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Ultrafast non-radiative decay of gas-phase nucleosides

Simone De Camillis, a Jordan Miles, a Grace Alexander, a Omair Ghafur, b Ian D. Williams, a Dave Townsend dc and Jason B. Greenwood* a

The ultrafast photo-physical properties of DNA are crucial in providing a stable basis for life. Although the DNA bases efficiently absorb ultraviolet (UV) radiation, this energy can be dissipated to the surrounding environment by the rapid conversion of electronic energy to vibrational energy within about a picosecond. The intrinsic nature of this internal conversion process has previously been demonstrated through gas phase experiments on the bases, supported by theoretical calculations. De-excitation rates appear to be accelerated when individual bases are hydrogen bonded to solvent molecules or their complementary Watson–Crick pair. In this paper, the first gas-phase measurements of electronic relaxation in DNA nucleosides following UV excitation are reported. Using a pump–probe ionization scheme, the lifetimes for internal conversion to the ground state following excitation at 267 nm are found to be reduced by around a factor of two for adenosine, cytidine and thymidine compared with the isolated bases. These results are discussed in terms of a recent proposition that a charge transfer state provides an additional internal conversion pathway mediated by proton transfer through a sugar to base hydrogen bond.

Introduction

DNA bases are strong ultraviolet (UV) chromophores which absorb at wavelengths in the range of 240–290 nm.1 These molecules are the basis for life despite the fact that the prebiotic Earth was bathed in ultraviolet radiation, suggesting that they are able to effectively quench excess absorbed energy very rapidly without generating photolesions. Similarly these ultrafast energy quenching mechanisms are relevant, for example, to the extent to which ultraviolet exposure leads to skin cancer and the efficacy of sunscreens.2 As such, the energy relaxation and dissipation mechanisms operating in these systems have been the subject of hundreds of experimental and theoretical studies over the last 20 years.

Measurements of ultrafast UV quenching in DNA were first made possible in solution through the development of the transient absorption and fluorescence up-conversion techniques,3,4 which have been used to observe ultrafast dynamics in DNA components from individual bases through to double helix strands. Theoretical techniques which can accurately describe the transient molecular dynamics observed have developed in parallel and have been essential in interpreting the experimental results.5 However, the complexity of the DNA structures means that further theoretical and experimental advances are required for a complete understanding of the ultrafast de-excitation dynamics. Complementary gas-phase experiments have been crucial in understanding the intrinsic properties of the isolated base molecules and validating the theoretical methodology. The first such measurements, undertaken by Kang et al.,6 studied the dynamics of low-lying ππ* states following initial excitation with 267 nm laser pulses and subsequent tracking of the evolution in multiphoton ionization probabilities by a delayed 800 nm pulse. These pump–probe results showed that the non-radiative de-excitation in isolated RNA and DNA nucleobases occurred within a few picoseconds or less. A subsequent study using a 400 nm probe and much better time resolution of around 80 fs7 revealed a bi-exponential decay mechanism. A fast component (<200 fs) was interpreted as wavepacket motion to the minimum of the ππ* potential energy surface and population transfer via a conical intersection to a state of nπ* character. Subsequent non-radiative transitions to the ground state were then assigned to a longer, picosecond process. A key reaction coordinate for this process corresponds to out-of-plane deformations of the aromatic ring system, which sends the nπ* and ππ* excited states towards conical intersections with the ground state.8–12

These formative experiments were followed up by more detailed studies which have mainly focussed on adenine since it is relatively easy to produce in the gas phase and has only one dominant tautomer. Of particular interest was the extent to
which additional electronic states formed via Rydberg/valence mixing were contributing to the de-excitation dynamics. Such states exhibit predominantly σ character in the Franck–Condon region, but evolve increasingly towards πσ* character at extended N–H distances. Initial studies of methylated and deuterated adenine showed no change in the lifetimes. However, time resolved photoelectron spectra (TRPES) obtained by Satzger et al. suggested that de-excitation from the πσ* state does proceed via a πσ* state (in competition with the ππ* internal conversion pathway). This contradicted theoretical calculations that predicted the πσ* state lies too high in energy to be accessed by a 267 nm pump. Ashfold et al. and, Evans and Ulrich agreed with this theoretical interpretation through measurements of H atom production and TRPES respectively, indicating πσ* states were only involved for wavelengths less than 238 nm. Similarly, for the other nucleobases a number of studies suggest that πσ* states can only be accessed at shorter wavelengths, but interpretation of the data is complicated by the presence of many more structural conformers.

There have been many more studies of other gas phase nucleobase derivatives (for example) which demonstrate that, except for simple methylation, de-excitation rates following UV excitation are generally slower for chemically modified bases and even for some isomers. These observations can, in some cases, be explained by the constraint put on out-of-plane deformations, but more generally modified nucleobases have energy barriers which restrict access to conical intersections. Indeed the canonical bases exploited for genetic encoding appear to have been chosen by natural selection as they are photochemically more stable than other variants.

For the nucleo-bases/sides/tides in solution, the lifetimes are found to be shorter than for the isolated bases in the gas-phase. The fact that Watson–Crick pairs in solution have even faster rates of decay that H-bonding to the base opens up additional quenching mechanisms involving proton transfer. For guanine–cytosine base pairs, Sobolewski and Domico have proposed that population of an inter-base charge transfer state moves an electron from guanine to cytosine followed in the same direction by a proton involved in one of the base-to-base H-bonds.

Gas phase studies of ultrafast dynamics in DNA components larger than the individual bases have yet to be undertaken. However, resonant two-photon ionization spectroscopic studies using nanosecond laser pulses have provided some indirect insight. For example, the observation of spectra which are broader for base pairs than for individual bases suggests that rapid decay via charge transfer states is also playing a role in the gas phase. For some nucleosides, it has also proved difficult to produce significant amounts of parent ions in the mass spectrum which suggests that ionization is being suppressed due to a faster excited state decay rate than for the bases. This observation recently motivated Tuna et al. to show that a charge transfer state with a conical intersection to the ground state exists in adenosine which can be reached as the distance between a proton on the ribose sugar and the N3 atom of the base reduces. This state was found to be present in both the syn and anti conformers and the conical intersections were significantly lower in energy than those intrinsic to the base. The only other theoretical work investigating electronic relaxation for a nucleoside was performed by Zgierski and Alavi who concluded for cytidine that the radiationless decay mechanisms were qualitatively similar to cytosine.

For DNA strands, the character of the excited state and hence its decay mechanisms depends strongly on the conformation of the helix. For tightly wound strands (A-type helix) where the bases are stacked on top of each other, the photo-excitation is de-localised between the two neighbouring bases which results in slower decay. If the bases are not well stacked (B-type helix), then the de-excitation is shorter and the decay mechanism is proposed to be similar to that of gas phase monomers. Therefore gas phase studies of DNA building blocks are important for understanding photo-protection mechanisms in long strands as well as providing benchmarks for the theoretical calculations which can be more easily compared with isolated molecules.

Given that studies of isolated nucleosides are a natural next step in bottom-up understanding of DNA photophysics, it is perhaps surprising that little theoretical work and no ultrafast pump–probe experiments have been undertaken to date. In the present study we have sought to remedy this situation by obtaining the first reported gas-phase measurements of ππ* excited state decay in all of the DNA nucleosides following UV excitation. The very low vapour pressures and strong propensity for thermal decomposition exhibited by these species at elevated temperatures has, up to now, made such an undertaking experimentally challenging. We have overcome these issues, however, through the use of a “soft” laser desorption method based on back-irradiation of thin foil targets coated with the molecular samples of interest. Similar approaches have previously been successfully applied to the study of dissociative electron attachment in thymidine.

**Experiment**

To produce gas-phase targets, nucleosides were desorbed directly from solid samples deposited on a 10 μm stainless steel foil which was integrated into the repeller electrode of a time-of-flight mass spectrometer. The samples (99% purity) were obtained from Sigma-Aldrich and used without further purification. The foil was irradiated with a 455 nm CW laser on the opposite side to that on which the samples were deposited to increase the temperature to ~150–250 °C. Unlike conventional molecular beams produced by ovens, evaporation using the present arrangement has the advantage that once desorbed the molecules do not undergo several cycles of condensation and re-evaporation, and they reach the interaction region quickly (in 10s of microseconds) due to the short travel distance from the foil (4–5 mm). This reduces the likelihood of thermal decomposition of the labile nucleoside species.

Femtosecond laser pulses used to excite and then subsequently ionize the nucleosides were obtained from a Coherent Libra system which produced 800 nm, 1 mJ pulses at a repetition rate of 1 kHz.
rate of 1 kHz. These were converted into the third harmonic (267 nm) using thin β-barium borate crystals as the non-linear medium. Residual fundamental pulses were delayed using a computer controlled translation stage and recombined with the 267 nm pulses to facilitate 267 nm + 800 nm pump-probe measurements. Both beams were focussed into the interaction region where they crossed the molecular plume at a distance of 10–15 mm from the focal position, with pulse energies of 8 µJ and 30 µJ and peak intensities of 2–5 × 10¹¹ W cm⁻² and 1–2 × 10¹³ W cm⁻², respectively. Ionization potentials (IPs) of the nucleosides under consideration here have not, to the best of our knowledge, been previously reported. However, IPs for all four species have been estimated from photoelectron studies performed on a series of methyl-substituted derivatives.⁴５ On the basis of this work it seems reasonable to assume that a minimum of three 800 nm photons are required to facilitate ionization from all of the excited bases. The pulse length for the 800 nm probe was measured by autocorrelation to be 130 fs, while ionization of Xe gas yielded a cross correlation of 190 fs which was consistent with the 267 nm probe also having a pulse length of 130 fs. For the pump-probe measurements, time of flight spectra were acquired for about half a minute at each time delay which was incremented in 27 fs steps between −500 fs and +2500 fs. A Xe cross correlation was performed before each acquisition to obtain an accurate zero delay position.

Results and discussion

We have previously reported mass spectra for each of the nucleosides using one-colour (1+1) resonance-enhanced multiphoton ionization (REMPI) at 267 nm with low interaction intensity (see ESI in ref. 43). These show that parent molecular ions make up more than 90% of the ion yield for thymidine and adenosine. This can be regarded as a lower limit on the proportion of intact molecules in the sample plume desorbed from the foil surface since at the pump intensities employed (5 × 10¹¹ W cm⁻²) there was a reasonable probability of the intact cation absorbing additional photons leading to dissociation. For guanosine and cytidine, while a strong parent peak was still observed in the spectrum, the largest peak corresponds to base ions. For cytidine this peak is nearly all due to protonated base ions which cannot originate from prior decomposition of the nucleoside into the nucleobase. For guanosine, about 40% of the ions in the mass spectrum were due to the parent, so we cannot rule out some decomposition prior to the laser interaction. However, this reduced parent yield may be a consequence of greater vibrational energy being deposited in the cation due to its lower ionization potential (8.0 eV⁴⁵). Therefore we are confident that the molecular plumes generated in our experiments are dominated by the intact, isolated nucleosides.

Mass spectra for the nucleosides obtained with the UV pump and IR probe pulses spatially and temporally overlapped are shown in Fig. 1 and have considerably more fragment ions than UV only REMPI spectra. We have attributed all significant peaks in the mass spectra to nucleoside fragments which agree with assignments found in other studies (for example NIST electron impact mass spectra⁴⁶). It is interesting to note that while the resonant excitation selectively ionizes an electron from the base, the presence of the intact sugar group ions (or fragments thereof) indicates that charge has been transferred from base to

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**Fig. 1** Mass spectra for the nucleosides integrated for delay times close to 0 fs. (a) Adenosine at 267 nm intensity 2 × 10¹¹ W cm⁻², 800 nm intensity 5 × 10¹³ W cm⁻²; (b) guanosine at UV = 3 × 10¹¹ W cm⁻², IR = 2 × 10¹³ W cm⁻²; (c) cytidine at UV = 3 × 10¹¹ W cm⁻², IR = 2 × 10¹³ W cm⁻²; (d) thymidine at UV = 6 × 10¹¹ W cm⁻², IR = 6 × 10¹² W cm⁻². M – molecular ion, B – base radical cation, B + H – protonated base, S₀ – intact sugar ion, Sₙ – ions corresponding to base + part of the sugar (numbering defined in ref 43), s – sugar fragment, b – base fragment.
sugar prior to dissociation. This is particularly noticeable for thymidine which is the only DNA nucleoside where the binding energy of the highest occupied molecular orbital in the base is greater than in the sugar,\textsuperscript{47} making the charge transfer energetically favourable. This subject has been discussed more fully by Poullay\textit{ et al.}\textsuperscript{42}

The variation in the thymidine parent ion yield and separately the sum of all the fragments is shown in Fig. 2 as a function of pump–probe delay time. It is evident that the excited state dynamics can be clearly observed through enhancement in the fragment ion yield for delays slightly greater than zero. As our analysis showed no difference in the dynamics between different fragments within the uncertainties, the yield from all fragments was summed to improve the statistics. Only a small proportion of this increase in fragment ions originates from loss in the parent ion, which is fairly insensitive to the pump–probe delay. Similar behaviour was found for the other nucleosides. The major contribution to the parent ions in the mass spectra arises from resonantly enhanced one-colour 1+1 ionisation by the pump pulse only. The lack of dynamics in the parent ion yield can be attributed to the high probability of dissociative ionisation out of the excited state by the moderately intense probe pulse so that signal enhancement is only observed for fragment ions. As it was not possible to guarantee a constant target density during the acquisition due to slow depletion of the sample on the foil, we have analysed the excited state dynamics by normalising the yield of all fragments to the parent ion yield. In this way we have accounted for the variations in target density without influencing the underlying dynamics. These results are presented for all the nucleosides in Fig. 3.

As noted by Kang\textit{ et al.},\textsuperscript{6} following excitation by the pump, the probe pulses can resonantly populate higher lying states when the two pulses are temporally overlapped – introducing a zero delay enhancement to the signal. Therefore, they fitted their results for gas-phase bases with the sum of a Gaussian function centred at zero pump–probe delay and a single exponential convolved by the cross-correlation of the laser pulses. For our data we use a similar approach but include a second exponential function as our better time resolution allows us to observe the fast decay component previously identified by Canuel \textit{et al.}\textsuperscript{7} To fully optimise the fitting process, we also introduced a third very long-lived component which arises due to parent ions generated directly by the pump being subsequently broken into fragment ions by absorption of probe photons.

The two lifetimes obtained for each nucleoside are presented in Table 1 along with the results of previous measurements performed at similar pump wavelengths on both the isolated bases (in the gas phase) and the nucleosides (in solution). The short lifetime components (described by a time-constant $\tau_1$) are consistent with the values obtained for the isolated bases by Canuel \textit{et al.}\textsuperscript{7} within the present uncertainties. We therefore attribute this extremely rapid decay ($\tau_1 = 50\text{–}230$ fs) to wavepacket motion away from the Franck–Condon region of the initially prepared $\pi \pi^*$ potential energy surface following excitation, which may also include internal conversion to an $\pi^* \pi^*$ state. The similarity of the base and nucleoside $\tau_1$ lifetimes suggests that there is little difference in their excited state potential energy surfaces near the Franck–Condon region.

In contrast, the second, slightly longer-lived component (described by a time-constant $\tau_2$) is around a factor of two shorter for adenosine, thymidine and cytidine than in the corresponding free base. This decay is a consequence of relaxation back to the ground state. In the case of guanosine, the situation appears somewhat different as we obtain a $\tau_2$ lifetime which is longer than for guanine. We note, however, that the lifetime previously reported for guanine (360 fs) is rather uncertain as later analysis including all fragment ions suggested that this was actually much longer (2300 fs).\textsuperscript{48} Theoretical calculations and simulated trajectories for guanine show that fast decay to the ground state (220 fs) is indeed present in the biologically relevant tautomer and proceeds directly from the initially populated $\pi \pi^*$ state without the involvement of an $\pi^* \pi^*$ state.\textsuperscript{12} The longer measured lifetime may be due to tautomers expected to be present in the experimental target. While the number of base tautomers present is restricted since the sugar is bound to the N9 atom, there are still at least three possible tautomers of the nucleoside.\textsuperscript{49} Without a means of pre-selecting different conformers and the fact that the present guanosine measurements have relatively poor statistics, it is not possible to determine if different tautomers are influencing the measured time constants (230, 2300 fs).

So and Alavi have previously reported excited state energies for adenosine, guanosine, cytidine and uridine (structurally very similar to thymidine) using density functional theory (B3LYP) in conjunction with the 311++G(d,p) and aug-cc-pVDZ basis sets.\textsuperscript{50} For comparative purposes, equivalent calculations were also reported for the isolated bases. In all cases, the addition of the sugar group appeared to have only a small influence on the energies of the lowest-lying singlet $\pi \pi^*$ and $\pi^* \pi^*$ states (the former of which is optically bright and is the principle state directly excited at 267 nm) – both in relative and absolute terms. Critically within the context of this current work, no clear systematic shifts in energy between these states were seen across the various
nucleoside systems. This is perhaps unsurprising as the lowest-
lying $\pi\pi^*$ and $n\pi^*$ states are predominantly localized on the
aromatic pyrimidine/purine ring systems in all cases. So and
Alavi\textsuperscript{50} also revealed states of $\pi\sigma^*$ character present in the low
energy excited state manifold in the isolated bases are no
longer seen in this energy region in the corresponding nucleo-
sides (with the exception of guanosine). Such states have
previously been implicated as playing a role in the dynamical
relaxation of the bases following UV excitation. As discussed
previously in the Introduction, the importance of this pathway
following excitation in the 267 nm region is not fully resolved,
although it is generally thought to be relatively minor.

For adenosine, thymidine and cytidine, the calculations
outlined above clearly suggest that the significantly shorter
overall excited state lifetimes observed (compared to the iso-
lated bases) cannot be rationalized simply in terms of changes

![Table 1](image)

<table>
<thead>
<tr>
<th>(a) $\tau_1$</th>
<th>A</th>
<th>G</th>
<th>C</th>
<th>T</th>
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</thead>
<tbody>
<tr>
<td>Base in gas-phase [7]</td>
<td>100</td>
<td>148</td>
<td>160</td>
<td>105</td>
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<tr>
<td>Nucleoside in gas-phase (present)</td>
<td>$50 \pm 90^{20}$</td>
<td>$230 \pm 80^{150}$</td>
<td>$120 \pm 40^{30}$</td>
<td>$100 \pm 10$</td>
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</table>

<table>
<thead>
<tr>
<th>(b) $\tau_2$</th>
<th>A</th>
<th>G</th>
<th>C</th>
<th>T</th>
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</thead>
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<tr>
<td>Nucleoside in solution [30]</td>
<td>530</td>
<td>690</td>
<td>760</td>
<td>700</td>
</tr>
<tr>
<td>Nucleoside in solution [31]</td>
<td>290</td>
<td>540</td>
<td>460</td>
<td>720</td>
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<tr>
<td>Nucleoside in gas phase (present)</td>
<td>$570 \pm 60$</td>
<td>$2300 \pm 3300^{900}$</td>
<td>$740 \pm 270^{160}$</td>
<td>$2900 \pm 700^{500}$</td>
</tr>
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in the relative excited state energy orderings and/or spacings. This would therefore imply that there is an additional decay mechanism operating in the nucleosides which is not present in the bases. To date there have been limited theoretical investigations of the decay mechanisms for isolated nucleosides. Calculations of the decay pathway in cytidine suggest that transition to the ground state occurs via a biradical state which is qualitatively similar to the cytosine base.\(^{40}\) In contrast, Tuna et al.\(^{39}\) calculate the presence of a charge transfer state in two separate conformers of adenosine which intersect with the ground state as a proton is transferred from the sugar to base via an intramolecular H-bond (5'-OH...N3 in the syn conformer and 2'-OH...N3 in the anti conformer). As these conical intersections are about 0.6 eV lower than the conical intersections corresponding to ring puckering in the base, this could lead to faster de-excitation. It is interesting to note that we do not observe a shorter lifetime in guanosine despite the similarity of its structure to adenosine. However, our measured lifetimes are consistent with previous qualitative observations that guanosine and adenine parent ions can be produced in resonant two photon spectroscopy experiments but not adenosine.\(^{37,38}\)

There is currently no theory supporting the existence of charge transfer states in cytidine and thymidine, but the influence of intra-molecular H-bonding between the base and sugar can be expected – as revealed in the DFT optimized ground state geometries previously reported by So and Alavi, where all nucleoside systems exhibit similar base-sugar interactions.\(^{50}\) It therefore seems reasonable to assume (as a starting point for future discussion) that this charge transfer state may potentially also be present in all cases and actively participate in the relaxation dynamics. Finally, it is interesting to note that the lifetimes for bases and nucleosides in solution are considerably shorter than for gas phase bases but in good agreement with our values for adenosine and cytidine. Similarly very short lifetimes can be found in H-bonded base pairs. These observations indicate that population of charge transfer states which stimulate proton migration to the base from the sugar, solvent or another base accelerates the de-excitation process.

**Conclusion**

The use of “soft” laser desorption methods have permitted the first reported dynamical study of excited state excess energy redistribution in gas-phase DNA nucleosides following UV irradiation. Time-resolved ion-yield measurements reveal that the lifetimes for internal conversion to the ground state following excitation are reduced by around a factor of two for adenosine, cytidine and thymidine when compared with the corresponding isolated bases. This is not the case for guanosine, which may be a consequence of several conformers/tautomers being present in the desorbed sample. Results are discussed in terms of a recent proposition that a charge transfer state in adenosine provides an additional internal conversion pathway mediated by proton transfer through a sugar to base hydrogen bond. Previously reported ground state geometry calculations for other DNA nucleosides appear to also suggest that this mechanism may operate more generally. However, more comprehensive theory looking at an expanded range of excited state properties in these systems would be highly desirable. This is in addition to more detailed gas-phase experimental studies using an expanded range of UV excitation wavelengths and/or exploiting more differential measurements (for example, time-resolved photoelectron imaging). We hope and anticipate that this communication will serve as a stimulus for such undertakings in the near future.

**Acknowledgements**

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