222219

- h -0.215524 -1.820048 1.964096
- o -1.421655 1.342254 2.861436
- h -0.718881 0.814773 2.445227
- h -1.272020 2.236418 2.512703

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