



A Novel Framework for Molecular Characterization of Atmospheric Organic Aerosol Based on Collision Cross Section and Mass-to-Charge Ratio

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1 Abstract

2 A new metric is introduced for representing the molecular signature of atmospheric
3 organic aerosols, the collision cross section (Ω), a quantity that is related to the structure
4 and geometry of molecules and is derived from ion mobility measurements. By
5 combination with the mass-to-charge ratio (m/z), a two-dimensional $\Omega - m/z$ space is
6 developed to facilitate the comprehensive investigation of the complex organic aerosol
7 mixture. A unique distribution pattern of chemical classes, characterized by functional
8 groups including amine, alcohol, carbonyl, carboxylic acid, ester, and organic sulfate, is
9 developed on the 2-D $\Omega - m/z$ space. Species of the same chemical class, despite
10 variations in the molecular structures, tend to situate as a narrow band on the space and
11 follow a trend line. Reactions involving changes in functionalization and fragmentation
12 can be represented by the directionalities along or across these trend lines, thus allowing
13 for the interpretation of mechanisms associated with the formation and evolution of
14 atmospheric organic aerosol. The characteristics of trend lines for a variety of
15 functionalities that are commonly present in ambient aerosols can be predicted by the
16 core model simulations, which provide a useful tool to identify the chemical class to
17 which an unknown species belongs on the $\Omega - m/z$ space. Within the band produced by
18 each chemical class on the space, molecular structural assignment can be achieved by
19 utilizing collision induced dissociation as well as by comparing the measured collision
20 cross sections in the context of those obtained via molecular dynamics simulations.

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28 1. Introduction

29 Organic aerosol (OA) constitutes a major fraction of sub-micrometer atmospheric
30 particulate matter and comprises a complex and dynamic system linking source emission,
31 chemical transformation, and phase partitioning (Hallquist et al., 2009). It consists of a
32 multitude of organic species that arise from primary emissions and secondary
33 productions. Once in the atmosphere, OA species actively evolve via gas-particle
34 conversion and multiphase chemistry. The complexity and dynamic behaviors of ambient
35 OA have rendered identification of major pathways contributing to OA budget difficult
36 and have limited our capability to evaluate its impact on human health and global
37 climate.

38 Several two-dimensional frameworks have been developed in an effort to deconvolve
39 the complexity of OA mixtures and visualize their atmospheric transformations. The Van
40 Krevelen diagram, which scatter plots the hydrogen-to-carbon atomic ratio (H:C) and the
41 oxygen-to-carbon atomic ratio (O:C), has been widely used to represent the bulk
42 elemental composition and the degree of oxygenation of organic aerosol (Heald et al.,
43 2010). The average carbon oxidation state (\overline{OS}_C), a quantity that necessarily increases
44 upon oxidation, can be estimated from the elemental ratios (Kroll et al., 2011). When
45 coupled with carbon number (n_C), it provides constraints on the chemical composition of
46 organic aerosol and defines key classes of atmospheric processes based on the unique
47 trajectory of the evolving OA composition on the $\overline{OS}_C - n_C$ space. The degree of
48 oxidation has also been combined with the volatility (expressed as the effective saturation
49 concentration, C^*), forming a 2-D volatility basis set to describe the coupled aging and
50 phase partitioning of organic aerosol (Donahue et al., 2012). These three spaces are
51 designed to represent fundamental properties of the OA mixture and provide insight into
52 the OA chemical evolution in the atmosphere. Organic aerosol components span large
53 varieties in the physicochemical properties. Species of similar volatility or elemental
54 composition can differ vastly in structures and functionalities. One weakness of these
55 frameworks is that they do not provide information on the OA components at molecular
56 level.



57 In this article we introduce a new framework that is based on the collision cross
58 section (Ω), a quantity that is related to the structure and geometry of a molecule. The
59 collision cross section of a charged molecule determines its mobility as it travels through
60 a neutral buffer gas such as N_2 under the influence of a weak and uniform electric field.
61 Species with open conformation undergo more collisions with buffer gas molecules and
62 hence travel more slowly than the compact ones (Shvartsburg et al., 2000; Eiceman et al.,
63 2013). Mobility measurements are usually performed with an Ion Mobility Spectrometer
64 (IMS), where ions are separated mainly on the basis of their size, geometry, as well as
65 interactions with the buffer gas. The combination of IMS with a Mass Spectrometer (MS)
66 allows for further selection of ions based on their mass-to-charge ratios. The resulting
67 IMS-MS plot provides separation of molecules according to two different properties:
68 geometry (as reflected by the collision cross section) and mass (as reflected by the mass-
69 to-charge ratio) (Kanu et al., 2008). The Ion Mobility Spectrometry - Mass Spectrometry
70 (IMS-MS) analytical technique has been widely employed in the fields of biochemistry
71 (McLean et al., 2005; Liu et al., 2007; Dwivedi et al., 2008; Roscioli et al., 2013; Groessl
72 et al., 2015) and homeland security (Eiceman and Stone, 2004; Ewing et al., 2001;
73 Fernandez-Maestre et al., 2010). To our knowledge, the application of IMS-MS to study
74 organic species in the atmosphere, however, has only been explored very recently
75 (Krechmer et al., 2016).

76 We propose a two-dimensional collision cross section vs. mass-to-charge ratio ($\Omega -$
77 m/z) space to facilitate the comprehensive investigation of complex OA mixtures. Despite
78 the typical complexity of the detailed molecular mechanism of OA production and
79 evolution, oxidized molecules that constitute OA can be characterized by their distinctive
80 functional groups (Zhang and Seinfeld, 2013). We show that the investigated organic
81 classes ($m/z < 600$), characterized by functional groups including amine, alcohol,
82 carbonyl, carboxylic acid, ester, and organic sulfate, exhibit unique distribution patterns
83 on the $\Omega - m/z$ space. Species of the same chemical class, despite variations in the
84 molecular structures, tend to develop a narrow band and follow a trend line on the space.
85 Reactions involving changes in functionalization and fragmentation can be represented
86 by directionalities along or across these trend lines. The locations and slopes of the
87 measured trend lines are shown to be predicted by the core model (Mason et al., 1972),



88 which characterizes the ion-neutral interactions as elastic sphere collisions. Within the
89 narrow band produced by each chemical class on the $\Omega - m/z$ space, molecular structural
90 assignment is achieved with the assistance of collision induced dissociation analysis.
91 Measured collision cross sections are also shown to be consistent with theoretically
92 predicted values from the trajectory method (Mesleh et al., 1996; Shvartsburg and
93 Jarrold, 1996) and are used to identify isomers that are separated from an isomeric
94 mixture.

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96 **2. Collision Cross Section Measurements**

97 **2.1 Materials**

98 A collection of chemical standards (ACS grade, $\geq 96\%$, purchased from Sigma
99 Aldrich, St. Louis, MO, USA), classified as amines, alcohols, carbonyls, carboxylic
100 acids, esters, phenols, and organic sulfates, were used to characterize the performance of
101 IMS-MS. These chemicals were dissolved in an HPLC-grade solvent consisting of a 70%
102 methanol / 29% water with 1% formic acid, at a concentration of approximately 10 μM .

103 **2.2 Instrumentation**

104 Ion mobility measurements were performed using an Electrospray Ionization (ESI)
105 Drift-Tube Ion Mobility Spectrometer (DT-IMS) interfaced to a Time-of-Flight Mass
106 Spectrometer (TOFMS). The instrument was designed and manufactured by TOFWERK
107 (Switzerland), with detailed descriptions and schematics provided by several recent
108 studies (Kaplan et al., 2010; Zhang et al., 2014; Groessl et al., 2015; Krechmer et al.,
109 2016). In the next few paragraphs, we will present the operating conditions of the ESI-
110 IMS-TOFMS instrument.

111 Solutions of chemical standards were delivered to the ESI source via a 250 μL gas-
112 tight syringe (Hamilton, Reno, NV, USA) held on a syringe pump (Harvard Apparatus,
113 Holliston, MA, USA) at a flow rate of 1 $\mu\text{L min}^{-1}$. A deactivated fused silica capillary
114 (360 μm OD, 50 μm ID, 50 cm length, New Objective, Woburn, MA, USA) was used as
115 the sample transfer line. The ESI source was equipped with an uncoated SilicaTip Emitter
116 (360 μm OD, 50 μm ID, 30 μm tip ID, New Objective, Woburn, MA, USA) and



117 connected to the capillary through a conductive micro union (IDEX Health & Science,
118 Oak Harbor, WA, USA). The ESI emitter was operated at both positive and negative
119 mode at a capillary voltage of $\pm (1.5 - 2.0)$ kV. The charged droplets generated at the
120 emitter tip migrate through a desolvation region in nitrogen atmosphere at room
121 temperature, where ions evaporate from the droplets and are introduced into the drift tube
122 through a Bradbury-Nielsen ion gate located at the entrance. The ion gate was operated in
123 the Hadamard Transform mode, with a closure voltage of ± 50 V and an average gate
124 pulse frequency of 1.2×10^3 Hz. The drift tube was held at a constant temperature (340 ± 3
125 K) and atmospheric pressure (~ 1019 mbar). A counter flow of N_2 drift gas was
126 introduced at the end of the drift region at a flow rate of 1.2 L min^{-1} . Ion mobility
127 separation was carried out at a typical field strength of $300 - 400$ V cm^{-1} , resulting in a
128 reduced electric field of approximately $1.4 - 1.8$ Td. After exiting from the drift tube,
129 ions were focused into TOFMS through a pressure-vacuum interface that includes two
130 segmented quadrupoles that were operated at ~ 2 mbar and $\sim 5 \times 10^{-3}$ mbar, respectively.
131 Collision Induced Dissociation (CID) of parent ions is achieved by adjusting the voltages
132 on the ion optical elements between the two quadruple stages (Kaplan et al., 2010).

133 The ESI-IMS-TOFMS instrument was operated in the m/z range of 40 to 1500 with a
134 total recording time of 90 s for each dataset. The Mass Spectrometer was calibrated using
135 a mixture of quaternary ammonium salts, reserpine, and a mixture of fluorinated
136 phosphazines (Ultramark 1621) in the positive mode and ammonium phosphate, sodium
137 dodecyl sulfate, sodium taurocholate hydrate, and Ultramark 1621 in the negative mode.
138 The ion mobility measurements were calibrated using tetraethyl ammonium chloride as
139 the instrument standard and 2,4-lutidine as the mobility standard, as defined shortly
140 (Fernández-Maestre et al., 2010). Mass spectra and ion mobility spectra were recorded
141 using the acquisition package “Acquility” (v2.1.0, <http://www.tofwerk.com/acquility>).
142 Post-processing was performed using the data analysis package “Tofware” (version 2.5.3,
143 www.tofwerk.com/tofware) running in the Igor Pro (Wavemetrics, OR, USA)
144 environment.



145 2.3 Calculations

146 The average velocity of an ion in the drift tube (v_d) is proportional to its characteristic
147 mobility constant ($K / \text{cm}^2 \text{V}^{-1} \text{s}^{-1}$) and the electric field intensity (E_d), provided that the
148 field is weak (McDaniel and Mason, 1973):

$$149 \quad v_d = K E_d \quad (1)$$

150 Experimentally, ion mobility constants can be approximated from the time of ion clouds
151 spent in the drift tube (t_d / s), given by the rearranged form of Equation (1):

$$152 \quad t_d = \frac{1}{K} \frac{L_d^2}{V_d} \quad (2)$$

153 where L_d (cm) is the length of the drift tube and V_d (V) is the drift voltage. In the present
154 study, drift time measurements were carried out at six different drift voltages ranging
155 from 5 kV to 8 kV in $\sim 1019 \text{ mbar}$ of nitrogen gas at 340 K (Figure S1 in the
156 supplement). The ion mobility constant (K) is derived by linear regression of the recorded
157 arrival time (t_a) of the ion clouds at the detector versus the reciprocal drift voltage:

$$158 \quad t_a = \frac{L_d^2}{K} \frac{1}{V_d} + t_0 \quad (3)$$

159 Note that the arrival time was determined from the centroid of the best-fit Gaussian
160 distribution, see Figure S2 in the Supplement. The y-intercept of the best-fit line
161 represents the transport time of the ion from the exit of the drift tube to the MS detector
162 (t_0), which exhibits strong m/z dependency that is attributable to a time-of-flight
163 separation in the ion optics, see Figure S3 in the Supplement.

164 It is practical to discuss an ion's mobility in terms of the reduced mobility constant
165 (K_0), defined as:

$$166 \quad K_0 = K \frac{273.15}{T} \frac{P}{1013.25} \quad (4)$$

167 where P (mbar) is the pressure in the drift region and T (K) is the buffer gas temperature.
168 In theory, the parameter K_0 is constant for a given ion in a given buffer gas and can be
169 used to characterize the intrinsic interactions of that particular ion-molecule pair. In
170 practice, however, K_0 values from different measurements might not be in good



171 agreement, primarily due to uncertainties in instrumental parameters such as
 172 inhomogeneities in drift temperature and voltage (Fernández-Maestre et al., 2010). In
 173 view of these uncertainties, the instrument standard (the reduced mobility of such a
 174 standard is not affected by contaminants in the buffer gas) is needed to provide an
 175 accurate constraint on the instrumental parameters, such as voltage, drift length, pressure,
 176 and temperature:

$$177 \quad K_0 \times t_d = \frac{L_d^2}{V_d} \frac{P}{1013.25} \frac{273.15}{T} = C_i \quad (5)$$

178 Tetraethyl ammonium chloride (TEA) is used here as the instrument standard
 179 (Fernández-Maestre et al., 2010). Given the well-known K_0 and measured t_d of the
 180 protonated TEA ion ($m/z = 130$), Equation (5) yields an instrument constant C_i
 181 to calibrate the IMS performance.

182 Unlike TEA, the reduced mobility of species that are more likely to cluster with
 183 contaminants can be significantly affected by impurities of the buffer gas. This category
 184 of species can be used as a ‘mobility standard’ to qualitatively indicate the potential
 185 contamination in the buffer gas. 2,4-Lutidine, with a well-characterized K_0 value of 1.95
 186 $\text{cm}^2 \text{V}^{-1} \text{s}^{-1}$, is used as such a mobility standard. As shown Figure S4 in the Supplement,
 187 the measured mobility of 2,4-Lutidine is 1.5% lower than its theoretical value, indicative
 188 of the absence of contaminations in the buffer gas.

189 In the low field limit, the collision cross section of an ion (Ω) with a buffer gas is
 190 related to its reduced mobility (K_0) through the modified zero field (so called Mason-
 191 Schamp) equation (McDaniel and Mason, 1973; Siems et al., 2012):

$$192 \quad \Omega = \frac{3ze}{16N_0} \left(\frac{2\pi}{k_B \mu T_0} \right)^{1/2} \frac{1}{K_0} \left[1 + \left(\frac{\beta_{MT}}{\alpha_{MT}} \right)^2 \left(\frac{v_d}{v_T} \right)^2 \right]^{-1/2} \quad (6)$$

193 where z is the net number of integer charges on the ion, e is the elementary charge, N_0 is
 194 the number density of buffer gas at 273 K and 1013 mbar, k_B is the Boltzmann constant,
 195 μ is the reduced mass for the molecule-ion pair, T_0 is the standard temperature, v_d is the
 196 drift velocity given by Equation (1), v_T is the thermal velocity, and α_{MT} and β_{MT} are
 197 correction coefficients for collision frequency and momentum transfer, respectively,
 198 given by:



$$\alpha_{\text{MT}} = \frac{2}{3} [1 + \hat{m}f_c + \hat{M}f_h] \quad \beta_{\text{MT}} = \left[\frac{2}{\hat{m}(1 + \hat{m})} \right]^{1/2} \quad (7)$$

200 where \hat{m} and \hat{M} are mass fractions of the ion and buffer gas molecule, respectively, and f_c
201 and f_h are the fractions of collisions in the cooling and heating classes, respectively. Note
202 that the reduced electric field used in this study is maximized at ~ 2 Td, at which the drift
203 velocity of any given ion is \sim two orders magnitude lower than its thermal velocity, thus
204 the values for f_c and f_h are assigned to be 0.5 and 0.5, respectively. As all measurements
205 in this study were carried out with nitrogen as the buffer gas, the reported collision cross
206 sections will be referred to Ω_{N_2} . Experimental Ω_{N_2} values for a selection of ionic species
207 are consistent with those reported in literatures (see Table S1 in the Supplement).

208

209 3. Collision Cross Section Modeling

210 Kinetic theory indicates that the quantity Ω is an orientationally averaged collision
211 integral ($\Omega_{\text{avg}}^{(l,l)}$), which depends on the nature of ion-neutral interaction potential
212 (McDaniel and Mason, 1973). Given the potential, the collision integral can be calculated
213 through successive integrations over collision trajectories, impact parameters and energy.
214 Here we adopt two computational methods, i.e., trajectory method and core model, to
215 simulate the average collision integral. The trajectory method is a rigorous calculation of
216 $\Omega_{\text{avg}}^{(l,l)}$ by propagating classical trajectories of neutral molecules in a realistic neutral/ion
217 potential consisting of a sum of pairwise Lennard-Jones interactions and ion induced
218 dipole interactions (Mesleh et al., 1996; Shvartsburg and Jarrold, 1996). The core model
219 treats the polyatomic ion as a rigid sphere where the center of charge is displaced from
220 the geometry center. The ion-neutral interaction is approximately represented by the cross
221 section of two rigid spheres during elastic collisions. The (12,4) potential, which includes
222 a long-range polarization term and a short-range repulsion term, is incorporated in the
223 core model (Mason et al., 1972).

224 The two models employed here represent opposite directions in the $\Omega_{\text{avg}}^{(l,l)}$ computation
225 methods. The trajectory method is a rigorous calculation of $\Omega_{\text{avg}}^{(l,l)}$ in a realistic
226 intermolecular potential yet the computation is time consuming. The core model, on the



227 other hand, substantially simplifies the calculation of $\Omega_{\text{avg}}^{(l,1)}$ as rigid sphere collisions at the
228 expense of simulation accuracy. We will show shortly that the core model is used for
229 locating individual chemical classes on the 2-D $\Omega_{\text{N}_2} - m/z$ space. Within the band
230 developed by each chemical class, molecular structure information can be deduced by
231 comparing the measured collision cross section with those calculated by the trajectory
232 method.

233 3.1 Trajectory Method

234 Molecular structures for L-leucine and D-isoleucine were initially constructed by
235 Avogadro v1.1.1 (Hanwell et al., 2012). For each molecule, both protonation and
236 deprotonation sites are created by placing a positive charge on the N-terminal amino
237 group and a negative charge on the C-terminal carboxyl group, respectively. The
238 geometry of each ion is further optimized using the Hartree-Fock method with the 6-
239 31G(d,p) basis set via GAMESS (Schmidt et al., 1993). Partial atomic charges were
240 estimated using Mulliken population analysis.

241 A freely available software, MOBCAL, developed by Jarrold and coworkers
242 (<http://www.indiana.edu/~nano/software.html>) was used for computing the collision
243 integrals. The potential term employed in the trajectory method takes the form:

$$244 \quad \Phi(\theta, \phi, \gamma, b, r) = 4\epsilon \sum_i^n \left[\left(\frac{\sigma}{r_i} \right)^{12} - \left(\frac{\sigma}{r_i} \right)^6 \right] - \frac{\alpha_p}{2} \left(\frac{ze}{n} \right)^2 \left[\left(\sum_i^n \frac{x_i}{r_i^3} \right)^2 + \left(\sum_i^n \frac{y_i}{r_i^3} \right)^2 + \left(\sum_i^n \frac{z_i}{r_i^3} \right)^2 \right] \quad (8)$$

245 where θ , ϕ , and γ are three angles that define the geometry of ion-neutral collision, b is
246 the impact parameter, ϵ is the depth of the potential well, σ is the finite distance at which
247 the interaction potential is zero, α_p is the polarizability of the neutral, which is 1.710×10^{-24}
248 cm^3 for N_2 (Olney et al., 1997), n is the number of atoms in the ion, and r_i , x_i , y_i , and z_i
249 are coordinates that define the relative positions of individual atoms with respect to the
250 buffer gas. Values of the Lenard-Jones parameters, ϵ and σ , are taken from the universal
251 force field (Casewit et al., 1992). The ion-quadruple interaction and the orientation of N_2
252 molecule are not considered here (Kim et al., 2008; Campuzano et al., 2012).



253 3.2 Core Model

254 The core model, consisting of a (12-4) central potential displaced from the origin, is
 255 used to represent interactions of polyatomic ions with N₂ molecules (Mason et al., 1972).
 256 The central potential includes the common long-range r^{-4} polarization energy, as well as
 257 the short-range r^{-12} overlap repulsion energy:

$$258 \quad \Phi(r) = \frac{\epsilon}{2} \left\{ \left(\frac{r_m - a}{r - a} \right)^{12} - 3 \left(\frac{r_m - a}{r - a} \right)^4 \right\} \quad (9)$$

259 where r is the distance between the ion-neutral geometric centers, a is the location of the
 260 ionic center of charge measured from the geometrical center of the ion, and r_m is the
 261 value of r at the potential minimum. At temperature of 0 K, the *polarization potential* can
 262 be expressed as:

$$263 \quad \Phi_{\text{pol}}(r) = -\frac{e^2 \alpha_p}{2r^4} \quad (10)$$

264

265 where α_p is the polarizability of the neutral. Thus ϵ is given by:

$$266 \quad \epsilon = \frac{e^2 \alpha_p}{3(r_m - a)^4} \quad (11)$$

267 The collision cross section can be expressed in dimensionless form by extracting its
 268 dependence on r_m :

$$269 \quad \Omega = \Omega^{(1,1)*} \pi r_m^2 \quad (12)$$

270 Tabulations of the dimensionless collision integral ($\Omega^{(1,1)*}$) can be found in literatures
 271 (Mason et al., 1972) as a function of dimensionless temperature (T^*) and core diameter
 272 (a^*), given by:

$$273 \quad T^* = \frac{kT}{\epsilon} = \frac{3kT(r_m - a)^4}{e^2 \alpha_p} \quad a^* = \frac{a}{r_m} \quad (13)$$

274 Polynomial interpolation of the tabulated $\Omega^{(1,1)*}$ yielded an analytical expression of the
 275 collision cross section, with r_m and a as adjustable parameters. This expression was then
 276 fit to the ion mobility datasets measured in N₂ buffer gas using a nonlinear least-square



277 regression procedure (Matlab code is available upon request) (Johnson et al., 2004; Kim
278 et al., 2005; Kim et al., 2008). Best-fit parameters, r_m and a , along with predicted vs.
279 measured collision cross section are given in Table S2 in the Supplementary Information.

280

281 4. Collision Cross Section vs. Mass-to-Charge Ratio 2-D Space

282 4.1 Distribution of *multi*-Functional Organic Species

283 Figure 1 (A) shows the distribution of organic species, classified as (*di/poly/sugar*)-
284 alcohol, *tertiary*-amine, *quaternary*-ammonium, (*mono/di*)-carbonyl, (*mono/di/tri*)-
285 carboxylic acid, (*di*)-ester, organic sulfate, and *multi*-functional compounds, on the
286 collision cross section vs. mass-to-charge ratio ($\Omega_{N_2} - m/z$) 2-D space. Note that analytes
287 that are detected in different ion modes (+/-) are plotted separately. One feature of the
288 distribution pattern is that species with higher density as pure liquids and carbon
289 oxidation state tend to occupy the lower region of the $\Omega_{N_2} - m/z$ space. This is not
290 surprising given that molecules of smaller collision cross sections tend to be much
291 denser, and potentially more functionalized, than those with extended and open
292 geometries. Furthermore, species of the same chemical class tend to occupy a narrow
293 region and follow a trend line on the $\Omega_{N_2} - m/z$ space. These observations form the basis
294 of potentially utilizing locations and trends on the 2-D space to identify chemical classes
295 to which an unknown compound belongs.

296 Small molecules ($m/z < 200$) with similar size and geometry are situated closely
297 together, as visualized by the ‘overlaps’ on the space. Improved visual separation of the
298 species within the overlapping region is obtained by transforming Ω_{N_2} to a quantity
299 $\Delta\Omega_{N_2}$, defined as the percentage difference between the measured collision cross section
300 for any given molecular ion and the calculated projection area for a rigid spherical ion- N_2
301 pair with the same molecular mass. A density of 1.2 g cm^{-3} , which represents the average
302 bulk density of ambient organic aerosol (Turpin and Lim, 2001), is used as the reference
303 value. Since this idealized ion- N_2 pair does not account for interaction potentials and
304 molecular conformation, it is only used as a reference state to improve visualization of
305 the $\Omega_{N_2} - m/z$ 2-D space, as shown in Figure 1 (B).



306 It is important to note that if these ‘overlapping’ molecules belong to different
307 chemical classes, they can be resolved based on the fragmentation pattern or
308 characteristic fragments upon collision induced dissociation (CID), as discussed in detail
309 in Section 4.3. Also note that isomeric and isobaric species can be identified by
310 comparing the measured collision cross sections with those obtained from trajectory
311 method simulations, see Section 4.4. As it is highly unlikely that two distinct molecules
312 will produce identical IMS, MS, as well as CID-based MS spectra, the 2-D framework
313 therefore virtually ensures reliable identification of compounds.

314 Reactions involving changes in functionalization and fragmentation can be represented
315 by an intrinsic directionality on the $\Omega_{N_2} - m/z$ space, as illustrated by the distribution
316 pattern of carboxylic acid series shown in Figure 2. Addition of one carbon always leads
317 to an increase in mass and collision cross section, with a generic slope of approximately 5
318 $\text{Å}^2/\text{Th}$. Although the addition of one oxygen in the form of a carbonyl group results in a
319 similar increase in the molecular mass, it leads to a shallower slope compared with that
320 from expanding the carbon chain. Addition of carboxylic or hydroxyl groups leads to a
321 substantial decrease in the collision cross section, due to the formation of a cyclic
322 conformation by the intramolecular hydrogen bonding ($\text{O} - \text{H} \cdots \text{O}^-$).

323 4.2 $\Omega_{N_2} - m/z$ Trend Lines

324 The $\Omega_{N_2} - m/z$ trend line visualized on the 2-D space describes the intrinsic increase
325 in collision cross sections resulting from the increase in molecular mass by extending the
326 carbon backbone or adding functional groups. It has been used for conformation space
327 separation of different classes of biomolecules including lipids, peptides, carbohydrates,
328 and nucleotides (McLean et al., 2005). Here we demonstrate for the first time the
329 presence of trend lines for small molecules of atmospheric interest, and the trend line
330 pattern for each chemical class can be predicted by the core model simulations.

331 Figure 3 shows the measured Ω_{N_2} as a function of mass-to-charge ratio for (A)
332 *tertiary*-amine and *quaternary*-ammonium, (B) (*di/poly/sugar*)-alcohol, and (C)
333 (*mono/oxo/hydroxy*)-carboxylic acid. Also shown are the predicted Ω_{N_2} by the core
334 model, with adjustable parameters optimized by the measured Ω_{N_2} for the subcategory



335 spanning the largest m/z range in each chemical class. Specifically, *quaternary-*
336 ammonium, propylene glycol, and *alkanoic-acid* are used in constraining the core model
337 performance to predict the $\Omega_{N_2} - m/z$ trend lines for amines, alcohols, and carboxylic
338 acids. Species in each chemical class, regardless of the variety in the carbon skeleton
339 structure, occupy a narrow range and appear along a $\Omega_{N_2} - m/z$ trend line. Such a
340 relationship can be further demonstrated by the goodness of the core model predictions,
341 i.e., the difference between predicted and measured Ω_{N_2} for compounds that are not used
342 to optimize the core model performance. For amine series, predicted Ω_{N_2} values for
343 lutidine and pyridine are 8.2% and 0.8% higher, respectively, than the measurements. For
344 alcohol series, the best-fit $\Omega_{N_2} - m/z$ trend line constrained by propylene glycol can be
345 used to predict the distribution of sugars and polyols within 3.5% difference on the space.
346 For carboxylic acid series, hydroxyl-hexadecanoic acid falls closely on the predicted
347 $\Omega_{N_2} - m/z$ trend line, despite the presence of an alcohol group on the C₁₆ carbon chain.
348 Predicted Ω_{N_2} values for *oxo-carboxylic acids* are 4.4% – 6.1% lower than the
349 observations. Benzoic acid exhibits a relatively large measurement-prediction gap (6.7%)
350 potentially due to the presence of an aromatic ring.

351 Overall, the demonstrated $\Omega_{N_2} - m/z$ trend lines for carboxylic acids, amines, and
352 alcohols provide a useful tool for classification of structurally related compounds on the
353 space. It is worth noting that the core model optimization and simulation can be certainly
354 extended to other functionalities with the availability of chemical standards. Mapping out
355 the locations and distribution patterns for various functionalities on the 2-D space would
356 greatly facilitate structural identification of unknown compounds of atmospheric interest.

357 4.3 Molecular Structure Elucidation of *multi-Functional Species*

358 The demonstrated $\Omega_{N_2} - m/z$ relationship provides a useful tool to identify the
359 chemical class to which an unknown species belongs. To further identify its molecular
360 structure, knowledge on the electrospray ionization mechanism for the generation of
361 *quasi-molecular ions*, as well as fragmentation patterns of the molecular ion upon
362 collision induced dissociation (CID), is required.



363 For species investigated in this study, the positive mass spectra collected for amines
364 and amino acids show major ions at m/z values corresponding to the protonated cations
365 ($[M+H]^+$). Sodiated clusters ($[M+Na]^+$) of esters were observed as the dominant peak in
366 the ESI(+) spectra. Aromatic aldehydes combine with a methyl group ($[M+CH_3]^+$) via the
367 gas-phase aldol reaction between protonated aldehydes and methanol in the positive
368 mode. Sugars and polyols can be readily ionized in both positive and negative mode with
369 the addition of a proton or sodium ion or deprotonation. Extensive formation of
370 oligomers is observed from the positive mass spectra of propylene glycol, with the
371 deprotonated propanol ($-OCH_2CH(CH_3)-$) as the primary building block. Monoanions
372 ($[M-H]^-$) were exclusively observed in the negative mass spectra of (*mono/di/tri/multi*)-
373 carboxylic acids due to the facile ionization afforded by the carboxylic group. An
374 exhibition of molecular formulas of ionic species is given in Table 1.

375 The instrument used in this study enables the collision induced dissociation of the
376 abovementioned precursor ions after ion mobility separation but prior to the mass
377 spectrometer (IMS-CID-MS). As a consequence, product ions exhibit the identical
378 mobility (drift time) with that of the precursor ion. IMS-CID-MS spectra for individual
379 compounds are then generated by the extraction of “mobility-selected” MS spectra that
380 contain both precursor and fragments. The major advantage of this approach is that it is
381 possible to obtain fragmentation spectra for all precursor ions simultaneously. This is in
382 contrast to MS/MS techniques which require the isolation of a small mass window prior
383 to fragmentation which can be a problem for very complex samples or time-resolved
384 analysis. Figure 4 shows the measured drift time for the precursor and product ions
385 generated from species representative of amines, aldehydes, carboxylic acids, esters, and
386 nitro compounds. Collision induced dissociation patterns of these species are used to
387 elucidate the fragmentation mechanisms for corresponding functional groups. The
388 deprotonated carboxylic acid is known to undergo facile decarboxylation to produce a
389 carbanion. If additional carboxylic groups are present in the molecule, combined loss of
390 water and carbon dioxide is expected (Grossert et al., 2005). Alternatively, the presence
391 of an $-OH$ group adjacent to the carboxylic group would usually result in a neutral loss of
392 formic acid (Greene et al., 2013), see the fragmentation pattern for 16-
393 hydroxyhexadecanoic acid as an illustration. Scission of the C–O bond in the ester



394 structure or the C–O bond between the secondary/tertiary carbon and the alcoholic
395 oxygen is observed for the ester series examined, consistent with previous studies (Zhang
396 et al., 2015). A primary fragmentation resulting in loss of CO was evident in the spectrum
397 of methylate derivative of protonated carbonyls (RCHOCH₃⁺) (Neta et al., 2014). The
398 IMS-CID-MS spectrum of deprotonated 4-nitrophenol is shown as a representative of
399 organic nitro compounds. Two dominant peaks at m/z 108 and m/z 92 are observed,
400 resulting from the neutral loss of NO and NO₂, respectively.

401 Signal intensities of the fragments from the CID pathway of the precursor ion depend
402 on the collision voltage, as shown in Figure 5. At low collision voltages, the precursor
403 ions predominate with transmission optimized at approximately 5 V potential gradient.
404 As the collision voltage increases, the intensity of the precursor ion decreases and that of
405 each product ion increases, eventually reaching a maximum level, and then decreases due
406 to subsequent fragmentation. The dependence of the product ion abundance on the
407 collision voltage provides information on the relative strength of the covalent bond at
408 which the parent molecule fragments. Consequently, the energy required to induce a
409 certain fragmentation pathway could potentially also serve as an additional parameter for
410 structure elucidation. For example, the predominance of the product ion at m/z 149
411 suggests that cleavage of the O–O bond in the ester moiety is the dominant fragmentation
412 pathway upon CID of dioctyl phthalate (C₂₄H₃₈O₄).

413

414 4.4 Resolving Isomeric Mixtures

415 Here we demonstrate the separation and identification of isomers on the $\Omega_{N_2} - m/z$
416 space using the mixture of L-leucine and D-isoleucine as an illustration. Leucine can be
417 directly ionized by electrospray in both positive and negative modes due to the presence
418 of amino and carboxyl groups. Figure 6 (A and B) shows a single peak that corresponds
419 to the protonated ([M+H]⁺, m/z = 132) and deprotonated ([M–H][–], m/z = 130) forms of
420 the leucine mixture, respectively, in the positive and negative MS spectra. Upon further
421 separation based on their distinct mobility in the N₂ buffer gas, the leucine mixture is
422 clearly resolved in the positive mode, while a broad peak is observed in the negative ion
423 mobility spectrum, see Figure 6 (C and D). Note that a typical IMS resolving power



424 (t/dt_{50}) of 100 leads to a baseline separation of leucine isomers that differ by 0.3 ms in the
425 measured drift time. Figure 6 (E-H) shows the IMS spectra for individual leucine
426 isomeric configurations, which provide precise constraints for the peak assignment in the
427 leucine mixture. Also given here are the measured vs. predicted collision cross sections
428 for each isomer, with predictions lower by 3.3 ~ 6.9% compared with the measurements.
429 However, despite the underprediction, the model using trajectory method correctly
430 predicts the relative collision cross sections of the isomers and therefore also the order in
431 which they appear in the IMS spectrum. The underprediction of Ω_{N_2} may result from the
432 simplification that linear N_2 molecules are considered as elastic and specular spheres in
433 the current model configuration (Larriba-Andaluz and Hogan Jr, 2014). Further
434 development of the model to more appropriately predict Ω_{N_2} values is needed.

435

436 5. Conclusions

437 We propose a new metric, collision cross section (Ω), for characterizing organic
438 species of atmospheric interest. Collision cross section represents an effective interaction
439 area between a charged molecule and neutral buffer gas as it travels through under the
440 action of a weak electric field, and thus relates to the chemical structure and 3-D
441 conformation of this molecule. The collision cross section of individual molecular ions is
442 calculated from the ion mobility measurements using an Ion Mobility Spectrometer. In
443 this study, we provide the derived Ω_{N_2} values for a series of organic species including
444 amines, alcohols, carbonyls, carboxylic acids, esters, organic sulfates, and *multi-*
445 *functional* compounds.

446 The collision cross section, when coupled with mass-to-charge ratio, provides a 2-D
447 framework for characterizing the molecular signature of atmospheric organic aerosol.
448 The $\Omega_{N_2} - m/z$ space is employed to guide our fundamental understanding of processes
449 of organic aerosol formation and evolution in the atmosphere. We show that different
450 chemical classes tend to develop unique narrow bands with trend lines on the $\Omega_{N_2} - m/z$
451 space. Trajectories associated with atmospheric transformation mechanisms either cross
452 or follow these trend lines through the space. The demonstrated $\Omega_{N_2} - m/z$ trend lines



453 provide a useful tool for resolving various functionalities in the complex OA mixture.
454 These intrinsic trend lines can be predicted by the core model, which provides a guide for
455 locating unknown functionalities on the $\Omega_{N_2} - m/z$ space.

456 Within each band that that belongs to a particular chemical class on the space, species
457 can be further separated based on their distinct structures and geometries. We
458 demonstrate the utility of collision induced dissociation technique, upon which the
459 resulted product ions share the identical drift time as the precursor ion, to facilitate the
460 elucidation of molecular structures of OA constituents. We employ the $\Omega_{N_2} - m/z$
461 framework for separation of isomeric mixtures as well by comparing the measured
462 collision cross sections with those predicted using the trajectory method. Further
463 advances in algorithms to correctly predict collision cross sections *ab initio* from
464 molecular coordinates are therefore also expected to significantly improve identification
465 of unknowns.

466

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473

474 **Appendix:**

475 a (Å): the location of the ionic center of charge from the geometrical center of the ion.

476 a^* : the dimensionless core diameter.

477 α_{MT} : the correction coefficient for collision frequency.

478 α_p (cm³): the polarizability of the neutral.

479 β_{MT} : the correction coefficient for momentum transfer.

480 C_i : the instrument constant that is used to calibrate the IMS performance.

481 ϵ (eV): the depth of the potential well.



- 482 E_d (V/cm): the electric field intensity in the drift tube.
483 Φ (eV): the ion-neutral interaction potential.
484 f_c : the fraction of collisions in the cooling classes.
485 f_h : the fraction of collisions in the heating classes.
486 k_B ($\text{m}^2 \text{kg s}^{-2} \text{K}^{-1}$): Boltzmann constant.
487 K ($\text{cm}^2 \text{V}^{-1} \text{s}^{-1}$): the characteristic mobility constant of a given ion.
488 K_0 ($\text{cm}^2 \text{V}^{-1} \text{s}^{-1}$): the reduced mobility constant of a given ion.
489 L_d (V/cm): the length of the drift tube.
490 \hat{m} : the mass fraction of the ion in the ion-molecule pair.
491 \hat{M} : the mass fraction of the buffer gas molecule (N_2) in the ion-molecule pair.
492 m/z (Th): the mass-to-charge ratio of any given ion.
493 N_0 ($\# \text{cm}^{-3}$): the number density of buffer gas at 273 K and 1013 mbar.
494 Ω (\AA^2): the collision cross section.
495 Ω_{N_2} (\AA^2): the collision cross section using N_2 as the buffer gas.
496 $\Omega_{\text{avg}}^{(1,1)}$: the orientationally averaged collision integral.
497 $\Omega^{(1,1)*}$: the dimensionless collision integral.
498 P (mbar): the pressure in the drift region.
499 r (\AA): the distance between the ion-neutral geometric centers.
500 r_m (\AA): the value of r at the potential minimum.
501 σ (\AA): the finite distance at which the interaction potential is zero.
502 T (K): the buffer gas temperature.
503 T_0 (K): the standard temperature.
504 T^* : the dimensionless temperature.
505 t_a (s): the recorded arrival time of the ion clouds at the detector.
506 t_d (s): the time of ion clouds spent in the drift tube.
507 t_0 (s): the transport time of ion clouds from the exit of the drift tube to the MS detector.
508 v_d (s): the average velocity of an ion in the drift tube.
509 v_T (m s^{-1}): the thermal velocity.
510 V_d (V): the voltage applied to the drift tube.
511 z : the net number of integer charges on the ion.

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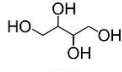
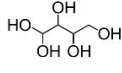
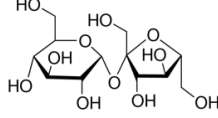


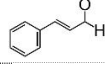
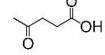
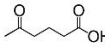
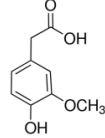
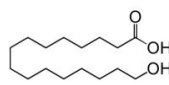
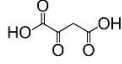
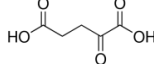
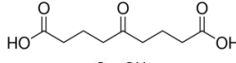
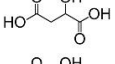
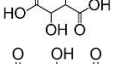
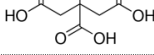
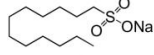
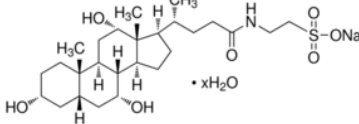
Table 1. Overview of organic standards investigated in this study.

Class	Chemical	Molecular Formula	Ion		Ω_{N_2} (\AA^2)	Molecular Structure
			Formula	m/z		
Amine	Tetraethyl ammonium chloride	$C_8H_{20}NCl$	$[M-Cl]^+$	130.16	122.1	
	Tetrapropyl ammonium chloride	$C_{12}H_{28}NCl$	$[M-Cl]^+$	186.10	143.8	
	Tetrabutyl ammonium iodide	$C_{16}H_{36}NI$	$[M-I]^+$	242.17	165.8	
	Tetrapentyl ammonium chloride	$C_{20}H_{44}NCl$	$[M-Cl]^+$	298.35	190.0	
	Tetraheptyl ammonium chloride	$C_{28}H_{60}NCl$	$[M-Cl]^+$	410.47	236.5	
	2,4-Lutidine	C_7H_9N	$[M+H]^+$	108.08	123.4	
	2,6-Di-tert-butylpyridine	$C_{13}H_{21}N$	$[M+H]^+$	192.17	145.0	
Amino acid	L-Leucine	$C_6H_{13}NO_2$	$[M+H]^+$ $[M-H]^-$	132.10 130.09	137.8 144.4	
	D-Isoleucine	$C_6H_{13}NO_2$	$[M+H]^+$ $[M-H]^-$	132.10 130.09	135.2 140.3	
mono Carboxylic Acid	Benzoic acid	$C_7H_6O_2$	$[M-H]^-$	121.03	128.6	
	Octanoic acid	$C_8H_{16}O_2$	$[M-H]^-$	143.11	144.7	
	2-Butyloctanoic acid	$C_{12}H_{24}O_2$	$[M-H]^-$	199.17	162.0	
	Tridecanoic acid	$C_{13}H_{26}O_2$	$[M-H]^-$	213.19	166.2	$CH_3(CH_2)_{10}CH_2COOH$
	Pentadecanoic acid	$C_{15}H_{30}O_2$	$[M-H]^-$	241.22	173.7	$CH_3(CH_2)_{12}CH_2COOH$
	Palmitic acid	$C_{16}H_{32}O_2$	$[M-H]^-$	255.23	177.9	$CH_3(CH_2)_{13}CH_2COOH$
	Stearic acid	$C_{18}H_{36}O_2$	$[M-H]^-$	283.26	185.4	$CH_3(CH_2)_{15}CH_2COOH$
Oleic acid	$C_{18}H_{34}O_2$	$[M-H]^-$	281.25	186.9	$CH_3(CH_2)_6CH=CH(CH_2)_7COOH$	



<i>di/multi</i> Carboxylic Acid	Succinic acid	C ₄ H ₆ O ₄	[M-H] ⁻	117.02	124.6	
	Glutaric acid	C ₅ H ₈ O ₄	[M-H] ⁻	131.03	128.4	
	Adipic acid	C ₆ H ₁₀ O ₄	[M-H] ⁻	145.05	131.5	
	Pimelic acid	C ₇ H ₁₂ O ₄	[M-H] ⁻	159.06	134.0	
	Azelaic acid	C ₉ H ₁₆ O ₄	[M-H] ⁻	187.09	143.5	
	Sebacic acid	C ₁₀ H ₁₈ O ₄	[M-H] ⁻	201.11	148.9	
	1,2,3-Propane tricarboxylic acid	C ₆ H ₈ O ₆	[M-H] ⁻	175.02	122.2	
	Cyclohexane tricarboxylic acid	C ₉ H ₁₂ O ₆	[M-H] ⁻	215.06	135.0	
Mellitic acid	C ₁₂ H ₆ O ₁₂	[M-H ₂ O-H] ⁻	322.96	154.6		
Ester	Dibutyl oxalate	C ₁₀ H ₁₈ O ₄	[M+Na] ⁺	225.11	170.0	
	Dibutyl phtahlate	C ₁₆ H ₂₂ O ₄	[M+Na] ⁺ [2M+Na] ⁺	301.14 579.29	192.4 255.5	
	Diocetyl phthalate	C ₂₄ H ₃₈ O ₄	[M+H] ⁺	391.28	203.6	
Alcohol	Propylene glycol	C ₃ H ₈ O ₂	[2M-2H ₂ O+Na] ⁺	215.12	144.8	
			[4M-3H ₂ O+Na] ⁺	273.17	156.4	
			[5M-4H ₂ O+H] ⁺	309.23	165.7	
			[5M-4H ₂ O+Na] ⁺	331.21	169.6	
			[6M-5H ₂ O+H] ⁺	367.27	179.1	
			[6M-5H ₂ O+Na] ⁺	389.24	181.6	
			[7M-6H ₂ O+H] ⁺	425.31	190.8	
			[7M-6H ₂ O+Na] ⁺	447.28	193.9	
			[8M-7H ₂ O+H] ⁺	483.35	204.7	
			[8M-7H ₂ O+Na] ⁺	505.32	206.2	
			[9M-8H ₂ O+H] ⁺	541.39	218.5	
[9M-8H ₂ O+Na] ⁺	563.36	219.3				
[10M-9H ₂ O+H] ⁺	599.42	231.3				
[10M-9H ₂ O+Na] ⁺	621.40	231.8				



	DL-Threitol	C ₄ H ₁₀ O ₄	[M+Na] ⁺	145.05	133.0	
	Xylitol	C ₅ H ₁₂ O ₅	[M-H] ⁻	151.06	131.2	
	Sucrose	C ₁₂ H ₂₂ O ₁₁	[M-H] ⁻ [M+Na] ⁺	341.11 365.11	167.6 175.1	
	Hexane-3,4-dione	C ₆ H ₁₀ O ₂	[M+H] ⁺ [M+CH ₃] ⁺	115.08 129.09	115.7 121.3	
Carbonyl	Acetophone	C ₈ H ₈ O	[M+CH ₃] ⁺	135.08	120.4	
	Cinnamaldehyde	C ₉ H ₈ O	[M+CH ₃] ⁺	147.08	123.9	
	Levulinic acid	C ₅ H ₈ O ₃	[M-H] ⁻	115.04	130.0	
	4-Acetylbutyric acid	C ₆ H ₁₀ O ₃	[M-H] ⁻	129.06	134.5	
	Homovanillic acid	C ₉ H ₁₀ O ₄	[M-H] ⁻	181.05	147.7	
	16-Hydroxy hexadecanoic acid	C ₁₆ H ₃₂ O ₃	[M-H] ⁻	271.22	183.7	
<i>multi</i> Functional Compound	Oxaloacetic acid	C ₄ H ₄ O ₅	[M-H] ⁻	131.06	118.3	
	Ketoglutaric acid	C ₅ H ₆ O ₅	[M-H] ⁻	145.01	120.9	
	Oxozelaic acid	C ₉ H ₁₄ O ₅	[M-H] ⁻	201.08	133.3	
	Malic acid	C ₄ H ₆ O ₅	[M-H] ⁻	133.01	111.4	
	Tartaric acid	C ₄ H ₆ O ₆	[M-H] ⁻	149.01	116.0	
	Citric acid	C ₆ H ₈ O ₇	[M-H] ⁻	191.02	123.0	
	Sodium Dodecyl sulfate	C ₁₂ H ₂₅ SO ₄ Na	[M-Na] ⁻	265.15	163.6	
Organic Sulfate	Sodium Taurocholate	C ₂₆ H ₄₄ SO ₇ NNa	[M-Na] ⁻	514.28	206.4	

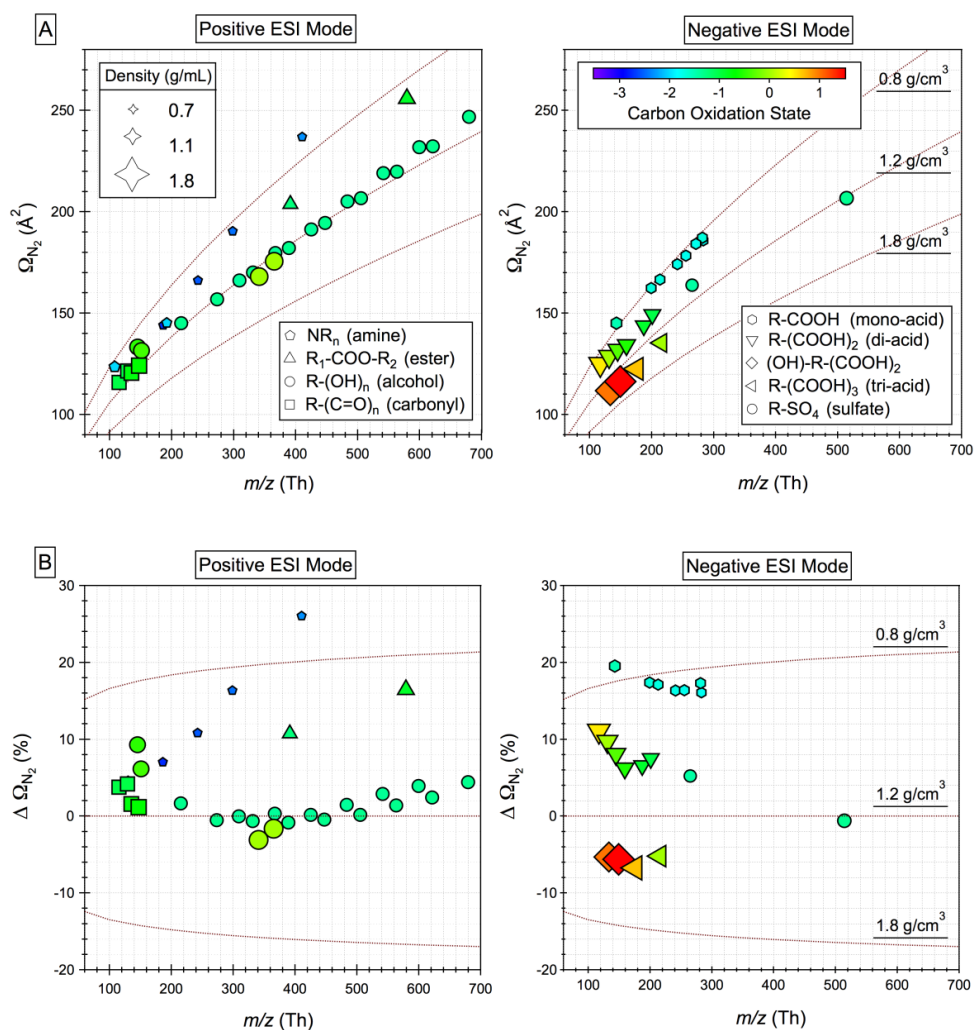


Figure 1. Distribution of organic species including alcohol (R-(OH)_n, $n = 2-8$), amine (NR₃), *quaternary*-ammonium (NR₄), carbonyl (R-(C=O)_n, $n = 1-2$), carboxylic acid (R-(COOH)_n, $n = 1-3$), ester (R₁-COO-R₂), organic sulfate (R-SO₄), and *multi*-functional compounds ((OH)-R-(COOH)₂) on the (A) $\Omega_{N_2} - m/z$ space and (B) $\Delta \Omega_{N_2} - m/z$ space.

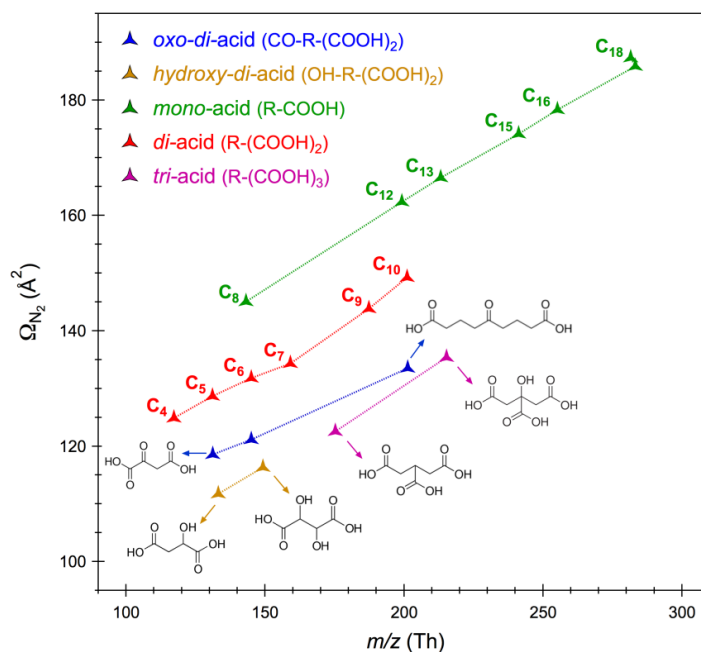


Figure 2. Trajectories associated with reactions involving functionalization (changes in the type and number of functional groups) and fragmentation (changes in the carbon chain length) through the 2-D $\Omega_{N_2} - m/z$ space using carboxylic acid series as an illustration.

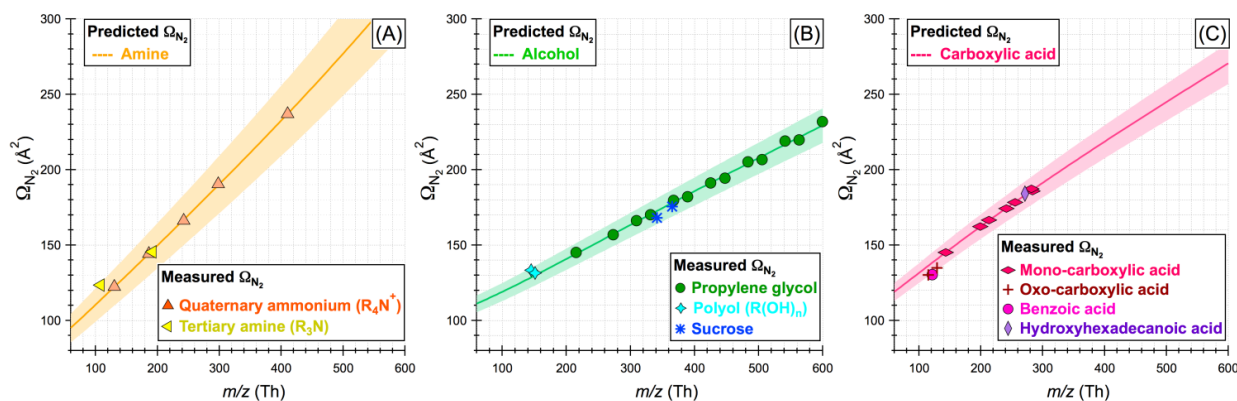


Figure 3. Measured collision cross sections (Ω_{N_2}) for (A) tertiary-amine and quaternary-ammonium, (B) (di/poly/sugar)-alcohol, and (C) (mono/oxo/hydroxy)-carboxylic acid as a function of the mass-to-charge ratio. Also shown are the predicted $\Omega_{N_2} - m/z$ trend lines for amine, alcohol, and carboxylic acid by the core model. Here, quaternary-ammonium, propylene glycol, and C_8 - C_{18} alkanolic-acid are used to optimize the adjustable parameters in the core model (The markers are in the same color as the trend lines). The colored shade in each figure represents the maximum deviations (8.21%, 3.54%, and 6.69% for amine, alcohol, and carboxylic acid, respectively) of the predicted Ω_{N_2} from the measured Ω_{N_2} for species that are not used to constrain the core model. A single plot showing the separation of these three chemical classes is given in Figure S5 in the supplement.

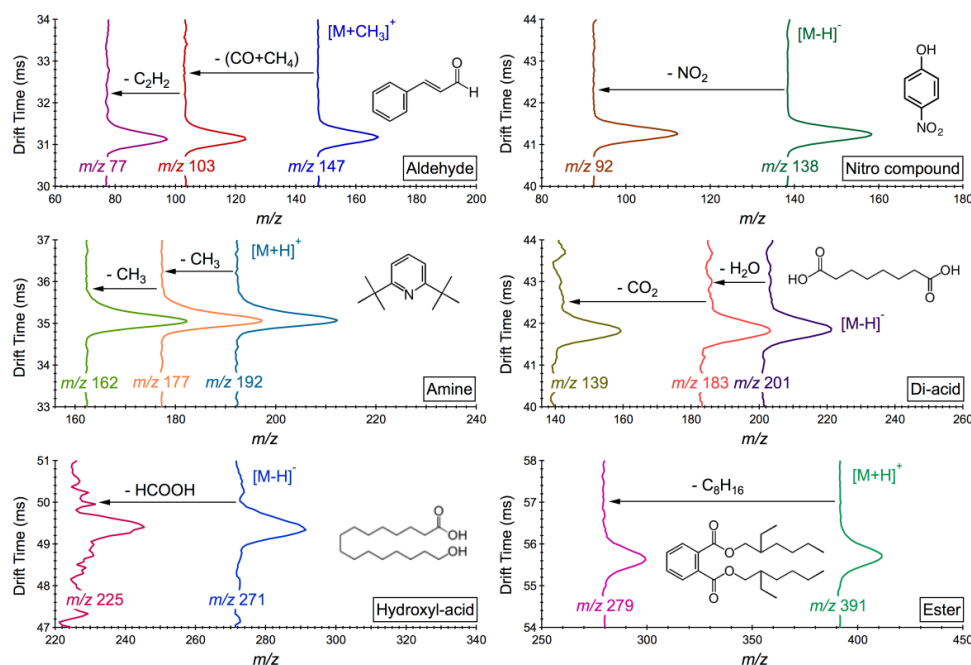


Figure 4. Collision induced dissociation patterns for molecular ions generated from cinnamaldehyde, dioctyl phthalate, 2,6-di-tert-butylpyridine, 4-nitrophenol, 16-hydroxyhexadecanoic acid, and sebacic acid on the mobility – mass framework with mass-to-charge ratio on the x -axis and drift time on the y -axis. The corresponding mobility selected MS spectra for each species is given in Figure S6 in the supplement.

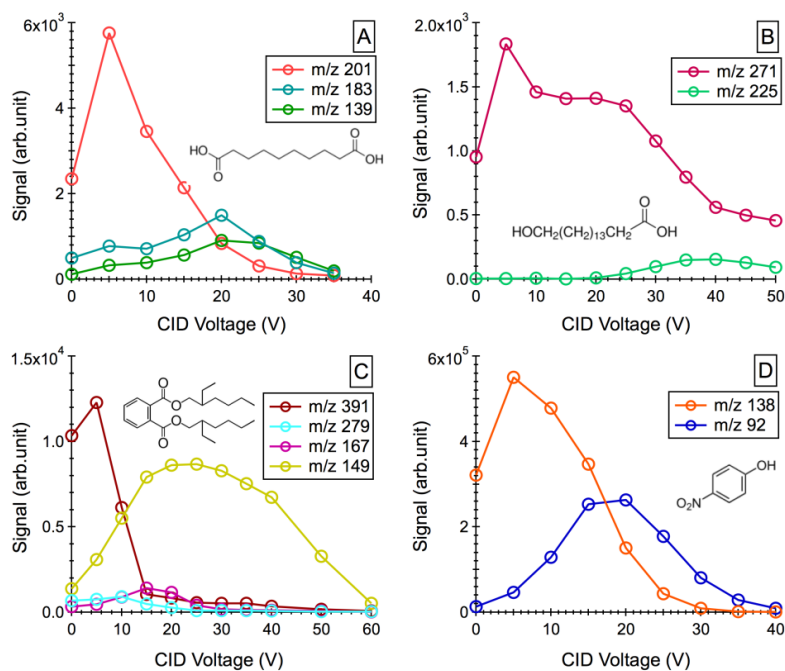


Figure 5. Product ion peak intensities as a function of collision voltage in the ‘mobility-selected’ MS spectra of (A) deprotonated sebacic acid, (B) deprotonated 16-hydroxyhexadecanoic acid, (C) sodiated dioctyl phthalate, and (D) deprotonated 4-nitrophenol.

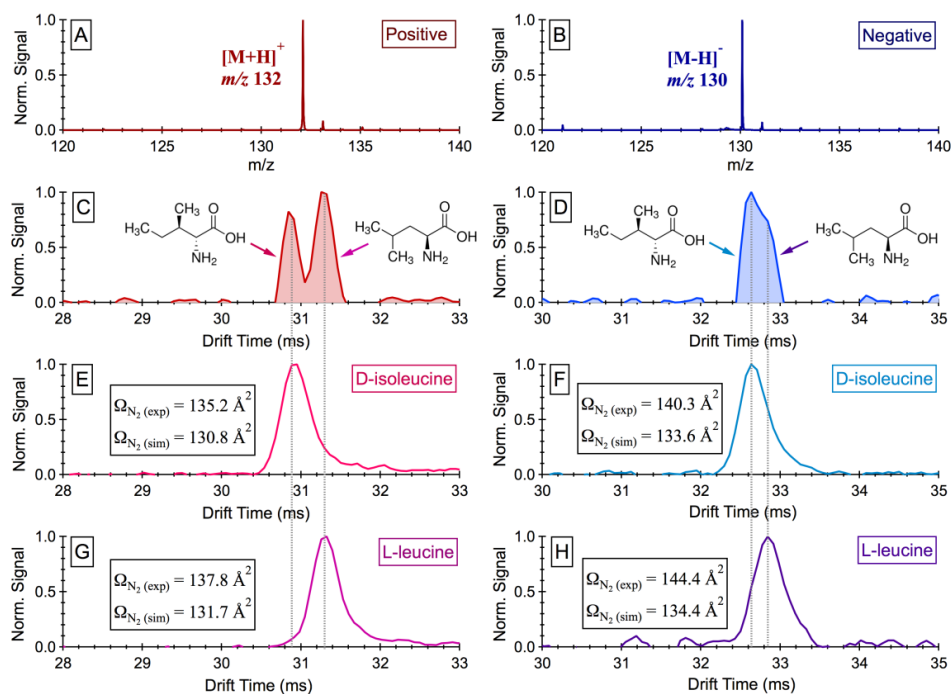


Figure 6. (A/B) ESI mass spectra collected for an equi-molar mixture (20 μM each) of L-leucine and D-isoleucine in positive and negative mode. (C/D) Measured drift time distributions for the leucine mixture in positive and negative mode. (E/F) Measured vs. predicted Ω_{N_2} (exp) for D-isoleucine, together with its drift time distributions in positive and negative mode. (G/H) Measured vs. predicted Ω_{N_2} for L-leucine, together with its drift time distributions in positive and negative mode. Note that all measurements were performed at ~ 303 K and ~ 1019 mbar with an electric field strength of 414 and 403 V cm^{-1} in the positive and negative mode, respectively.