Ion Mobility Unlocks the Photofragmentation Mechanism of Retinal Protonated Schiff Base

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Abstract

Retinal protonated Schiff base (RPSB) is a key molecular component of biological photoreceptors and bacterial photosynthetic structures, where its action involves photoisomerization around bonds in the polyene chain. In a vacuum environment, collisional activation or exposure to visible light causes the RPSB molecule to disintegrate, producing charged molecular fragments with \( m/z = 248 \) Da, that cannot be formed by simple cleavage of the polyene chain. Photofragments resulting from laser excitation of RPSB at a wavelength of 532 nm are analysed in an ion mobility mass spectrometer (IMMS) and found to be the protonated Schiff base of \( \beta \)-ionone. Density functional theory calculations at the M062X/cc-pVDZ level support a fragmentation mechanism in which RPSB undergoes an electrocyclization/fragmentation cascade with the production of protonated Schiff base of \( \beta \)-ionone and toluene.

Keywords: retinal, photodissociation, electrocyclization, ion mobility, mass spectrometry
TOC Graphic: Retinal protonated Schiff base is photodissociated in an ion mobility mass spectrometer. The drift mobility of the 248 Da fragments unambiguously identifies them as isomers of β-ionone protonated Schiff base resulting from electrocyclization of the polyene chain.
Retinal protonated Schiff base (RPSB - Figure 1.1A) is a key molecule in the visual transduction cycle and in bacterial photosynthesis, where, in both cases, its action involves photoisomerization around bonds in the polyene chain. When embedded in a host protein, RPSB is remarkably photostable and can be repeatedly photo-excited without decomposing. However, in a vacuum environment RPSB is more fragile, and exposure to visible light or collisions with neutral gas molecules, causes its disintegration.\textsuperscript{1–5} Intriguingly, the dominant charged fragment at $m/z=248$ Da cannot be generated through a simple chain cleavage and must result from a process involving cyclization of the polyene chain.\textsuperscript{5} The previously proposed decomposition mechanism,\textsuperscript{5} illustrated in Figure 1.1, involves a sub-millisecond rearrangement through a Diels-Alder mechanism giving an energized, long-lived tricyclic structure, that disintegrates on a millisecond timescale to produce a charged bicyclic 248 Da fragment and a toluene molecule (or an isomer thereof).

Here we consider an alternative mechanism for production of the 248 Da fragment from RPSB. As shown in Figure 1.2, a sequential electrocyclization/fragmentation process leads directly to $\beta$-ionone protonated Schiff base (Figure 1.2D) and a neutral toluene molecule. This process is analogous to the elimination of toluene and xylene from carotenes observed in mass spectrometric studies.\textsuperscript{6,7} Density functional theory (DFT) calculations at the M062X/cc-pVDZ level suggest that the fragment energies for the two mechanisms are comparable (–58 kJ/mol for 1D and –70 kJ/mol for 2E with respect to the energy of RPSB).

To examine the RPSB fragmentation process we use ion mobility mass spectrometry (IMMS) to probe the 248 Da photofragments, which have different structures (and collision cross sections) for the two mechanisms. IMMS is a versatile tool for exploring conformations of charged molecules, relying on the sensitivity of a molecule’s collision cross section to its structure; under the influence of an electric field, compact molecules travel more swiftly through a buffer gas than extended, unfolded molecules.\textsuperscript{8–13} IMMS has been employed to elucidate the structures of a broad range of charged molecules and clusters including carbon and silicon clusters,\textsuperscript{14,15} carbohydrates,\textsuperscript{16} peptides,\textsuperscript{17,18} and proteins.\textsuperscript{10} Recently we showed
Figure 1: RPSB fragmentation mechanisms proposed by Toker et al. (1),\(^5\) and in this work (2). In mechanism 1, RPSB undergoes a [4+2] cycloaddition to form a tricyclic intermediate structure, followed by loss of toluene. In mechanism 2, sequential 8\(\pi\)/6\(\pi\) electron electrocyclizations lead to an intermediate containing a four membered ring, followed by elimination of toluene through a 4\(\pi\) cycloreversion to give the protonated Schiff base of \(\beta\)-ionone. The two mechanisms involve loss of different sections of the polyene chain.
that it can also be used to monitor the photoisomerization of charged molecules in the
gas phase including carbocyanine dyes and RPSB.\textsuperscript{19–21} Here we demonstrate that IMMS
can also be used to explore the structures of molecular photofragments and shed light on
photofragmentation mechanisms.

In our experiment, RPSB ions produced by electrospray ionization are irradiated in the
drift region of a purpose-built ion mobility spectrometer in which ions are propelled through
N\textsubscript{2} buffer gas by a modest electric field (Figure 2).\textsuperscript{19–21} This arrangement enables us to
discriminate charged photofragments on the basis of both mass and collision cross section,
providing information on their geometrical structures. If the mechanism proposed in ref.
5 prevails, one would expect to observe a relatively mobile 248 Da bicyclic photofragment
(1D in Figure 1), and perhaps a stabilized version of the 340 Da intermediate (1C). On
the other hand, the sequential electrocyclization/fragmentation reaction should yield slower
monocyclic \(\beta\)-ionone PSB photofragments (2E in Figure 1) and possibly the stabilized inter-
mediates 2C and 2D.
Figure 3: Laser-off ion count plotted against mass (m/z) and cross section for collisions with N₂ (σ). Red circles represent calculated data for isomers derived from the mechanism proposed in this work, whereas red squares correspond to isomers associated with the mechanism proposed by Toker et al.⁵ The ion count inside the dashed rectangle has been multiplied by a factor of 10 for clarity. For expanded views of this plot, see the SI.

The ion count for electrosprayed RPSB solution is plotted as a function of mass and collision cross section with N₂ in Figure 3. The ion population is dominated by 340 Da, σ=227 Å² ions associated with all-trans RPSB, but also includes 248 Da ions, as previously identified by Toker et al.⁵ These ions are produced in the source region of our apparatus (presumably through decomposition of RPSB ions in the electrospray desolvation capillary or first ion funnel). The ion mobility spectrum for the 248 Da ions (obtained with the mass filter tuned to 248 Da) is given in Figure 4a, clearly showing that there are in fact three 248 Da isomers with collision cross sections of 179.7, 176.8 and 167.5 Å², respectively. Measured collision cross sections for the dominant 248 Da ions (179.7, 176.8 Å²) are closer to the predicted cross section for the monocyclic β-ionone PSB (2E; σ_{calc}=173.9 Å² for the all-trans isomer) than the more compact, bicyclic structure proposed by Toker et al.⁵ (1D; σ_{calc}=166.3 Å²).

Notably, we find no evidence for the stabilized tricyclic intermediate (1C), which is pre-
dicted to have a collision cross section 15% less than RPSB, and which, if it is formed rapidly and is long-lived, should be stabilized through collisions in our IMMS apparatus. Evidence for the formation of 2C and 2D is equivocal; predicted cross sections for the various stereoisomers of 2C and 2D (see SI) are close to observed signal although there is no exact correspondence.

As demonstrated previously, when the drifting RPSB ions in our apparatus are exposed to light at low intensity (λ=532 nm, I=1-2 mJ/cm²/pulse) photodissociation is minimal and instead RPSB photoisomerizes from the all-trans form to various cis forms. However, at much higher intensities (I≈30 mJ/cm²/pulse) we find that 248 Da photofragments are generated from RPSB. These photofragments have a very similar ion mobility spectrum to that of the 248 Da fragments formed in the ion source (Figure 4b).

To confirm the identity of the 248 Da fragment we synthesized β-ionone PSB (Figure 1, 2E) and electrosprayed it directly into the IMMS apparatus. As shown in Figure 4c, the ion mobility spectrum for the electrosprayed sample corresponds unambiguously with the ion mobility spectra for the 248 Da fragments generated from RPSB in the ion source and through photodissociation from RPSB, conclusively identifying these fragments as β-ionone PSB. The three IMS peaks, which have slightly different relative intensities in Figure 4(a-c), are presumably associated with β-ionone geometric isomers with different numbers of cis bonds along the polyene chain (calculated energies and predicted collision cross sections for β-ionone PSB isomers are given in the SI). At this stage, we assign the slowest peak to the all-trans isomer, the next fastest peak to molecules with a single cis linkage and the fastest peak to isomers with a double cis linkage.

The consecutive 8πe and 6πe electrocyclizations of retinal through mechanism 2 (Figure 1) are reminiscent of processes occurring in synthetic organic chemistry and in natural systems. To support the proposed rearrangement we used DFT M06-2X/cc-pVDZ calculations to investigate a simplified RPSB model that mimics the C9-N polyene chain, replacing the β-ionone and N-butyl substituents by vinyl and methyl groups, respectively (Figure 5). The
Figure 4: Ion mobility spectra for 248 Da ions formed from: (a) activation of RPSB in the ion source; (b) photodissociation of RPSB in the drift tube; (c) electrosprayed protonated Schiff base of β-ionone. In each case the peaks are fitted to 3 Gaussian functions constrained to have fwhm=3 Å². Calculated collision cross sections for isomers 1D and 2E (β-ionone) with N₂ are shown in (a).
process involves several trans-cis isomerizations to achieve the required “curled” precursor conformation for the $8\pi$e cyclization (1). The reaction then proceeds in a stepwise manner, whereby the first ring closure, which leads to an 8-membered ring (structure 2 in Figure 5) is associated with a barrier of 58 kJ/mol (TS1). The second electrocyclization has an activation energy of 77 kJ/mol (TS2) and leads stereospecifically to a cis-bicyclo[4.2.0]octane framework (structure 3 in Figure 5). This cycloadduct is lower in energy by 34 kJ/mol compared to starting material 1. Subsequent dissociation through cycloreversion via TS3 occurs through a concerted bond cleavage (as confirmed by Intrinsic Reaction Coordinate calculations) and produces toluene and a protonated Schiff base. Overall, the electrocyclization/fragmentation cascade is energetically highly favourable with the product association complex 4 being about 157 kJ/mol lower in energy than the starting material 1. Mechanism 2 is possible both thermally (as shown in Figure 5) and photochemically. The only possible difference lies in the stereoselectivity of the cyclization and fragmentation steps. However, as outlined below, we believe that photoexcited RPSB rapidly internally converts from the
S₁ to the S₀ state so that the photo-induced electrocyclic reaction observed under our experimental conditions occurs for highly vibrationally excited ions on the ground state PES. Under these circumstances, the stereochemical progress of the 8πe and 6πe electrocyclizations should follow courses appropriate for thermal conditions (conrotatory for the 8πe cyclization and disrotatory for the 6πe cyclization). Although, the sequential electrocyclizations should be highly stereospecific processes, apart from the cis fusion of the two rings, it is not possible to predict the relative stereochemistry at the cyclobutane ring for the full 2D (Figure 1), since the outcome of the electrocyclizations depends also on the geometry of the C9-C10 and C=N double bond in RPSB. As explained above, under our experimental conditions, we cannot unequivocally determine whether 2C and 2D are sufficiently long-lived to enable their detection, and, therefore, detailed knowledge of their stereochemistry is not essential at this stage.

It should also be noted that DFT M06-2X/cc-pVDZ computations for a simplified model system for 1C did not reveal a fragmentation pathway leading to 1D. Instead, the calculations suggest that heterolytic cleavage of the C-NR₃⁺ bond in 1C is a more viable process.

The dominance of RPSB photoisomerization over photodissociation at low light levels causes us to believe that in our apparatus photodissociation of RPSB involves consecutive absorption of two or more 532 nm photons.²¹ Possibly, the process occurs through absorption of one photon followed by rapid internal conversion, absorption of a second photon, again followed by internal conversion, trans-cis isomerizations, 8πe electrocyclization, 6πe electrocyclization, and dissociation, all on the ground state potential energy surface. In contrast, RPSB dissociation in the ion storage ring is believed to involve absorption of a single photon.⁵ The magnitude of the calculated energy barriers for the simplified model suggests that single photon photodissociation of RPSB should be possible over the 500-610 nm range, although the process may be sufficiently slow that in our experiment the reactant RPSB ions are collisionally quenched in the drift tube, where the collision rate is ≈5×10⁸ s⁻¹, before surmounting the initial barrier to cyclization. On the other hand, in the ion ring experiment,
where the pressure is $\leq 10^{-10}$ torr and the collision rate is much lower. RPSB ions that have absorbed a single photon have much more time to rearrange, and are able to dissociate on a millisecond timescale.

In summary, we have used IMMS to probe the structure of charged molecular fragments from the chromophore RPSB in the gas phase and have unequivocally identified the 248 Da fragment as the protonated Schiff base of $\beta$-ionone. The proposed mechanism, which occurs after both thermal and photo-excitation, involves isomerization of the polyene chain, followed by a sequential electrocyclization/fragmentation cascade that leads to elimination of toluene. Eventually, the IMMS approach described in this paper should be applicable to structural and mechanistic investigations of a broad range of gas-phase photochemical reactions, augmenting more cumbersome and expensive methods including isotopic labelling.

**Experimental Methods**

The ion mobility apparatus is the same as used recently to investigate the photoisomerization of carbocyanine dyes and RPSB cations. In the current study, electrosprayed RPSB cations produced from a $10^{-5}$ M solution of trans RPSB in 1:1 methanol:H$_2$O (electrospray voltage 3 kV, flow rate 5 $\mu$L/min), were accumulated in an ion funnel before being launched in a 400 $\mu$s pulse into a 0.9 m drift tube containing N$_2$ buffer gas ($P=14$ torr). The electric field in the drift tube (44 V/cm) was sustained by 90 ring electrodes. At the end of the drift tube, the ions were collected radially by a second ion funnel before passing through a 0.3 mm orifice into an octopole ion guide from which they exited through a 3 mm orifice into a quadrupole mass filter. Ions were sensed by a channeltron detector connected to a discriminator and a multichannel scaler.

The apparatus was run at 20 Hz, with alternate ion packets exposed to the 532 nm output from a pulsed frequency-doubled Q-switched Nd:YAG laser. The ions’ arrival time distribution (ATD) was built up as a histogram of ion counts versus time. The mobility
resolution of the instrument was typically $t_d/\Delta t_d=60$, which can be compared to a maximum, diffusion-limited resolution under the prevailing conditions of 120.$^{19}$ Under typical operating conditions with a drift field of 44 V/cm and N$_2$ buffer gas pressure of 14 torr, the effective temperature of the drifting RPSB ions is predicted to be $\approx 300$ K.$^{24}$

**Computational Methods**

To connect the measured collision cross sections with molecular structures, we determined equilibrium geometries for different 248 and 340 Da isomers using DFT calculations at the M062X/cc-pVDZ level.$^{25}$ Collision cross sections were determined from the calculated equilibrium structures using the trajectory method as instituted in the MOBCAL program.$^{26,27}$ Atom-atom potential energy parameters for the interaction between N$_2$ and the colliding molecule were taken from ref. 12. Calculated energies and collision cross sections for isomers of 1C, 1D, 2C, 2D and 2E are given in the SI.

**Acknowledgments**

This research was supported under the Australian Research Council’s Discovery Project funding scheme (Project Numbers DP110100312 and DP120100100). The computations were supported by the National Computational Merit Allocation Scheme (Project m88). We thank Professor Matthew Bush for providing a modified version of the Mobcal code with N$_2$ collision parameters described in ref. 12, Luke Gamon for assistance with synthesis and collection of GC-MS data for the protonated Schiff base of $\beta$-ionone, and Sioe See Volaric for assistance with collection of LC-MS data.
Supporting Information Available

Further experimental and computational information, including details for the synthesis of RPSB and β-ionone PSB, and calculated energies and collision cross sections for intermediates and products shown in Figure 1, is available as SI. This material is available free of charge via the Internet at http://pubs.acs.org/.

References


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Title:
Ion Mobility Unlocks the Photofragmentation Mechanism of Retinal Protonated Schiff Base

Date:
2014-09-18

Citation:
Coughlan, NJA; Adamson, BD; Catani, KJ; Wille, U; Bieske, EJ, Ion Mobility Unlocks the Photofragmentation Mechanism of Retinal Protonated Schiff Base, JOURNAL OF PHYSICAL CHEMISTRY LETTERS, 2014, 5 (18), pp. 3195 - 3199

Persistent Link:
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File Description:
Accepted version