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Structural elucidation of soluble organic matter: application to Titan's haze

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16 Abstract

The origin and evolution of organic matter in the solar system intertwines astrobiology and 17 18 planetary geochemistry issues. To observe the contribution of atmospheric processes in the 19 formation of complex organic matter, one of the most intensively studied objects in the outer 20 solar system is Titan, the largest moon of Saturn. Its reducing atmosphere of methane and 21 nitrogen hosts a thick, permanent, nitrogen-rich, organic haze, whose complex composition 22 remains largely unknown. Due to the measurement of species at large mass-to-charge ratio and infrared information from the Cassini-Huygens mission, polyaromatic hydrocarbons (PAHs) 23 24 based structure for this haze has been suggested. Here, we propose a snapshot of the global 25 chemical structure based on the analysis of laboratory analogue of Titan's haze with ion 26 mobility spectrometry coupled with mass spectrometry. This robust analysis, validated with 27 other geochemical complex mixtures, such as petroleum, allows for the observation of the size 28 and three-dimensional structure of detected species. By comparison, with standards molecules, 29 we exclude several structures such as pure PolyHCN and pure polycyclic aromatic 30 hydrocarbons to be present in the principal trend of the laboratory tholins. Using theoretical 31 calculations, we propose a plausible structure consistent with our results, which is a branched 32 triazine-pyrazole. We observe that the larger the aerosols molecules are, the more they tend 33 towards a structure containing small aromatics cores linked together by short chains. We 34 suggest that the use of such an analytical approach could help advance our understanding of 35 other complex organic compounds in the Solar System such as soluble organic matter in meteorites. 36

37 Keywords:

- 38 Titan's haze chemistry,
- 39 Ion mobility mass spectrometry,
- 40 Structural analyses,
- 41 Soluble organic matter,
- 42 Icy world,

43 Graphical Abstract



45 1. **Introduction**

Organic matter is present all over the solar system. Understanding its formation and history 46 47 remains a major challenge for the fields of astrobiology and planetary geochemistry. A key 48 approach to do so is to study the elemental composition and chemical structure of this organic 49 material. In this work, we propose to tackle the question of the chemical structure of complex, 50 soluble organic matters of interest for Earth field samples, meteorites, interplanetary dust 51 collected in the Earth's upper atmosphere, sample return from space missions and in situ future space developments oriented towards complex organic worlds such as Titan and the ocean 52 53 worlds. The case study chosen here is the challenging analysis of Titan's aerosols analogues. 54 These are reduced nitrogen-rich C_xH_yN_z organic samples synthesized in the laboratory from 55 processes simulating the formation of the atmospheric organic haze surrounding Titan. The 56 high degree of complexity of this material, with only carbon, nitrogen and hydrogen atoms, the 57 absence of oxygen, and its formation in cold conditions leads us to consider this model organic 58 material of interest not only for Titan, but for understanding the formation of primitive organic 59 material in the outer reduced solar system, for example comets (Jost et al., 2017) and the early Earth (Trainer et al., 2006). In addition, the hydrolysis of these analogues has been shown to 60 61 lead to a variety of amino acids very close to the result of the Miller-Urey experiment, making 62 it a key component for astrobiology (Neish et al., 2009; Neish et al., 2010).

These proxies have been extensively analyzed with state-of-the-art instruments, e.g., ultrahigh resolution mass spectrometers, such as FT-ICR and Orbitrap to push our understanding of this organic matter (Cable et al., 2012; Danger et al., 2016; Danger et al., 2013; Somogyi et al., 2005; Szopa et al., 2006; Toupance et al., 1975). Unfortunately, even with Earth best instruments, revealing the structure of these analogues remains a challenge. Several analyses were performed on Titan's aerosol analogues, including infrared spectroscopy (Cable et al., 2014; Gautier et al., 2012; Imanaka et al., 2004), capillary electrophoresis (Cable et al., 2014) 70 and nuclear magnetic resonance (He and Smith, 2015). These previous analyses revealed the 71 presence of molecular families such as nitriles, amines and hydrocarbons. Mass spectrometry 72 analysis revealed a great deal of structural information on laboratory tholins, including a 73 polymeric trend with CH₂ and HCN as repetition pattern (Anicich et al., 2006; Bonnet et al., 2013; Gautier et al., 2016; Somogyi et al., 2005; Somogyi et al., 2016; Vuitton et al., 2010). 74 75 Recently, mass spectrometry analyses were performed with a laser ionization desorption source 76 (Gautier et al., 2017; Maillard et al., 2018; Somogyi et al., 2012). This source allowed for the 77 comparison of both liquid and solid state of non-totally soluble tholins and showed the chemical 78 difference between these two fractions with the presence of half the amount of hydrogen in the 79 insoluble fraction (Maillard et al., 2018).

80 The main limitation of mass spectrometry analyses has been the absence of conformational and 81 isomeric information about the analyzed sample. Tandem mass spectrometry remains a solution 82 to recover structural information but is difficult to apply here due to the extreme complexity of 83 such organic samples. One way to get around this issue is to use chromatographic separation 84 (in gaseous or liquid state) upstream of the mass spectrometer to recover isomeric information. 85 The retrieved retention time is dependent of the chemical properties of the molecule and allows, 86 using standard molecules, identification of species. This technique revealed a strong potential 87 for the elucidation of several compounds such as triazole, triazine and cyanoguanidine (Gautier 88 et al., 2016). However, chromatography does not allow obtaining structural information without 89 comparison to a standard molecule and therefore remains unable to solve the structure of the 90 thousands of molecules comprising laboratory tholins.

Ion mobility spectrometry (IMS) is a gas phase separation method that can be coupled to mass
spectrometry (IMS-MS) (Mason and Schamp, 1958; Revercomb and Mason, 1975). IMS-MS
has been used recently for the analysis of complex mixtures such as petroleum (Castellanos et
al., 2014; Fernandez-Lima et al., 2009; Maleki et al., 2016). IMS is based on the separation of

95 ions in a neutral gas, according to their charge state, size and shape. Three main ion mobility techniques allow the determination of collision cross section (CCS): drift tube (May et al., 96 97 2014), Trapped Ions Mobility Spectrometry (TIMS) (Tose et al., 2018) and Travelling Waves Ion Mobility Spectrometry (TWIMS) (Fasciotti et al., 2013; Hines et al., 2016) technology. 98 99 Direct measurement of the ion collision cross section (CCS) can be done with a drift tube. For 100 TIMS or TWIMS, it is possible to determine the CCS (Campuzano et al., 2012) from the 101 experimentally determined drift time (t_D) after calibration. The CCS is an intrinsic property of 102 the ions (for a particular buffer gas) that can be compared to theoretical CCS values obtained 103 from putative tridimensional structures.

104 The following study presents the advantage of IMS-MS for the study of complex organic 105 matter, in our case analogous of the Titan's hazes, to provide a snapshot of their global structure.

106 **2. Methods**

107 2.1. Tholins production

108 Titan's aerosol analogues, also hereafter called *tholins*, were produced with the PAMPRE 109 experiment (French acronym for Aerosols Production in microgravity with reactive plasma) 110 following the same procedure detailed in previous publications (Gautier et al., 2011; Szopa et al., 2006). The reactor is composed of a stainless steel cylindrical reactor in which a RF-111 112 Capacitively Coupled Plasma discharge is established thanks to an RF 13.56 MHz frequency 113 generator. A gas mixture containing 95% of nitrogen and 5% of methane was injected in the 114 chamber as a continuous flow through polarized electrodes. It is then extracted by a primary 115 vacuum pump to ensure that gases are homogeneously distributed. The plasma discharge was 116 maintained at a pressure of 0.9 ± 0.1 mbar and at room temperature. A brown powder, called 117 tholins, was recovered after 1 day. The harvesting procedure was carried out under ambient air. 118 Tholins are then stored in an inert atmosphere and protected from light. Some oxidation occurs 119 during the harvest but these new oxygenated species can be sorted and are not studied during 120 the mass spectrometry analysis. It should be noticed that the pressure and temperature are lower in Titan's ionosphere (respectively $\sim 10^{-5}$ - 10^{-8} mbar and 200K) than in our experiment, but the 121 122 ionization rate is the same ($\sim ppm_y$). As ion-molecule reaction rates are relatively insensitive to 123 the temperature, the lower temperature in Titan's ionosphere (200K instead of 293K in the 124 laboratory) is not an important issue in our case. The higher pressure ensures a faster kinetics 125 in the experiment without being high enough to enable molecular reactions, whereas the similar 126 ionization rate enables a realistic contribution of ions into the whole ion-neutral coupled 127 chemical network.

129 2.2. Sample preparation

130 To recover the soluble part of the sample, 4 mg of laboratory tholins were dissolved in 1 mL of methanol 131 in a vial. The vial was vigorously stirred for 3 minutes to solubilize the maximum amount of species. 132 The brown mixture was then filtered using a 0.2 µm polytetrafluoroethylene (PTFE) membrane filter on 133 a filter holder. Filtered solution was transferred in a vial and then analyzed after half dilution with a 134 50/50 water/methanol mixture to be analyzed under the same conditions than previous studies by 135 electrospray ionization. Based on electrospray mass spectrometry and elemental analysis, we 136 have evidenced the presence of less than 10% of oxygen containing species. Oxidation occurred 137 most likely during sample harvesting and storage but we cannot rule out that some oxidation 138 might occur also during ionization. The oxygen containing species have been discarded for data 139 treatment. It must be notified that around 40% of the initial quantity of laboratory tholins is 140 soluble in methanol, so the following study will only focus on a part of the sample and will not 141 be fully representative of the entire sample (Carrasco et al., 2009).

142 2.3. Chemicals

Methanol (LCMS grade) was purchased from Fisher scientific. All other chemical products (tetra-alkylammonium, polyglycine and drug-like compounds) were purchased from Sigma Aldrich.

145 2.4. Ion mobility - mass spectrometry experiments

146 Laboratory tholins were analyzed in positive ion mode on a hybrid quadrupole-time of flight 147 mass spectrometer equipped with a Travelling Wave Ion Mobility cell (TWIMS) and 148 electrospray ion source (Synapt G2 HDMS, Waters, Manchester, UK). Although it has been 149 shown that different species ionize in negative ionization mode (Somogyi et al., 2012), it was 150 decided, in this work, to focus on the positive mode in order to present an introduction to ion 151 mobility analysis. Following ionization parameters were applied: For the source, the capillary 152 was set at 3 kV, the temperature at 100 °C, the Sampling cone at 25 V. The extraction cone was also at 5 V and the desolvation temperature at 250 °C. Nitrogen cone gas flow was set at 10 153

154 L/Hour and nitrogen desolvation gas flow at 400 L/Hour. For Ion mobility parameters, helium 155 cell gas flow was set at 180 mL/min; Nitrogen gas flow was set at 90 mL/min, the IMS wave 156 velocity at 400 m/s. and the IMS wave height at 13 V. Trap wave velocity was set at 311 m/s 157 and Trap wave height at 6 V. Finally, transfer wave velocity was set at 191 m/s and transfer 158 wave height at 4 V. The m/z values in recorded spectra were first externally calibrated using 159 sodium formate and then internally calibrated using several well-known ions (Maillard et al., 160 2018). Drift times were extracted for each species using DriftScope 2.8 and MassLynx 4.1. 161 Mobility dimension was calibrated using tetraalkylammonium ions (TAA) (Campuzano et al., 162 2012) with reference values in helium. Even if the instrument is working under nitrogen, it is 163 commonly used to calibrate the ccs values in helium. Indeed, it was proven that for small 164 molecules, there is no bias in the calibration (Bleiholder et al., 2015). Therefore, every 165 recovered laboratory tholins CCS is calculated in the same gas. Calibration peaks were fitted 166 with a Gaussian shape using Origin 2016. The resulting uncertainty after calibration for cross sections was estimated at ± 0.8 Å² by taking into account the error of the fitting (Figure S3). 167 168 Consequently, CCS for laboratory tholins will be stated without decimal. Table S2 shows the 169 comparison between experimental measurements and database CCS values in order to validate 170 our calibration. All reported CCS values are given following the recommendation of the ion 171 mobility community in terms of rating, calibration and results reporting (Gabelica et al., 2018).

Due to the resolution of the time of flight mass analyzer (m/ Δ m 40 000), we choose to focus on species below *m/z* 250 to prevent isobaric interferences and misassignments (See Figure S1-S2). We have shown that at this mass range, detected species are similar in the soluble and the insoluble fractions (Maillard et al., 2018).

176 2.5. Collision Cross Section calculation using theoretical model

177 From two-dimensional structures of chosen references ions, three-dimensional structures were

178 geometrically optimized (including partial charges) with Avogadro 1.1.1 using MMFF94 force

179 field, 10,000 steps, a steepest descent algorithm and a convergence of 10e⁻⁷. Conformation
180 search was performed when molecules were containing chiral center.

181 Theoretical calculation was achieved with MOBCAL (Mesleh et al., 1996; Shvartsburg and 182 Jarrold, 1996) with the trajectory method following Lennard-Jones parameters (Campuzano et 183 al., 2012) : hydrogen with atomic energy at 0.6175 meV and van der Waals distance at 2.2610 Å. carbon with atomic energy at 1.3266 meV and van der Waals distance at 3.0126 Å and finally 184 185 nitrogen with atomic energy at 1.4740 meV and van der Waals distance at 3.3473 Å. Each 186 calculated species were obtained thanks to 400,000 points (10 cycles. 1000 points in Monte 187 Carlo integrations of impact and 40 points in velocity integration.). All calculations were 188 performed considering helium as buffer gas. Helium was chosen because it is a much easier gas 189 to model than nitrogen.

191 **3. Results**

192 *3.1. IMS-MS experiments*

193 Two pieces of observable information are recovered from the analysis of complex organic 194 matter by IMS-MS: first, the usual mass-to-charge (m/z) dimension, and then the drift time 195 dimension. The second dimension corresponds to the time taken by the ions to get across the 196 mobility cell (in practice, it includes a short period of time spent by the ions in the ion optics to 197 reach the detector). This time is characteristic of the ion size, shape and charge in the gas phase. 198 The resulting spectrum is given in Figure 1. This three-dimensional map allows a first overview 199 of the analyzed sample. The major sequence (blue area) corresponds to singly charged species. 200 In the case of laboratory tholins, a typical wave-formed structure is recovered (Bonnet et al., 201 2013; Gautier et al., 2016; Pernot et al., 2010) with a succession of high intensity clusters 202 characteristic of known repetition patterns such as CH₂ and HCN (Somogyi et al., 2016). The 203 minor sequence below the major one on the map (green area) is composed of doubly charged 204 species naturally separated from singly charged species due to the ion mobility. These doubly 205 charged species are centered around m/z 275, which represents molecules of mass 550 u. Their 206 presence is not surprising because laboratory tholins contains a large amount of nitrogenous 207 compounds (Sciamma-O'Brien et al., 2010), which easily allow double protonation. These 208 double charged species will not be discussed in this study. A third sequence (white area) is 209 observable on Figure 1 far below the doubly charged one. This represents the larger singly 210 charged species collected from the previous scan. Indeed, these species would need a longer 211 time to be expelled from the mobility cell than the time stated for the next ion packet to be 212 released in the cell. This phenomenon is often called "wrap-around". It could have been avoided 213 by changing parameters, such as wave velocity and wave height, but the used experimental 214 conditions were chosen to optimize the ion mobility separation in the desired drift time range. 215 In this study, we focused on the separation of ions between m/z 50 and m/z 250, as discussed in the methods section. Therefore, we optimized the drift time separation for this m/z range. The following results will focus on the extracted peaks in the major sequence. Their collision cross section (CCS) are recovered using tetra-alkylammonium salt (TAA) ions (Campuzano et al., 2012) as reference for CCS calibration in helium.





Figure 1: CCS vs m/z plots obtained from laboratory tholins recorded on Synapt G2 (intensity in logarithm scale). Three areas are surrounded: (blue) main trend corresponding to singly charged species, (green) secondary trend corresponding to doubly charged species, (white) wrap around phenomenon corresponding to low mobility singly charged species coming from the precedent scan.

227 All recovered peaks are presented in Figure 2, which provides CCS in helium vs mass-to-charge 228 ratio for each detected peak. With this overview, we obtain the general trend for our sample. 229 To have a snapshot of the global structure of laboratory tholins sample, we added two well-230 known families from our experimental measurements: tetra-alkylammonium ions (TAA), 231 which were used as reference CCS, and polyglycine. In addition, polycyclic aromatic 232 hydrocarbon was added from precedent measurements (PAH) (Campuzano et al., 2012; Lim et 233 al., 2018). All these families are listed in the table S2 and a comparison is given between 234 experimental measurements and database CCS in order to validate our results.

Furthermore, two other families are described using theoretical calculation: linear and branchedpolyHCN (See Figure S4 for detailed formulas of plotted species).

237 PAHs represent a family of planar compounds containing only carbon and hydrogen. This 238 family is generally used in DBE vs. carbon number plot because it is the most unsaturated 239 known family (Cho et al., 2011) and was suspected to be present in Titan's aerosols and in meteoritic organic matter (Waite et al., 2007). Due to the planar conformation of PAH 240 241 molecules, it is interesting to observe the trend of the analyzed sample in comparison to this 242 family. It can be noticed that, even if the amount of heteroatoms is not the same between 243 laboratory tholins and PAHs, the PAH line remains under all detected species of laboratory 244 tholins. This means that the planar conformation, and so PAHs, does not seem to be the 245 preferred structure of our sample at this m/z range. TAA family is composed of molecules 246 containing an ammonium core with four carbon chain arms. This group, in opposition to PAHs, 247 tends to form spherical conformations with the folding of the carbon chains. This line remains 248 above the laboratory tholins one, leading to the conclusion that this specific spherical 249 conformation is not the one preferred in our sample. Finally, polyglycine was also considered 250 as possible laboratory tholins structures. Polyglycine is a peptide, which tend to form a helical 251 structure according to our 3D optimization (See Figure S5a for a structure). The shorter Gly2 252 peptide presents a CCS close to that of tholins but larger peptides present a more compact 253 structure. Small polyglycines are not long enough to form a helical structure. This explain why 254 they fit with the tholins sample. As a short summary of the comparison of these tree families 255 with our sample, the preferential conformation present in these laboratory tholins seems to be 256 seems to be none of the three observed conformations. Two ideal PolyHCN polymers were 257 added thanks to calculated CCS as presented in the next section. The linear PolyHCN can be 258 assimilated to a straight stick shape (3D structure in figure S5b). The branched one form a non-259 perfectly flat network (3D structure in figure S5c). These 3D structures are also given in figure

S5b and S5c. This family has been proposed to be the one composing Titan's aerosols. In
agreement with Vuitton *et al.*, 2010 (Vuitton et al., 2010), laboratory tholins are clearly not pure
ideal polyHCN because representative lines of this family are far above the detected sample.
We can then exclude this structure from our measurement. They are potentially present in small
quantities in the sample but not detected in these measurements.

In this first overview, we have recovered important information concerning the global structure
of laboratory tholins sample. Spherical and pure planar shapes were excluded as well as helical
structures.



268

Figure 2: Comparison of collision cross section (A²) *vs m/z* of ions of laboratory tholins sample
with experimental measurements (*), polyglycin (purple) and tetraalkylammonium salts (red).
Taken from database (+), polyaromatics hydrocarbons (black) and calculated (o) polyHCN
(dashed lines)

A middle range zoom, described in Figure 3a, allows for the observation of four clusters between m/z 120 and m/z 180. Ion mobility spectrometry naturally acts as a Kendrick mass defect diagram (Hughey et al., 2001; Kendrick, 1963). In our case, two main repetition patterns are recovered: CH₂ and HCN. This result is consistent with previous studies (Anicich et al., 2006; Somogyi et al., 2005).

279 A short-range zoom is given in Figure 3b. This sub-figure focuses on a single cluster located 280 between m/z 134 and m/z 144. All molecular formulas are determined for the detected species, 281 allowing the direct observation of the collision cross section evolution according to the 282 compounds formula. Two main areas are shown according to the number of nitrogen atoms in 283 each species. The first region (in blue) is located at the top of the cluster. It contains ions with 284 a lower amount of nitrogen atoms (between N2 and N3). The other region (in red), which 285 contains ions with a higher nitrogen content (between N4 and N7) is located under the first one. 286 Therefore, the number of nitrogen atoms plays a major role in the conformation of our organic 287 matter samples and also allows the production of different isomers that are not separated with 288 the current instrumental resolution. The collision cross section decreases with an increasing 289 number of nitrogen atoms. Such evolution of polymeric species as a function of the number of 290 polar heteroelements has been observed previously (Farenc et al., 2017; Woods et al., 2004). It 291 suggests that laboratory tholins molecules have some flexibility that can yield to partial folding 292 as the number of heteroelements increase by the formation of intramolecular hydrogen bonding. 293 This overview allowed the study of the global molecular structure of the laboratory tholins

294 sample. Several typical structures (such as polyHCN, PAH, TAA and polyglycine) are excluded 295 without doubt from our sample due to the difference in the CCS values at this m/z range. After 296 working by exclusion, the next step is to use the CCS calculation tool to propose possible 297 structures of laboratory tholins.



Figure 3: Middle range zoom of the figure 2 for detailed comparison of collision cross section (A²) *vs m/z* of laboratory tholins ions. (a) Close zoom of the figure 2 on clusters between m/z120 and m/z 170, b) Zoom on one cluster between m/z 134 and m/z 144 with molecular formula.

302

303 *3.2. CCS calculation of potential structures*

304 Due to the actual limited knowledge of such organic material molecular structures, finding a 305 family of standard compounds that would fit the entire range of collision cross sections can be 306 difficult. To solve this problem, CCS calculation is the easiest way to probe possible structures. 307 Theoretical CCS in helium are obtained using MOBCAL software (Mesleh et al., 1996; 308 Shvartsburg and Jarrold, 1996) modified with new Lennard-Jones parameters (Campuzano et 309 al., 2012). The calculation method is first trained on standards to ensure its robustness (See 310 table S2 for further details). CCS values of two series of compounds are represented on Figure 311 4. The blue curve presents a family of compounds based on a triazine core substituted with 312 different numbers of branches. These branches are composed of an imine connected to a carbon 313 chain. The green curve shows a family based on pyrazole cores connected with imine groups. 314 Triazine and pyrazole cores have been chosen because their presence has been previously 315 identified in laboratory tholins (Gautier et al., 2016; Quirico et al., 2008). As observed, the CCS 316 of these two set of compounds corresponds to the CCS of tholins compounds. Thus, these two 317 families could be consistent with laboratory tholins structure, in opposition with compounds 318 presented in the previous section. It should be pointed out that several structures can match a 319 particular CCS value, the structures presented here are likely hypothesis in agreement with 320 laboratory tholins.



Figure 4: Comparison of CCS *vs m/z* with (blue) calculated CCS of triazine family (green)
calculated CCS of pyrazole family.

325 **4. Discussion**

326 Triazine and pyrazole were identified in laboratory tholins by NMR studies (He and Smith, 327 2014) and liquid chromatography coupled to high resolution mass spectrometry (Gautier et al., 328 2016). The first described triazine family in this study was based on these precedent works 329 (Gautier et al., 2016; He and Smith, 2014) and seems to fit well with the global trend of laboratory tholins samples between PAHs and TAA families. However, the CCS values of this 330 331 family correspond to the highest CCS values of laboratory tholins typical ions. In opposition, 332 the second family based on a pyrazole core described in precedent works have CCS values in 333 agreement with the lowest CCS values for the laboratory tholins species. We propose, regarding 334 these results, a structure based on a Nitrogenated-PAH cores (Triazine and pyrazole) linked 335 together with short chains. The core of this set is completely conjugated. This would induce a 336 strong absorption in the UV/visible, consistent with Titan's aerosols color. A growing path 337 perspective of this family is given in Figure 5. This hypothetical structure is consistent with 338 recent works that highlighted the wide diversity of production pathways based on aromatic 339 starting cores (Gautier et al., 2017; Mahjoub et al., 2016).

340 Regarding Titan, it was proposed that the aerosols should have a PAHs structure based on 341 several observations obtained thanks to the Visual and Infrared Mapping Spectrometer (VIMS) 342 boarded on the Cassini probe (Dinelli et al., 2013; López-Puertas et al., 2013). This scenario 343 was largely approved because the aromatisation of small compounds is usually favoured due to 344 their strong stability. While our results agree with the presence of aromatic rings in the aerosol 345 structure, we observe that a pure PAH structure cannot fit the aerosol CCS. However, an 346 aromatic structure based on N-PAH cores could still be in agreement with VIMS observation 347 and would fit the structure of studied laboratory tholins sample much better.



Figure 5: Growing path possibility of families containing triazine and pyrazole cores. As example here, a compound with raw formula $C_{52}H_{78}N_{25}$ Note: This structure by itself was not

- detected directly in the laboratory tholins but represents a general idea of molecules that could
- be embedded in them.
- 353
- 354

5. Conclusions

356 In summary, this work introduces ion mobility spectrometry coupled with mass spectrometry 357 analysis for the structural study of soluble organic matter by presenting a case study on 358 analogues of the Titan's haze. This new analysis allows for an isomeric separation of the 359 sample. In addition, the collision cross section, additional structural information, is determined 360 for each ion. The collision cross sections give an overview of the main structural shape of the 361 organic matter present in the sample. Moreover, as this information is intrinsic (for a certain 362 gas) and predictable, the calculation of theoretical CCS is possible for several series of 363 compounds. Our method allows the exclusion of several structures, in the case of laboratory 364 tholins such as pure polyHCN, PAH, polyglycine and tetra-alkylammonium salts at this range 365 of mass-to-charge ratio. These are definitely not present in the major structure of laboratory 366 tholins. To conclude, we proposed two families composed of small aromatic cores linked 367 together with short chains that could be potentially be present in laboratory tholins sample. We 368 suggest using the methodology detailed here to study other material containing complex organic 369 matter, such as meteoritic soluble organic matter or Earth's kerogens. We would like to point 370 out that this first introduction provides an opportunity for further studies to complete this 371 research, including the study of negative ions as well as the insoluble fraction of tholins, both 372 proved to be an important part of the material (Maillard et al., 2018; Somogyi et al., 2012).

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Supplementary Information for

Structural elucidation of soluble organic matter: application to Titan's haze

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This PDF file includes: Figs. S1 to S5

- Tables S1 to S2



Figure S1: Comparison between Synapt G2 resolving power and FTICR at m/z 181. All species are resolved with both analyser.



- **Figure S2:** Comparison between Synapt G2 resolving power and FTICR at *m/z* 301. Above *m/z* 250,
- 579 tholins species are not resolved with the Synapt G2 analyser.

	CCS _{He} (А	Ų) M	charge	t _D (ms)	Ω'
Methyl	48.5	74.0970	1	5.18	94.51
Ethyl	65.9	130.1596	1	7.43	129.86
Propyl	88.9	186.2222	1	10.65	175.98
Butyl	111.2	242.2848	1	14.4	220.66

Table S1: TAA calibrant CCS values. Drift time was measured for each ion and added in the table.



Figure S3: Calibration curve for the tetraalkyammonium salts in helium with a power fitting.







Tetraalkylammonium salts (CCS calibrant)							
	Experimental CCS (Å ²)		Calculated CCS (Å ²)				
	Literature	This work					
Methyl	49ª	49	48				
Ethyl	66ª	66	66				
Propyl	89 ^a	89	90				
Butyl	111ª	111	114				
Polyc	yclic Aromatic H	lydrocarbons					
Benzene	/	/	45				
Naphtalene	59ª	/	59				
Antracene	72ª	/	73				
dibenz[A,H]anthracene	98 ^c	/	100				
Polyglycines							
Glc ₂	62 ^b	64	60				
Glc₃	76 ^b	76	78				
Glc ₄	86 ^b	87	88				
Glc₅	97 ^b	/	101				
Drug-like compounds							
N-Ethylaniline	63ª	63	62				
Acetaminophen	67ª	69	64				
Alprenolol	97 ^a	98	98				
Ondansetron	106ª	109	108				
a : Campuzano et al. Anal. C	hem. 2012,						

Table S2: Validation of calibration parameters

b: Wyttenbach et al. J. Am. Chem . Soc. 1998

c: Dongwan et al. Bull. Kor. Soc. 2018