A velocity map ion imaging study of difluorobenzene-water complexes: Binding energies and recoil distributions

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The binding energies of the \( p \)-, \( m \)-, and \( o \)-difluorobenzene-H\(_2\)O complexes have been measured by velocity map ion imaging to be 922 \( \pm \) 10, 945 \( \pm \) 10, and 891 \( \pm \) 4 cm\(^{-1}\), respectively. The lack of variation provides circumstantial evidence for water binding to the three isomers via the same interaction, viz. an in-plane O–H \( \cdot \cdot \cdot \) F hydrogen bond to one of the fluorine atoms on the ring, with a second, weaker interaction of the water O atom with an \( o\)-hydrogen, as determined previously for the \( p\)-difluorobenzene-H\(_2\)O complex [Kang et al., J. Phys. Chem. A 109, 767 (2005)]. The ground state binding energies for the difluorobenzene-H\(_2\)O complexes are \(-5\%\) to \(-11\%\) larger than that for benzene-H\(_2\)O, where binding occurs to the \( \pi \) electrons out-of-plane. However, in the \( S\)\textsubscript{1} state the binding energies of the \( o \)- and \( p \)-difluorobenzene-H\(_2\)O complexes are smaller than the benzene-H\(_2\)O value, raising an interesting question about whether the geometry at the global energy minimum remains in-plane in the excited electronic states of these two complexes. Recoil energy distributions for dissociation of \( p \)-difluorobenzene-H\(_2\)O have been measured from the \( 3^7\), \( 5^7 \), and \( 3^7\) \( \cdot \) \( \bar{3}^7\) levels of the excited electronic state. These levels are 490, 880, and 1304 cm\(^{-1}\), respectively, above the dissociation threshold. Within the experimental uncertainty, the recoil energy distributions are the same for dissociation from these three states, with average recoil energies of \( \sim 100 \) cm\(^{-1}\). These recoil energies are 60\% larger than was observed for the dissociation of \( p \)-difluorobenzene-Ar, which is a substantially smaller increase than the 400\% seen in a comparable study of dissociation within the triplet state for pyrazine-Ar, -H\(_2\)O complexes. The majority of the available energy is partitioned into vibration and rotation of the fragments. © 2008 American Institute of Physics. [DOI: 10.1063/1.2896081]

I. INTRODUCTION

The study of molecular complexes and clusters in the gas phase provides an important link with condensed phase behavior, with clusters considered as models for reagent surrounded by a specific number of solvent molecules. Complexes involving water are of significance, with particular interest in water complexed to an aromatic molecule because of the relevance to biological processes. Benzene-water complexes have been extensively studied experimentally and theoretically to determine the nature of the interaction as a model for hydrophobic interactions, as discussed in several reviews. The binding energy and geometry of the benzene-water dimer have been determined. A weak \( \pi \)-hydrogen bond is formed between the aromatic \( \pi \) electrons and one of the hydrogen atoms on the water molecule. The water molecule is located above the aromatic ring with both O–H bonds pointing toward the ring.

Substituents on the ring can significantly alter the aromatic-water interaction. Fluorine’s influence on the aromaticity of the ring has recently been explored in detail. There is a net destabilizing of the aromatic benzene system by the addition of fluorine, although \( \pi \) back bonding increases the electron density in the ring. In the case of \( p \)-difluorobenzene (\( p\)DFB) complexes, it has been established experimentally that the water molecule does not interact with the \( \pi \) cloud. Instead, it forms an in-plane O–H \( \cdot \cdot \cdot \) F hydrogen bond to one of the fluorine atoms on the ring, with a second, weaker interaction of the water O atom with one of the two \( o\)-hydrogens. This is consistent with earlier calculations. The remaining H atom appears to lie out of the plane. This change in the nature of the aromatic-water interaction has led us to explore the complexes of water with the series of difluorobenzenes (DFBs), viz. \( para\), \( meta\), and \( ortho\). The \( meta\) - and \( ortho\)-DFB-water complexes have been probed previously using resonance enhanced multiphoton ionization (REMPI). A rotational contour analysis of the origin bands suggests that the complexes involve an in-plane O–H \( \cdot \cdot \cdot \) F hydrogen bond analogous to that seen in \( p\)DFB-water.

The aim of our study is twofold. The first is to determine the binding energies of water to the DFBs and hence provide data on the strength of the water-difluorinated benzene interaction and on the extent to which the change in the relative positions of the fluorine atoms, which alters the aromaticity, affects the strength of the interaction. The second is to explore the dissociation dynamics of the van der Waals complex, specifically the partitioning of energy into translation. This latter study is focused on the \( p\)DFB-water complex as there are data for the \( p\)DFB-Ar complex for comparison.

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For pDFB-Ar dissociation we found that (a) the translational energy release is low (average recoil energy \(~63\) cm\(^{-1}\)), (b) there is significant angular momentum in the pDFB fragment, and (c) the recoil energy distribution is insensitive to the initial vibrational energy and destination state density up to \(~3000\) cm\(^{-1}\) available energy. Yoder and Barker found in studies of dissociation of aromatic complexes from triplet states that H-bonded complexes released significantly more energy in translation than those involving dispersion interactions. We here explore whether this trend is seen for the \(S_1\) pDFB-H\(_2\)O versus pDFB-Ar complexes.

These issues will be explored using the technique of velocity map ion imaging. This technique is ideal for studies of van der Waals molecule dissociation where the translational energy released is small (typically \(<1000\) cm\(^{-1}\)) because the image resolution is linear in velocity, and so the energy resolution is highest at low translational energies.

A number of studies of the pDFB-H\(_2\)O complex have been reported,\(^7\) however, studies of the ortho and meta isomers appear restricted to a Ph.D thesis.\(^10\) The \(S_1\) \(\rightarrow\) \(S_0\) transition of the pDFB-H\(_2\)O complex has been studied using both one and two-color 1+1 REMPI.\(^8,14\) The pDFB-H\(_2\)O origin transition is 169 cm\(^{-1}\) higher in energy than the pDFB monomer origin: The complex is more weakly bound in the excited electronic state. Similar blueshifts were reported for the other vibronic bands: \( \tilde{v}_0^0\) (+169 cm\(^{-1}\)), \( \tilde{v}_0^{1}\) (+176 cm\(^{-1}\)), and \( \tilde{v}_0^{2}\) (+158 cm\(^{-1}\)). The lack of progressions in the intermolecular modes suggested that the complex has similar geometries in the ground and excited electronic states. The geometry was explored in detail in a rotationally resolved study of the \(S_1\) \(\rightarrow\) \(S_0\) spectrum by Kang et al.\(^7\) They found a tunneling splitting of bands in the spectrum, showing that the attached water undergoes a combined inversion-internal rotation motion. The H-bonding interaction has also been explored through the O–H stretch in the infrared.\(^15\) The ionization potential of pDFB-H\(_2\)O was determined using two color 1+1 REMPI to be 72.450 ± 50 cm\(^{-1}\),\(^14\) although Brenner et al. note that as their calculations suggest a significant difference in geometry between the neutral and ionic states, the measured ionization threshold may be higher than the adiabatic ionization potential.\(^8\)

The \(S_1\) \(\rightarrow\) \(S_0\) transitions of the mDFB-H\(_2\)O and oDFB-H\(_2\)O complex have been studied using mass resolved REMPI.\(^10\) The mDFB-H\(_2\)O origin transition is 92 cm\(^{-1}\) blueshifted from the mDFB monomer, while the oDFB-H\(_2\)O origin transition is 105 cm\(^{-1}\) blueshifted from the oDFB monomer. Thus, like pDFB-H\(_2\)O, both complexes are more weakly bound in the excited electronic state, although the change in bond strength on electronic excitation is reduced in these two isomers.

**II. EXPERIMENTAL DETAILS**

The experimental details for the velocity map imaging experiments have been given previously.\(^16\) Complexes are formed in a supersonic free jet expansion of the DFB (1%) and water (1%) in Ar at 6 atm backing pressure. Complexes are excited in the extraction region of a time of flight mass spectrometer (ToFMS) operating in velocity map imaging mode\(^13\) using the frequency doubled output of a Nd:YAG pumped dye laser (pulse duration of about 4 ns; frequency doubled line width of about 0.4 cm\(^{-1}\)). As is discussed in the following sections, the complexes dissociate within the \(S_1\) state, producing \(S_1\) DFB products that can absorb a second photon and be ionized. The ions are accelerated down the ToFMS and detected using a gated microchannel plate/phosphor screen combination with a charge coupled device camera. Images are downloaded shot-by-shot and analyzed by a laboratory computer to accumulate a histogram of ion count versus position. The images are isotropic, indicating that dissociation is slow compared with molecular rotation, and are reduced to plots of intensity versus radius (so-called radial plots) for analysis. Conversion of the radial plots to the original three-dimensional distribution is undertaken using the inverse Abel function method and the total energy distribution determined as described in Ref. 16. Ion images were calibrated using photoelectron images. The laser power was maintained such that only one to two ions were observed with each laser shot to eliminate Coulomb repulsion effects.

Spectra were obtained using the method of velocity and mass resolved REMPI.\(^17\) Here, the center of each image and a surrounding annulus are integrated separately. DFB ions originating from dissociation of the complex gain kinetic energy and contribute intensity away from the center of the image, i.e., in the annulus. Table I summarizes the transition frequencies measured for pDFB-H\(_2\)O in key regions. The blueshift is reduced at higher vibrational energies.

<table>
<thead>
<tr>
<th>Transition</th>
<th>Wavenumber (cm(^{-1}))</th>
<th>Shift from pDFB band (cm(^{-1}))</th>
<th>Note</th>
</tr>
</thead>
<tbody>
<tr>
<td>( \tilde{v}_0^0 )</td>
<td>37 359</td>
<td>163</td>
<td>( h\alpha )</td>
</tr>
<tr>
<td>( \tilde{v}_0^{1})</td>
<td>37 376</td>
<td>180</td>
<td></td>
</tr>
<tr>
<td>( \tilde{v}_0^{2})</td>
<td>37 812</td>
<td>168</td>
<td></td>
</tr>
<tr>
<td>( \tilde{v}_0^1 )</td>
<td>37 824</td>
<td>168</td>
<td></td>
</tr>
<tr>
<td>( \tilde{v}_0^2 )</td>
<td>37 828</td>
<td>170</td>
<td></td>
</tr>
<tr>
<td>( \tilde{v}_0^3 )</td>
<td>38 250</td>
<td>161</td>
<td>( c )</td>
</tr>
<tr>
<td>( \tilde{v}_0^4 )</td>
<td>38 255</td>
<td>166</td>
<td></td>
</tr>
<tr>
<td>( \tilde{v}_0^5 )</td>
<td>38 638</td>
<td>164</td>
<td></td>
</tr>
<tr>
<td>( \tilde{v}_0^6 )</td>
<td>38 640</td>
<td>164</td>
<td></td>
</tr>
<tr>
<td>( \tilde{v}_0^7 )</td>
<td>39 065</td>
<td>158</td>
<td></td>
</tr>
<tr>
<td>( \tilde{v}_0^8 )</td>
<td>39 069</td>
<td>162</td>
<td></td>
</tr>
</tbody>
</table>

\( ^a \) The pDFB-H\(_2\)O \( 0^0_0 \) band is 169 cm\(^{-1}\) higher in energy than that for pDFB (Refs. 8 and 14).

\( ^b \) Broad bands with unresolved substructure.

\( ^c \) Probable Fermi resonance.
III. RESULTS

A. DFB-water binding energies

The binding energies for the DFB-H2O complexes are determined using the technique of velocity map imaging discussed in Ref. 18. In essence, the complex is excited to a vibrational level not far above the S1 dissociation energy and the translational energy distribution of the products measured. Because of the limited energy available to the products, the translational energy distribution is truncated. The binding energy is determined from the maximum translational energy released.

One of the keys to performing the experiment is determining an appropriate vibronic transition of the complex, which would allow measurement of dissociation in the ion with the values for the neutral subsequently calculated, one requires rapid dissociation on the S1 surface to prevent the complication of ionization of intact complexes and their subsequent dissociation. A second requirement is that the vibrational level excited must be sufficiently close to dissociation that the energy at which the recoil energy truncates is readily identified.

For pDFB-H2O, limits for the binding energy have been determined experimentally.8,14 An upper limit of 818 cm−1 was set by the appearance of fragment ions following S0 excitation. The energy of two photons at this wavelength is well above the pDFB-H2O ionization potential so that fragmentation could occur from both excited and ionic states; however, a two-color 1+1 REMPI experiment established that the dissociation observed in the one-color experiment occurs from S1. The lack of dissociation products following S0 excitation implied a lower limit of 410 cm−1 for S1 dissociation. However, this was refined upwards as the failure of the ion to decompose after 1+1 REMPI via S0 suggests a lower limit of 2234 cm−1 for the ion. Combined with the spectral shift for ionization, this gave a lower bound of 794 cm−1 in S1. As we discuss later, this lower limit is less definite than the upper bound.

Figure 1 shows the recoil energy distribution for the products produced following excitation of S1 in pDFB-H2O. The distribution is truncated because the translational energy available to the fragments in the dissociation process is limited. The cutoff is not as sharp as seen for pDFB-Ar,-Kr complexes,16 presumably because some excess energy is partitioned into rotational motion of the H2O fragment. To determine the position of the cutoff, we fit the signal and background components to quadratic functions. (The background comes from multiphoton ionization of pDFB remaining in the chamber between laser shots.) The intersection of these two curves (the solid lines in the figure) gives the maximum translational energy in the products, which in this case is 65±10 cm−1. From this value and the vibrational energy of S1, the binding energy in the excited state is calculated to be 753±10 cm−1. Binding energies for the remaining states are calculated from the shift in ionization potential [1421±50 cm−1 (Ref. 14)] and the 169 cm−1 S1→S0 complex shift. They are summarized in Table II.

In the case of mDFB-H2O, Springfield noted a loss of features above 600+931 cm−1 at the parent mass in the REMPI spectrum, indicating rapid dissociation.10 The 600+931 cm−1 feature itself is also significantly diminished relative to the corresponding feature in the mDFB spectrum. These observations, coupled with the pDFB-H2O dissociation energy, strongly suggest that dissociation occurs within 931 cm−1. An image taken at the mDFB mass with excitation on the 600+931 cm−1 band of mDFB-H2O reveals the truncation characteristic of excitation just above the dissociation onset. Following the approach outlined above for pDFB complex, we determine a maximum excess energy of 78±10 cm−1. The corresponding binding energies are given in Table II.

For oDFB-H2O, Springfield found a reduction in intensity at the parent mass in the REMPI spectrum above a

![Figure 1. The recoil energy distribution following dissociation of pDFB-H2O from the 55 level (600+817 cm−1). The points are the experimental data while the solid curves are a fit to the signal and background regions of the distribution (see text).](image-url)

**TABLE II. Experimentally determined binding energies for the DFB–H2O complexes.**

<table>
<thead>
<tr>
<th>Isomer</th>
<th>Electronic state</th>
<th>Dissociation energy (cm−1)</th>
<th>This work</th>
<th>Martrenchard et al.</th>
</tr>
</thead>
<tbody>
<tr>
<td>para</td>
<td>S0 (D0)</td>
<td>927±10</td>
<td>963&lt;D0</td>
<td>987</td>
</tr>
<tr>
<td></td>
<td>S1 (D1)</td>
<td>758±10</td>
<td>794&lt;D1</td>
<td>818</td>
</tr>
<tr>
<td></td>
<td>D0 (D0)</td>
<td>2348±51</td>
<td>2334&lt;D0</td>
<td>2458</td>
</tr>
<tr>
<td>meta</td>
<td>S0 (D0)</td>
<td>945±10</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>S1 (D1)</td>
<td>853±10</td>
<td></td>
<td></td>
</tr>
<tr>
<td>ortho</td>
<td>S0 (D0)</td>
<td>891±4</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>S1 (D1)</td>
<td>786±4</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

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The signal was insufficient to obtain images of satisfactory quality for the higher energy, more intense of the pair. On the contrary, the higher vibrational levels investigated dissociation also occurs in $S_1$. The recoil energy distributions obtained from the images are shown in Fig. 2. The points represent the inverse Abel transformed data and the solid line is the fit by a function of the form

$$P(E) = \sqrt{E} \sum_{i=1}^{2} A_i e^{-k_i E},$$

(1)

which is equivalent to the sum of two Maxwell–Boltzmann distributions. The parameters for the fits to the distributions are given in Table III. As we have observed previously, this functional form provides an excellent fit to the data.

The three recoil distributions are very similar. They quickly rise to a maximum at 15–25 cm$^{-1}$, following which they decay in an exponential-like fashion to zero by ~450 cm$^{-1}$. The average kinetic energy released to the fragments from each energy level investigated is given in Table IV. As we have found for the $p$DFB–Ar, $p$DFB+–Ar, and benzene+–Ar complexes, most of the energy remains as internal energy of the fragments. There is essentially no change in the average kinetic energy released with increasing available energy.

**IV. DISCUSSION**

**A. DFB-water binding energies**

The $S_0$ binding energies for the three difluorobenzene isomers with water are very similar, lying in the range of 919 cm$^{-1} \pm 3\%$. It has been established in $p$DFB–H$_2$O that water forms an in-plane O–H···F hydrogen bond to one of the fluorine atoms on the ring, with a weaker interaction of the water O atom with an ortho hydrogen. The lack of variation in binding energies is circumstantial evidence for water binding to the three DFB isomers via the same interaction although, as we discuss below, the interaction is not significantly stronger than that seen for the out-of-plane interaction with the $\pi$ electrons in benzene–H$_2$O. An in-plane structure for $p$DFB–H$_2$O and $p$DFB+–H$_2$O is consistent with the rotational analysis of Springfield. The experimental data does not allow us to comment on the possible existence of a second F···H–O bond in the ortho and meta in-plane complexes.

Interestingly, the variation between isomers doubles, and their ordering changes, in the $S_1$ state. $p$DFB has the weakest bond in $S_0$, while $p$DFB has the weakest interaction in $S_1$. The binding is largest for $m$DFB in both states. This is presumably a result of the change in electron density in the excited electronic states of the different isomers. The lack of advantage at high energy.

<table>
<thead>
<tr>
<th>Transition</th>
<th>$A_1$</th>
<th>$k_1$</th>
<th>$A_2$</th>
<th>$k_2$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$\tilde{3}^1$</td>
<td>0.001 73</td>
<td>0.0228</td>
<td>0.000 767</td>
<td>0.0115</td>
</tr>
<tr>
<td>$\tilde{2}^0$</td>
<td>0.001 82</td>
<td>0.0651</td>
<td>0.001 66</td>
<td>0.0139</td>
</tr>
<tr>
<td>$\tilde{3}^0$</td>
<td>0.001 51</td>
<td>0.0136</td>
<td>0.001 70</td>
<td>0.0450</td>
</tr>
</tbody>
</table>

**Table IV. Values for the average translational energy released during the dissociation of $p$DFB–H$_2$O.**

<table>
<thead>
<tr>
<th>Initial Level</th>
<th>$S_1$ Vibrational Energy (cm$^{-1}$)</th>
<th>Excess Energy (cm$^{-1}$)</th>
<th>Average recoil Energy (cm$^{-1}$)</th>
</tr>
</thead>
<tbody>
<tr>
<td>$\tilde{3}^1$</td>
<td>1243</td>
<td>490</td>
<td>101</td>
</tr>
<tr>
<td>$\tilde{2}^0$</td>
<td>1633</td>
<td>880</td>
<td>100</td>
</tr>
<tr>
<td>$\tilde{3}^0$</td>
<td>2057</td>
<td>1304</td>
<td>97</td>
</tr>
</tbody>
</table>
ionization energies for the ortho and meta species prevents us from determining the binding of water to the corresponding ions. However, a value can be determined for pDFB, where the interaction is $\sim 2.5$ times larger than for the neutral complex.

The pDFB–H$_2$O binding energy limits determined in the REMPI study of Martrenchard et al. are included in Table II. For the neutral, our imaging values lie below the lower bounds determined previously while our value for the cation is within the bounds set by Martrenchard et al. The lower bound set by Martrenchard et al. is too high. It was deduced from the failure to observe dissociation from the ion after $1 + 1$ REMPI via $6^1\Omega_2$. When considering dissociation from the ion, not all of the energy provided by the two photons need reside in the ion. A distribution of population will be produced in the ion, governed by the Franck–Condon factors for the ionization process, and the photoelectron will remove any excess energy. By this means, it is possible to have situations where the photon energy absorbed exceeds the dissociation limit, but dissociation is not observed as the ions are formed in vibrational levels below dissociation. It appears that this is the situation for excitation of pDFB–H$_2$O via $6^1\Omega_2$.

The fluorine substitution on the aromatic ring leads to a different binding site (in-plane to F) compared to that for the parent system benzene–H$_2$O (B–H$_2$O; out of plane to the $\pi$-electron system). It is thus interesting to compare the DFB–H$_2$O binding energies with that for B–H$_2$O. The most precise determination of the $S_1$ binding energy for B–H$_2$O has reported $853 \pm 32$ cm$^{-1}$. The $S_0$ binding energies for the DFB–H$_2$O complexes are $891 \pm 4$, $945 \pm 10$, and $922 \pm 10$ cm$^{-1}$ for ortho-, meta-, and para–DFB. The DFB–water values are 4.5%, 10.8%, and 8.1% larger for ortho–DFB, which suggests that the out-of-plane interaction might be enhanced in the DFBs compared with benzene. While the three DFB–H$_2$O binding energies are larger than the B–H$_2$O value, the increase is quite small, suggesting that the in-plane configuration is only marginally lower in energy than the out-of-plane one. The increase in binding energy for oDFB–H$_2$O is particularly small suggesting that there is little difference in energy between the in-plane and out-of-plane geometries in this case.

While the $S_0$ binding energies for the DFB–H$_2$O complexes are uniformly larger than the B–H$_2$O value, this is not the case for the $S_1$ values. The $S_1$ values are B–H$_2$O $803 \pm 32$ cm$^{-1}$, oDFB–H$_2$O $786 \pm 4$ cm$^{-1}$, mDFB–H$_2$O $853 \pm 10$ cm$^{-1}$, and pDFB–H$_2$O $753 \pm 10$ cm$^{-1}$. The oDFB–H$_2$O and pDFB–H$_2$O values are smaller than the B–H$_2$O value. This raises the interesting possibility that the out-of-plane geometry is favored in the $S_1$ state for the oDFB–H$_2$O and pDFB–H$_2$O complexes. Excitation may not access the global minimum in the $S_1$ state of these complexes since the Franck–Condon factors will favor excitation to the in-plane geometry which may simply be a local minimum. These systems would be excellent candidates for high level ab initio calculations to investigate this issue.

Calculations have not been reported for the dissociation energies of the mDFB–H$_2$O and oDFB–H$_2$O complexes; however, they have been published for pDFB–H$_2$O (Ref. 9) and are summarized in Table V. The calculations were all at the MP2 level of theory but varied with respect to basis set. Within the quoted theoretical uncertainty all of the basis sets give a value matching the experimental value although the MP2/6–31++G(3df,2p) value is closest.

### B. Recoil energy distributions

The recoil energy distributions for the three vibrational levels of pDFB–H$_2$O are displayed in Fig. 3 with the intensity on a logarithmic scale. A distribution for the dissociation of pDFB–Ar from the $3^2S^1$ level has been included for comparison.

![A comparison of the recoil distributions for pDFB–H$_2$O with a representative distribution for pDFB–Ar.](image-url)
Comparison. The $S_1$ pDFB-Ar recoil distributions are all very similar, so the $3^{1S}_{1}$ distribution is representative of this complex. The pDFB-H$_2$O distributions are very similar as a group, although clearly different to the Ar case, showing a larger proportion of molecules with higher translational energy release.

Recoil energy distributions for dissociation occurring within the triplet state ($T_1$) of a number of aromatic van der Waals clusters have been determined using a velocity resolved time-of-flight technique by Yoder and Barker. Figure 4 shows translational energy distributions obtained by these authors for the dissociation of pyrazine-Ar and pyrazine-H$_2$O clusters, reproduced from the data provided in their paper. A long tail to significantly higher energy occurs in the distribution for the pyrazine-H$_2$O complex compared with the pyrazine-Ar complex. Interestingly, the pyrazine-Ar distribution is indistinguishable from the pDFB-H$_2$O distributions. Experiments indicate that pyrazine-H$_2$O contains a hydrogen bond between one of the hydrogen atoms on the water molecule and a nitrogen atom in the pyrazine ring. Similar to the pDFB-H$_2$O complex, the water molecule lies in the plane of the ring.

Yoder and Barker have postulated that the difference in the distributions for the pyrazine-Ar and pyrazine-H$_2$O clusters is due to the distinction between hydrogen bonding and nonlocalized dispersive van der Waals interactions. The relatively weak dispersive interactions in aromatic-Ar means the argon molecule is able to migrate over the aromatic ring. Trajectory calculations on the pyrazine-Ar complex have shown that, as energy is transferred from the initially excited pyrazine vibrations to the intermolecular modes, the Ar atom samples other minima on the surface until the energy in the intermolecular modes exceeds the binding energy and the molecule dissociates. For pyrazine-water, Yoder and Barker suggest that since a hydrogen bond between the water molecule and a lone pair of electrons on a highly electronegative atom is localized; no change in geometry of the cluster occurs without breaking the bond. They have argued that this allows more energy to be partitioned into translational energy rather than rotational energy of the fragments because the initial geometry of the cluster is constrained prior to dissociation.

The distributions reported by Yoder and Barker reveal average recoil energies of 100 and 111 cm$^{-1}$ for their two pyrazine-Ar distributions and 425 cm$^{-1}$ for pyrazine-H$_2$O. Our velocity map imaging measurements of pDFB van der Waals molecules give average recoil energies of $\sim$62 and $\sim$100 cm$^{-1}$ for Ar and H$_2$O clusters, respectively. More energy is released as translational energy in dissociation of the pDFB-H$_2$O cluster relative to the Ar cluster, but the increase is only $\sim$60%, far from the $\sim$400% increase seen in the pyrazine case.

The reason for the much higher average translational energy release for the pyrazine-H$_2$O complex compared to pDFB-H$_2$O is unclear. The major differences between the pyrazine cluster dissociation and our measurements for pDFB are that dissociation of the pyrazine clusters is from the lowest triplet state and involves much higher energies. Prior to dissociation the vibrational energy of pyrazine clusters is 4056 cm$^{-1}$. Although the dissociation energy is not known for this case, and hence the excess energy is unknown, it is clearly well above the $\sim$1300 cm$^{-1}$ available for the highest vibrational level we have investigated for pDFB-H$_2$O. However, for pDFB-Ar we found that there is essentially no change in the average translational energy released with increasing initial vibrational energy and the limited data set for the water complex also shows no change with increasing vibrational energy. We thus believe it unlikely that the differences arise from the extra vibrational energy available for pyrazine-water.

Previously, we have noted that the dissociation of aromatic–rare gas complexes leads to significant rotational excitation of the aromatic fragment. It appears that this is also the case for dissociation of pDFB-H$_2$O complexes. Consider dissociation from $3^{1S}$, $3^{3T}$ is 490 cm$^{-1}$ above dissociation, so this is the maximum energy available to the products. 490 cm$^{-1}$ is well below the water vibrations and the excess energy will appear as translation of the pDFB and H$_2$O fragments, rotation of water and pDFB, and vibration of pDFB. Few vibrational levels of pDFB are accessible. The recoil energy distribution, however, shows no structure, particularly at high translational energy, which corresponds to low vibrational energy where there is a large gap between vibrational levels. This lack of structure indicates significant rotational excitation of pDFB and/or water fragments.

V. CONCLUSIONS

We have determined the binding energies of the pDFB-H$_2$O, mDFB-H$_2$O, and oDFB-H$_2$O complexes. The ground state binding energies for the three difluorobenzene isomers with water are very similar, varying by $\sim$6.5%. This lack of variation is circumstantial evidence for water binding to the three isomers via the same interaction. Based on the established binding for pDFB-H$_2$O, we suggest that in all cases water forms an in-plane O–H−···F hydrogen bond to one of the fluorine atoms on the ring, with a weaker interac-
tion of the water O atom with an ortho hydrogen, although we cannot rule out an interaction of the water H with the second F atom in the ortho and meta cases. The ground state binding energies for the DFB-H$_2$O complexes are about 5%—11% larger than that for benzene-H$_2$O. However, in the $S_1$ state the binding energies of the o- and p-difluorobenzene-H$_2$O complexes are smaller than the benzene-H$_2$O value, raising an interesting question about whether the geometry at the global energy minimum remains in-plane in the excited electronic states of these two complexes.

Distributions for the kinetic energy released in the dissociation of pDFB-H$_2$O have been measured from the $3^1T$, $5^1T$, and $3^13^1T$ levels of the excited electronic state. These levels are 490, 880, and 1304 cm$^{-1}$, respectively, above the dissociation threshold. Within the experimental uncertainty, the translational energy distributions are the same for dissociation from $3^1T$, $5^1T$, and $3^13^1T$. The corresponding average recoil energies were determined to be 101, 100, and 97 cm$^{-1}$, respectively. The majority of the energy remains as vibrational and rotational energy of the fragments. The lack of structure in the recoil distribution for dissociation from $3^1T$ indicates that there is significant rotational excitation in the fragments. For the relatively low initial vibrational energies studied, there is no change in translational energy released with increasing internal energy of the excited complex. At the energies investigated, no long tail was observed in the distributions as was observed in a previous study of the translational energy released in dissociation of a hydrogen bonded complex.12

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