Characterization of O_2 ($^1\Delta_g$)-Derived Oxidation Products of Tryptophan: A Combination of Tandem Mass Spectrometry Analyses and Isotopic Labeling Studies

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The fragmentation mechanisms of singlet oxygen $[O_2 ({}^{L}\Delta_g)]$ -derived oxidation products of tryptophan (W) were analyzed using collision-induced dissociation coupled with ¹⁸O-isotopic labeling experiments and accurate mass measurements. The five identified oxidized products, namely two isomeric alcohols (*trans* and *cis* WOH), two isomeric hydroperoxides (*trans* and *cis* WOOH), and N-formylkynurenine (FMK), were shown to share some common fragment ions and losses of small neutral molecules. Conversely, each oxidation product has its own fragmentation mechanism and intermediates, which were confirmed by ¹⁸O-labeling studies. Isomeric WOH lost mainly $H_2O + CO$, while WOOH showed preferential elimination of $C_2H_5NO_3$ by two distinct mechanisms. Differences in the spatial arrangement of the two isomeric WOHs led to differences in the intensities of the fragment ions. The same behavior was also found for *trans* and *cis* WOOH. FMK was shown to dissociate by a diverse range of mechanisms, with the loss of ammonia the most favored route. MS/MS analyses, ¹⁸O-labeling, and $H_2^{18}O$ experiments demonstrated the ability of FMK to exchange its oxygen atoms with water. Moreover, this approach also revealed that the carbonyl group has more pronounced oxygen exchange ability compared with the formyl group. The understanding of fragmentation mechanisms involved in O₂ ($^{1}\Delta_{o}$)-mediated oxidation of W provides a useful step toward the structural characterization of oxidized peptides and proteins. (J Am Soc Mass Spectrom 2009, 20, 188-197) © 2009 Published by Elsevier Inc. on behalf of American Society for Mass Spectrometry

Reactive oxygen species are believed to be formed within cells during aerobic metabolism. Many lines of evidence have linked the oxidative damage generated by these species with the development of pathological conditions, such as atherosclerosis, neurodegenerative diseases, and cancer [1]. Singlet molecular oxygen $[O_2 ({}^{1}\Delta_g)]$ is a reactive form of oxygen that can be produced by the action of peroxidases [2, 3], UVA light in the presence of a sensitizer [4, 5] and also during the dimerization reactions of peroxyl radicals in lipid peroxidation [6, 7]. Electron-rich biomolecules, such as lipids, proteins, and DNA, can be important targets for $O_2 ({}^{1}\Delta_e)$ oxidation [8].

Mass spectrometry (MS) has been applied to the identification of oxidized products resulting from the reaction of reactive oxygen species with a variety of biomolecules. For instance, this technique has been employed to analyze oxidized lipids derived from the Fenton reaction, such as oxidation products of linoleic acid [9], phospholipids [10, 11], and also cholesterol ozonolysis derived products [12]. In DNA, a representative example is O_2 ($^1\Delta_g$) oxidation of 2'-deoxyguanosine, which has been shown to yield 8-oxo-7,8-dihydro-2'-deoxyguanosine in double stranded DNA and also other oxidized products upon reaction with an isolated base [13, 14].

Detection of oxidized amino acid residues in proteins is a growing field of interest, since accumulation of oxidized proteins in vivo has been related to aging and age-related diseases [15]. Among the common amino acids, tryptophan (W) is considered a highly susceptible residue to oxidizing agents [16, 17]. Many oxidation products of W have been studied by MS. These products include oxindolylalanine (Oia) and dioxindolylalanine [18], generated by the treatment of W with mixtures of dimethyl sulfoxide and hydrochloric acid in acetic acid and monitored by fast atom bombardment mass spectrometry. Oxindolylalanine and *N*-acetyloxindolylalanine were also studied by tandem mass spectrometry in a quadrupole ion trap mass spectrometer. These molecules were shown to undergo

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a range of fragmentation reactions, including the loss of water via a number of distinct pathways [19]. Furthermore, hydroxytryptophan, dihydroxytryptophan and/ or *N*-formylkynurenine (FMK) and W-W dimers generated by the Fenton reaction with W were detected by the combination of spin trapping and electrospray mass spectrometry techniques [20]. Electrospray mass spectrometry analyses were also used to propose the formation of FMK as an ozonation product of W [21]. Indeed, the fragmentation pathways of kynurenine (kn) were evaluated using this methodology and deuterium exchange experiments [22].

However, other important oxidation products of W remain unexplored. This is the case for the products generated by the action of O_2 (${}^{1}\Delta_g$) and other oxidizing conditions, such as the addition of a hydroxyl radical to the pyrrole moiety, followed by loss of water and addition of superoxide radical or molecular oxygen [23, 24]. In a previous work, we have shown that tryptophan oxidation by O_2 (${}^{1}\Delta_g$) gives rise to *cis*- and *trans*-tryptophan hydroperoxides (WOOH), which can further decompose into the corresponding isomeric alcohols (WOH) and FMK (Scheme 1) [25]. In this study, we explore in depth the use of product and precursor ion spectra, associated with 18 O-labeling experiments, to derive structural information regarding these oxidation products of W.

Experimental

Materials

L-Tryptophan (W), deuterium oxide (D_2O), $H_2^{-18}O$ (96.9%), formic acid, and rose bengal were supplied by Sigma Aldrich (St. Louis, MO). The $^{18}O_2$ gas cylinder came from Isotec-Sigma (St. Louis, MO). Acetonitrile (HPLC grade) was purchased from Merck (Rio de Janeiro, Brazil). The water used in the experiments was treated with the Nanopure Water System (Barnstead, Dubuque, IA).

Synthesis of ¹⁶O- or ¹⁸O-Tryptophan *Photoproducts*

L-Tryptophan (20 mM) was dissolved in D_2O containing 10 μ M rose bengal and irradiated using light from a



Scheme 1. O₂ ($^{1}\Delta_{g}$)-derived oxidation products of tryptophan.

tungsten lamp (500 watt) filtered through a 360 nm cutoff filter. Irradiation was applied for 3 h in an ice bath under a continuous flux of oxygen. For the photo-oxidation in an ¹⁸O₂-saturated atmosphere, oxygen contained in the system was removed by successive freezing and thawing under vacuum. This procedure was repeated at least five times to ensure complete removal of ¹⁶O₂. Thereafter, the whole system was connected to an ¹⁸O₂ gas cylinder under 0.5 atm. After the irradiation, rose bengal was removed by passage through an Acrodisc syringe filter (Pall Gelman, Ann Arbor, MI) with a 0.2 μ m Supor membrane.

Purification of Photoproducts by HPLC

The reaction products were purified using a semipreparative C18 reverse-phase column ($250 \times 10 \text{ mm}$, particle size 10 μ m, Phenomenex, Torrance, CA) connected to a Shimadzu HPLC system (Shimadzu, Tokyo, Japan). The separation was carried out at a flow rate of 4.6 mL/min with a gradient of water–acetonitrile, using isocratic elution with 2% acetonitrile for 8 min. Then a linear gradient from 2% to 6% in 1 min was employed, maintaining 6% up to 15 min and reaching 40% in the next 3 min. After that, the column was washed with 40% acetonitrile for 3 min, returning to 2% in 3 min, followed by column stabilization for another 5 min at 2%. The product peak fractions were collected and immediately lyophilized.

HPLC/Mass Spectrometry Analysis

HPLC/MS analyses were carried out in a Shimadzu HPLC system (Tokyo, Japan) coupled to a Quattro II mass spectrometer (Micromass, Manchester, UK) with a Z-spray source. The separation of photoproducts was achieved using a 250 \times 4.60 mm (particle size 5 μ m) C18 reverse-phase column (Phenomenex). The mobile phase consisted of 4% acetonitrile in 0.05% formic acid at a flow rate of 1.0 mL/min. A small fraction of the eluent was directed into the mass spectrometer at a flow rate of 100 μ L/min. Mass spectrometry analyses were done with the electrospray ionizer in the positive ion mode. The source and desolvation temperatures were kept at 100 °C and 150 °C, respectively. The optimal flow rates of the drying and nebulizing gases were found to be 300 L/h and 15 L/h, respectively. The cone voltage was set at 10 V. Full-scan data were acquired over a mass range 100-500 m/z. Tandem mass spectrometry (MS/MS) experiments were performed for each molecular ion, using the collision energy at 10 eV. Also, precursor and product ion scans were conducted on the major ions observed in the MS/MS of unlabeled and ¹⁸O-labeled W oxidation products. For this purpose, the most abundant protonated fragments were generated in the ionization source by increasing the cone voltage to 30 V. After that, each of the generated ions was selected and fragmented again, with a collision energy ranging from

5 to 20 eV. Also, the product ions were selected in MS-1 and scanned to identify all of the precursor ions.

Accurate Mass Measurements

Accurate mass measurements were performed on a micrOTOF electrospray ionization time-of-flight (ESI-TOF) mass spectrometer (Bruker Daltonics, Billerica, MA). Purified ¹⁸O-labeled or unlabeled oxidized W derivatives were dissolved in a mixture of 0.01% formic acid:acetonitrile (1:1) and infused into the mass spectrometer at a flow rate of 180 μ L/h. The voltage of the capillary was 4 kV, the dry gas was kept at 5.0 l/min, and the source temperature was maintained at 160 °C. Calibration of the instrument was achieved through the use of sodium formate. Mass spectra were acquired over a *m*/z range 100–400.

Oxygen Exchange Experiments

Purified ¹⁶O-FMK was dissolved in 96.9% $H_2^{18}O$ and incubated at 37 °C with stirring for up to 4 h. This solution was then analyzed by MS/MS.

Results and Discussion

Photosensitization of W gives rise mainly to four product peaks with retention times of 5.03, 6.60, 7.92, and 10.28 min in the UV chromatogram, together with the unreacted W peak at 21.54 min (Figure 1). The ion chromatogram obtained by selecting the m/z 221 reveals that two products have insertions of one oxygen atom in the tryptophan moiety [W (m/z 205) + 16, Figure 1b], while selection of the m/z 237 shows three products with insertion of two oxygen atoms (W + 32, Figure 1c) in the parent molecule. These products were previously characterized as two isomeric alcohols (m/z 221), two isomeric hydroperoxides (m/z 237) and FMK (m/z 237). The *cis/trans* configurations of the alcohols and hydroperoxides were established regarding the relative position of the hydroxyl or hydroperoxyl and carboxylic acid groups [25].

Accurate mass measurements were carried out for each W-derived oxidation product synthesized with isotopes 16 and 18 of oxygen. The accurate mass was also determined for the main fragment ions obtained for each labeled or unlabeled photoproduct. The results are summarized in Table 1. In all cases, the measured masses of the molecular and fragment ions were within ± 5 ppm of those expected from the proposed composition.

The tandem mass spectrum obtained for the $[M + H]^+$ molecular ion at 5.03 min (*trans* alcohol) is shown in Figure 2a. The *cis* isomer exhibits almost the same profile (see discussion below). To understand the mechanisms of fragmentation, the two isomeric alcohols were also synthesized with isotope 18 of oxygen. Figure 2b shows the successful incorporation of one ¹⁸O into the *trans* alcohol, which then exhibited a molecular ion at *m*/*z* 223.

Indeed, successive fragment and precursor ion scans were conducted on the major ions observed in the MS/MS of unlabeled and ¹⁸O-labeled alcohols. For this purpose, protonated fragments were generated in the ionization source. After that, each of the ions was selected and fragmented again, with a collision energy ranging from 5 to 20 eV. Also, the product ions were selected in MS-1 and scanned to identify all of the precursor ions. The results of these experiments are illustrated in Scheme 2 for the ¹⁸O-labeled alcohols. Three fragmentation pathways can be proposed upon analysis of the product and precursor ion scans. The loss of small neutral molecules from the protonated molecular ion or fragments is a common event for all routes. Indeed, the three pathways share some intermediates and final products.

Elimination of H_2O and CO (m/z 177, for the labeled molecule) from the protonated alcohols was the major



Figure 1. HPLC-ESI/MS obtained after photosensitization of tryptophan. (a) Chromatogram with UV detection at 215 nm. (b) Ion chromatogram selecting m/z 221. (c) Ion chromatogram selecting m/z 237.

Product	Assignment	Suggested elemental composition	Observed <i>m/z</i>	Calculated <i>m/z</i>	Error (ppm)
trans WOH	$[M + H]^+$	C ₁₁ H ₁₃ N ₂ ¹⁶ O ₃	221.0924	221.0926	-1.1
	$[M - (CO, H_2O) + H]^+$	C ₁₀ H ₁₁ N ₂ ¹⁶ O	175.0867	175.0872	-2.6
<i>trans</i> W ¹⁸ OH	$[(M + 2) + H]^+$	C ₁₁ H ₁₃ N ₂ ¹⁶ O ₂ ¹⁸ O	223.0962	223.0969	-3.1
	$[(M + 2) - (CO, H_2O) + H]^+$	C ₁₀ H ₁₁ N ₂ ¹⁶ O	177.0906	177.0914	-4.5
cis WOH	$[M + H]^+$	C ₁₁ H ₁₃ N ₂ ¹⁶ O ₃	221.0933	221.0926	3.0
	$[M - (CO, H_2O) + H]^+$	C ₁₀ H ₁₁ N ₂ ¹⁶ O	175.0875	175.0872	2.0
<i>cis</i> W ¹⁸ OH	$[(M + 2) + H]^+$	C ₁₁ H ₁₃ N ₂ ¹⁶ O ₂ ¹⁸ O	223.0968	223.0969	-0.4
	$[(M + 2) - (CO, H_2O) + H]^+$	C ₁₀ H ₁₁ N ₂ ¹⁶ O	177.0911	177.0914	-1.6
trans WOOH	$[M + H]^+$	C ₁₁ H ₁₃ N ₂ ¹⁶ O ₄	237.0866	237.0876	-4.0
	$[M - C_2H_5NO_3 + H]^+$	C ₉ H ₈ N ¹⁶ O	146.0599	146.0606	-4.7
<i>trans</i> W ¹⁸ O ¹⁸ OH	$[(M + 4) H]^+$	C ₁₁ H ₁₃ N ₂ ¹⁶ O ₂ ¹⁸ O ₂	241.0953	241.0960	-3.1
	$[(M + 4) - C_2H_5NO_2^{18}O + H]^+$	C ₉ H ₈ N ¹⁸ O	148.0649	148.0648	0.5
<i>cis</i> WOOH	$[M + H]^+$	C ₁₁ H ₁₃ N ₂ ¹⁶ O ₄	237.0866	237.0876	-4.0
	$[M - C_2H_5NO_3 + H]^+$	C ₉ H ₈ N ¹⁶ O	146.0599	146.0606	-4.7
<i>cis</i> W ¹⁸ O ¹⁸ OH	[(M + 4) H] ⁺	C ₁₁ H ₁₃ N ₂ ¹⁶ O ₂ ¹⁸ O ₂	241.0958	241.0960	-1.0
	$[(M + 4) - C_2H_5NO_2^{18}O + H]^+$	C ₉ H ₈ N¹ ⁸ O	148.0647	148.0648	-0.9
FMK	$[M + H]^+$	C ₁₁ H ₁₃ N ₂ ¹⁶ O ₄	237.0872	237.0876	-1.5
	$[M - NH_3 + H]^+$	C ₁₁ H ₁₀ N ¹⁶ O ₄	220.0614	220.0610	1.8
¹⁸ O-FMK	$[M + H]^+$	C ₁₁ H ₁₃ N ₂ ¹⁶ O ₃ ¹⁸ O	239.0910	239.0918	-3.3
	$[(M + 2) - NH_3 + H]^+$	C ₁₁ H ₁₀ N ¹⁶ O ₃ ¹⁸ O	222.0660	222.0652	3.4

Table 1. Accurate mass measurements of the molecular ions and main fragments obtained for the O_2 ($^{1}\Delta_g$)-derived oxidation products of W

decomposition pathway. This result is consistent with the fragmentation patterns of the majority of amino acids, which show an initial protonation at the amino group, followed by proton transfer to the carboxylic acid moiety and fragmentation [26]. This common behavior is also in agreement with the "mobile proton model", which describes the dissociation of the molecule as a result of charge-directed cleavages that are initiated by intramolecular proton transfers [27, 28]. Interestingly, the fragmentation pattern of protonated alcohols is in contrast to that of protonated W, which almost exclusively loses NH₃ [29].

The major labeled protonated fragment observed for both alcohols (m/z 177) was shown to dissociate by three distinct routes. First, this ion loses C₂H₅N, yielding the labeled protonated ion at m/z 134. Also, H₂¹⁸O is lost, which gives rise to the ion at m/z 157, and the molecule further decomposes to the ion at m/z 130. Finally, the protonated fragment at m/z 177 eliminates neutral ammonia, generating the ion at m/z 160, which yields the ion at m/z 130 by the loss of C¹⁸O.

Besides the concomitant loss of H_2O and CO, the isomeric alcohols also eliminate H_2O . There are two possible sites of water elimination in the alcohol molecules: the hydroxyl oxygen of the carboxylic acid moiety or the alcohol group. The isotopic labeling approach was used to discriminate between these two possibilities. In this way, loss of labeled water from the alcohol moiety would result in a protonated fragment at m/z203. In contrast, elimination of unlabeled water from the carboxylic acid would generate a protonated fragment at m/z 205. The result showed that loss of water was



Figure 2. ESI MS/MS of *trans* alcohol synthesized under ${}^{16}O_2$ (a) or (b) ${}^{18}O_2$ atmospheres.



Scheme 2. Proposed fragmentation pathways for ¹⁸O-labeled alcohols.

exclusively from the alcohol moiety and loss of the hydroxyl oxygen of carboxylic acid moiety was not observed. This behavior was already reported for protonated glycine [26, 30]. The direct elimination of two neutral species (H_2O and CO) indicates that the intermediate ion is produced near or above the dissociation limit or that it is not a stable species.

The isomeric alcohols show the same fragment ions and the intensity of the ion at m/z 203 is the most notable difference. This fragmentation process occurs preferentially in the *cis* alcohol and could be explained considering the differences in the spatial arrangement of these two isomers. In the *cis* isomer, the distance between the protonated carboxylic acid and the alcohol oxygen could be small and could favor a proton transfer between these two interacting functional groups. The same behavior can also be observed in hydroperoxides, with the loss of hydrogen peroxide favored by the spatial arrangement of the *cis* isomer.

Finally, loss of NH₃ was a minor decomposition route for both alcohols, as judged by the relative intensity of the fragment at m/z 206 (or 204 in the case of the unlabeled molecule) (Figure 2 and Scheme **2**). Following this decomposition route, the fragment at m/z 206 undergoes successive cleavages, leading to the fragments at m/z 188 (loss of H₂O, m/z 186, in the case of the unlabeled molecule) and 160 (–CO, m/z 158, in the case of unlabeled WOH) and finally to the fragment at m/z130 (–C¹⁸O).

In the same way as the isomeric alcohols, *trans* and *cis* hydroperoxides, which elute at 7.92 and 10.28 min, respectively, in the UV chromatogram of Figure 1, were analyzed by MS/MS and ¹⁸O-isotopic labeling. Repre-

sentative results are shown in Figure 3a and b, respectively, for unlabeled and ¹⁸O-labeled *trans* hydroperoxides. As already discussed above, *cis* hydroperoxides exhibit the same fragmentation profile, except for the higher intensity of the fragment at m/z 220 (loss of H₂O₂), compared with the *trans* isomer.

Three fragmentation routes can be proposed for hydroperoxides. Two of these routes give rise to the stable ion at m/z 148 (or 146 in the case of the unlabeled molecule). When analyzing the precursor and fragment ion scans, it is possible to see that the ions at m/z 222 (or 220 for unlabeled molecule) and 168 (or 166 in the case of W¹⁶O¹⁶OH) are precursors of this stable ion (Scheme 3, the decomposition of ¹⁸O-labeled hydroperoxides). The chemistry associated with the dissociation of the ion at m/z 222 is consistent with a homolytic scission of the peroxide bond, yielding an alkoxyl radical, followed by a rearrangement and loss of an aminyl radical of glycine, generating the ion at m/z 148. Peroxides have relatively low bond energy compared with average bond energies of oxygen atoms bound to carbon or nitrogen. Consequently, the weak O-O bond linkage can be homolytically cleaved [31]. The homolytic scission of the peroxide linkage was previously reported in the electrospray collision-induced fragmentation of the lysine peroxycarbamates [32].

Loss of the immonium form of glycine ($C_2H_3O_2N$) from the molecular ion, followed by loss of labeled water is the other way to form the ion at m/z 148. The unlabeled form of this ion (m/z 146) has been identified as oxoquinoline, a characteristic fragment of Oia [18, 33]. Nevertheless, the fragmentation profile of the oxoquinolonium ion (protonated oxoquinoline) [18] is dif-



Figure 3. ESI MS/MS of *trans* hydroperoxide synthesized under ${}^{16}O_2$ (a) or (b) ${}^{18}O_2$ atmospheres.

ferent from the fragment at m/z 146 derived from hydroperoxides. The most abundant ion (m/z 118) arising from the former probably corresponds to the elimination of CO, which is consistent with the structure of the oxoquinolonium ion. In contrast, in the case of hydroperoxide, the main ion derived from the fragment at m/z 146 corresponds to the elimination of water. The last ion was obtained from the fragmentation of dioxindolylalanine [18]. The proposed structure for this ion is shown in Scheme **3**. The last hydroperoxide fragmentation pathway involves the sequential loss of H_2O_2 , yielding the ion at m/z 203, which then exhibits the same fragmentation pattern found for the isomeric alcohols.

Figure 4a and b show the fragmentation of FMK molecules synthesized with isotopes 16 and 18 of oxygen, respectively. As previously noted, unlabeled protonated FMK (Figure 4a) exhibits a molecular ion at m/z 237. The same ¹⁸O-isotopic labeling approach used to elucidate the fragmentation pathways of alcohols and



Scheme 3. Proposed fragmentation pathways for ¹⁸O-labeled hydroperoxides.



Figure 4. ESI MS/MS of FMK. (a) Fragment ion spectrum of unlabeled FMK. (b) Fragment ion spectrum of ¹⁸O-labeled FMK. (c) Fragment ion spectrum of unlabeled FMK incubated in the presence of $H_2^{18}O$.

hydroperoxides was employed for FMK. However, an unexpected problem related to this approach had to be overcome. In contrast to the successful incorporation of two ¹⁸O atoms found in WOOH (m/z 241), the mass spectrum of ¹⁸O-FMK shows a single atom of ¹⁸O (m/z 239, Figure 4b).

FMK contains carbonyl and amide functional groups. These functional groups have been reported to present high rates of oxygen exchange in water [34–36]. In this way, one can suggest that doubly labeled FMK molecules are able to exchange ¹⁸O atoms with solvent molecules, yielding a population of FMK containing just one ¹⁸O or even lacking ¹⁸O atoms. Thus, another strategy was employed to evaluate the ability of FMK to exchange its oxygen atoms. The strategy consisted of the incubation of unlabeled purified FMK (m/z 237) in the presence of $H_2^{18}O$. The fragment ion spectrum depicted in Figure 4c shows the incorporation of one ¹⁸O from labeled water in the FMK moiety. This experiment confirms that FMK is able to exchange oxygen atoms with the solvent. One interesting result is the inversion of some labeled fragments compared with ¹⁸O-labeled FMK (Figure 4b), showing that the amide and carbonyl fragments have different rates of oxygen exchange (see discussion below).

FMK has already been analyzed by MS [20, 21]. However, using the ¹⁸O-labeling approach, along with precursor and fragment ion experiments, it is possible to characterize the fragmentation mechanisms of this molecule. In contrast to isomeric WOH and WOOH, loss of NH₃ (m/z 222 or 220 in the case of unlabeled

molecule) is a favored pathway in the case of FMK. The same fragmentation behavior was obtained with similar molecules, namely W [29] and kn [22], although Oia [19] showed the preferential loss of H_2O and CO. The different behavior for structurally related molecules shows that small differences in the molecular arrangement, as well as the introduction of some functional groups, can greatly alter the proton affinity of the molecule. This is in agreement with the concepts of the mobile proton model [28], the role of neighboring groups to help cleavage reactions [37], and the fact that fragmentation reactions of protonated amino acids tend to be charge directed [29, 37].

Two parallel dissociation pathways can be proposed for the elimination of ammonia from protonated FMK (Scheme 4). First, a cyclized fragment, produced by the nucleophilic attack of the side-chain nitrogen with the α -carbon, can be proposed. This proposal is consistent with the concept of the neighboring group facilitating cleavage reactions [37] and was shown to occur in the elimination of ammonia from protonated W [29]. Second, a noncyclized structure, with initial formation of a carbocation, followed by proton transfer to oxygen of carboxylic acid group, can also be proposed. In fact, these two proposed fragmentation routes were based on the fragmentation profile of the deaminated fragment produced from deuterated kn [22].

Following NH_3 loss from protonated FMK, the generated fragment ion at m/z 222 (220, unlabeled) shows an initial loss of water followed by elimination of CO,



Scheme 4. Main proposed fragmentation pathways for ¹⁸O-labeled FMK.

yielding the protonated fragments at m/z 204 (202, unlabeled) and 176 (174, unlabeled) (Scheme 4). These fragmentation patterns are consistent with the maintenance of ¹⁸O-labeled fragments and corroborate the routes proposed for protonated kn. However, FMK has an additional formyl group compared with kn. For this reason, besides the analogy between the fragmentation pathways of these two molecules, the FMK deaminated fragment was also shown to lose CO. Loss of this neutral CO yields the protonated fragment at m/z 192 for both labeled and unlabeled FMK. Thus, this fragmentation route appears to refer to the loss of a labeled formyl group from deaminated fragment. Similarly to the protonated fragment at m/z 222 (220, unlabeled), the fragment at m/z 192 also showed successive losses of H_2O and CO, yielding protonated fragments at m/z 174 and 146.

To confirm the ¹⁸O-labeling in the formyl group, the spectrum obtained with labeled FMK was compared with spectrum obtained upon incubation of unlabeled FMK in the presence of $H_2^{18}O$ (Figure 4b and c). In the last case (Figure 4c), it is possible to see the inversion of some labeled fragments in the MS, compared with ¹⁸O-labeled FMK (Figure 4b). This inversion could be explained by different exchange susceptibilities between carbonyl and amide groups. The former is prone to a nucleophilic attack by water molecules, while the last one is relatively nonreactive due to its resonance stabilization. In this view, if the carbonyl group is more susceptible to oxygen exchanges, labeled FMK will easily change its ¹⁸O, but the more stable amide group will

keep its labeled oxygen atom. On the other hand, in the presence of $H_2^{18}O$, the amide group of unlabeled FMK will remain unaltered, while the carbonyl oxygen will now display an ¹⁸O atom. The explanation was confirmed by MS/MS experiments. For instance, in the case of unlabeled FMK incubated in the presence of $H_2^{18}O$, loss of ammonia and formyl groups produced a labeled fragment at m/z 194 (Figure 4c), instead of an unlabeled fragment at m/z 192 for the ¹⁸O- labeled molecule (Figure 4b). Also, in the unlabeled FMK maintained in the presence of $H_2^{18}O$, the direct loss of a formyl group produced a protonated fragment at m/z 211 (Figure 4c), in contrast to labeled FMK, which displayed a protonated fragment at m/z 209 (Figure 4b).

The last fragmentation route observed for FMK is the formation of a protonated ion at m/z 166 (or 164 in the case of the unlabeled molecule). The presence of this ion could be explained by the formation of an electrostatically-bound ion molecule complex before elimination of the imine form of glycine (Scheme 5). In fact, this pathway was previously proposed to explain the production of complementary fragment ions from kn [22]. The use of deuterated kn supports this conclusion.

Further fragmentation of the labeled protonated ion at m/z 166 yields the fragment ion at m/z 136 (loss of C¹⁸O). On the other hand, fragmentation of unlabeled FMK incubated in the presence of H₂¹⁸O generates the fragment ion at m/z 138 (loss of CO), supporting the differences in oxygen exchange susceptibility between carbonyl and amide groups and the formation of the electrostatically-bound ion molecule complex.



Scheme 5. Proposed fragmentation pathways for ¹⁸O-labeled FMK via the formation of an ion molecule complex.

Conclusions

The use of collisional activation reactions, associated with accurate mass measurements and ¹⁸O-isotopic labeling, has proven to be useful for the structural characterization of O_2 ($^1\Delta_{\circ}$)-derived oxidation products of W. Isomeric alcohols, hydroperoxides, and FMK share some common fragment ions and losses of small neutral molecules, such as H₂O, CO and NH₃. However, it was shown that small differences in the structure can greatly alter the fragmentation mechanism of structurally related molecules. In this way, tandem mass spectra revealed that each of the W oxidation products had its own fragmentation mechanism. The most stable fragment ion for isomeric WOH resulted from concomitant loss of H_2O and CO. On the contrary, the most stable fragment ion resulting from hydroperoxide dissociation was different. Two distinct routes contribute to yielding this stable fragment ion. The first one results from loss of the immonium form of glycine and water. The other possibility involves the scission of the peroxide bound to yield an alkoxyl radical, which through a rearrangement causes the loss of the aminyl radical of glycine. FMK was shown to dissociate by a diverse range of mechanisms, with the loss of ammonia the most favored route. ¹⁸O-labeling studies confirmed the fragmentation mechanisms and intermediates. Also, this approach showed the ability of FMK to exchange its oxygen atoms with water. Moreover, MS/MS and $H_2^{18}O$ experiments demonstrated that the oxygen atom from ketone is more susceptible to exchange with water than the formyl group.

The understanding of dissociation routes of $O_2(^{1}\Delta_g)$ mediated oxidation products of W reaffirms the importance of fragmentation mechanisms in the structural characterization and provides a useful step toward the analysis of W oxidation products in peptides and proteins. This work can assist in the investigation of oxidized modifications seen in some pathological conditions, such as protein aggregates found in neurodegenerative diseases.

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