Characteristics of Hydrogen Bonding between Chalcogen Matthew H. V. Graneri, Duncan A. Wild and Allan J. McKinley and Pnictogen Species **School of Molecular Sciences**



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Hydrogen bonding is one of the most prevalent types of intermolecular interactions in nature. While many O-H···B and O···H-A systems have been investigated, complexes involving non-oxygenated N-R₃ and sulfur-based molecules exclusively have been received less attention... We conducted matrix isolation Fourier transform infrared (MI-FTIR) spectroscopy studies using gas mixtures of argon, H₂S and several nitrogen species in order to investigate these interactions.

(b)

$H_2S \cdot NH_3$

Hydrogen sulfide (H_2S) and ammonia (NH_3) are emitted into the atmosphere through both natural and anthropogenic processes, yet their interaction is largely unexplored. This is particularly surprising, as NH₄SH ice clouds are believed to be responsible for the colour of Jupiter's Great Red Spot,¹ which has not yet been fully explained.

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$H_2S \cdot Amines$

Clouds have a significant effect on global energy uptake in the atmosphere. The presence and behaviour of clouds is largely dependent on cloud condensation nuclei $(CCN)^3$ – hydrophilic aggregates of atmospheric particles upon which water vapour can condense. CCN generally include sulfur-containing molecules, such as H₂SO₄.⁴ It has recently been found that nitrogen-containing molecules, including NH₃, methyl-, dimethyl-, and trimethylamine (MA, DMA, TMA, respectively) significantly enhance the ability of H₂SO₄ to attract water molecules.⁵ We looked at the interactions



Figure 1: MI-FTIR difference spectra of mixtures containing various isotopologues of H₂S and NH₃ in solid argon. Spectra were recorded following deposition at 10 K. The spectra in (a) show the v₁ and v_3 stretching regions of H₂S, while the spectra in (b) show the v_2 'umbrella' vibration region of NH_3 .²

Table	1: Fre	equency	shifts	(in cm ⁻¹) as a	result c	of comp	lexation	between
H ₂ S ar	nd NH ₃	with res	spect to	monom	ner vibra	ational fi	requence	cies. ²	

Vibration	$^{1}\text{H}_{2}\text{S} \cdot ^{14}\text{NH}_{3}$	${}^{1}\text{H}_{2}\text{S} \cdot {}^{15}\text{NH}_{3}$	$D_2S^{-14}NH_3$	$^{1}\text{H}_{2}\text{S} \cdot \text{ND}_{3}$	$D_2S \cdot ND_3$
NH_3, V_2	+ 31.8	+ 31.3	+ 32.7	+ 23.9	+ 24.5
H_2S, v_2	+ 15.1	+ 15.0	+ 11.1		
NH_3 , V_4 (sym)	- 21.1	-21.0		- 8.1	- 7.7
NH_3 , V_4 (asym)	- 6.8	- 6.8			

between these amines and H_2S , the sulfur analogue of H_2O , in order to explore the interactions between sulfur and nitrogen-containing species and create a 'series' of data which could be used to predict the binding of other such molecules.

The binding energy of the complex can be qualitatively estimated by the shift in the v_1 vibration of H_2S . The trend determined from our spectra (DMA > TMA > MA) does not match that of the gas phase (GP) basicity data, nor does it match our calculated values (see Figure 3, Table 2). This may be due to interference from the Ar matrix. We are in the process of investigating this possibility.

New peaks appear at higher energy wavelengths to each H₂S•amine peak (see Figure 3). They are present in every H₂S•amine spectrum. Their origins are currently unknown.



Figure 3: MI-FTIR difference spectra of mixtures containing H₂S and MA, DMA or TMA in solid argon. Spectra were recorded following deposition at 10 K.

Table 2: Comparison of data related to MA, DMA and TMA or their complexes with H_2S .

H_2S, v_1	- 155.3	- 156.0	- 105.8	- 160.3	- 108.9
H_2S, v_1^*	- 145.4	-146.4		- 148.9	
NH ₃ , v ₁	- 21.5	- 22.1	- 22.3		
NH_3 , $V_{3 (sym)}$	- 24.1	- 24.3	- 24.2		
NH ₃ , v _{3 (asym)}	- 13.4	- 13.5	- 13.9		

Splitting of the v_1 vibration of 1H_2S upon complexation indicates the presence of two different trapping sites; the absence of the splitting in the D₂S/NH₃ spectra suggests that in one site (marked *), the ¹H₂S subunit undergoes hindered rotation, tunnelling through the Ar matrix, while, in the other site, the rotation is completely suppressed.²



Figure 2: The H₂S•NH₃ complex, optimised at CCSD(T)/aug-cc-pV(T+d)Z level of theory.

Complex	$\Delta v_1(H_2S)$	Amine Basicity, kJ mol ⁻¹ (GP) ⁶	Amine Basicity (Aq., pK _{aH}) ⁷	Binding Energy (kJ mol ⁻¹)*
$^{1}H_{2}S \cdot NH_{3}$	-155.3	819.0		7.3
$^{1}H_{2}S \cdot MA$	-263.8	864.5	10.62	12.4
¹ H ₂ S•DMA	-305.8	896.5	10.64	14.6
$^{1}H_{2}S \bullet TMA$	-277.8	918.1	9.76	15.4

*CCSD(T)-RI/F12/AVQZ-F12//B97M-D4/AVQZ



Figure 4: Most tightly bound conformers calculated for the H₂S•amine complexes. The structure in (a) is of the H_2S ·MA complex, calculated at CCSD(T)/aug-cc-pV(T+d)Z level of theory. The structures in (b) and (c) are of the H₂S•DMA and H₂S•TMA complexes, respectively, calculated at B97M-D4/aug-cc-pV(Q+d)Z level of theory.

$H_2O \cdot PH_3$

PH₃ is considered a biosignature gas in the search for extra-terrestrial life in anoxic environments.⁸ Surprisingly, the 1:1 H_2O •PH₃ complex had not been investigated, despite the importance of H_2O for the propagation of life. We sought to rectify this using MI-FTIR to identify the complex.

A binding energy of 5.1 kJ mol⁻¹ was calculated at $CCSD(T)/CBS_{T,Q,5}$ level of theory; much weaker than its S/N analogue, H_2 S•N H_3 , with 8.6 kJ mol⁻¹ at the same level of theory.²



isotopologues of H_2O and PH_3 in Ar matrices.

1. M. J. Loeffler, R. L. Hudson, N. J. Chanover and A. A. Simon, *Icarus*, 2016, 271, 265-268.

2. M. H. V. Graneri, D. A. Wild and A. J. McKinley, J. Mol. Spectrosc., 2021, 378, 111440.

3. B. A. Albrecht, Science, 1989, 245, 1227-1230.

4. M. Kulmala, T. Petäjä, M. Ehn, J. Thornton, M. Sipilä, D. R. Worsnop and V.-M. Kerminen, Annu. Rev. Phys. Chem., 2014, 65, 21-37.

5. T. Kurtén, V. Loukonen, H. Vehkamäki and M. Kulmala, Atmos. Chem. Phys., 2008, 8, 4095-4103.

6. E. P. L. Hunter and S. G. Lias, J. Phys. Chem. Ref. Data, 1998, 27, 413-656.

7. S. Tshepelevitsh, A. Kütt, M. Lõkov, I. Kaljurand, J. Saame, A. Heering, P. G. Plieger, R. Vianello and I. Leito, Eur. J. Org. Chem., 2019, 2019, 6735-6748.

8. C. Sousa-Silva, S. Seager, S. Ranjan, J. J. Petkowski, Z. Zhan, R. Hu and W. Bains, Astrobiology, 2020, 20, 235-268.

Figure 5: The $H_2O \cdot PH_3$ complex, optimised at CCSD(T)/aug-cc-pV(T+D)Z level of theory.