Dissolved organic matter in sea spray: a transfer study from marine surface water to aerosols

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Abstract

Atmospheric aerosols impose direct and indirect effects on the climate system, for example, by adsorption of radiation in relation to cloud droplets size, on chemical and organic composition and cloud dynamics. The first step in the formation of primary marine aerosols, i.e., the transfer of dissolved organic matter from the marine surface into the atmosphere was studied, and we present a molecular level description of this phenomenon using high resolution analytical tools (Fourier transform ion cyclotron resonance = FT-ICR MS and NMR). We could experimentally confirm the chemo-selective transfer of natural organic molecules, especially of aliphatic compounds from the surface water into the atmosphere via bubble bursting processes. Transfer from marine surface water to the atmosphere involves a chemical gradient governed by the physico-chemical properties of the involved molecules when comparing elemental compositions and differentiating CHO, CHNO, CHOS and CHNOS bearing compounds. Typical chemical fingerprints of concentrated compounds were CHO and CHOS type of molecules, smaller molecules of higher aliphaticity and lower oxygen content and typical surfactants. A non-targeted mass spectrometric analysis of the samples showed that many of these molecules correspond to homologous series of oxo-, hydroxyl-, methoxy-, branched fatty acids and mono-, di- and tricarboxylic acids as well as monoterpenes and sugars. These surface active biomolecules were preferentially transferred from surface water into the atmosphere via bubble bursting processes to form a significant fraction of primary organic aerosols. This way of production of sea spray leaves a specific biological signature of the surface water in the corresponding lower atmosphere that can be transported laterally in the context of global cycling.

1 Introduction

On a large scale, sea spray, which transports dissolved gases, salts, and biological materials from the sea to the atmosphere with resulting climatic impact, is largely attributed to aerosols produced by an estimated $10^{18}$ to $10^{20}$ bubbles that rupture every second
across the oceans (Woodcock et al., 1953; MacIntyre, 1972; Wu, 1981). Marine aerosols account for the majority of the global natural aerosol flux and consequently may have a significant impact on the Earth’s radiative balance and biogeochemical cycling (O’Dowd et al., 2004; O’Dowd and de Leeuw, 2007).

 Preferential enrichment of surfactants at the air-water interface occurs as a result of the amphiphilic properties of surfactants, with the water-soluble moiety plunging into the solution and the hydrophobic component in contact with the air. Concentration of specific surfactant materials – in the sea-surface microlayer and in atmospheric aerosols – has been well described (Barger and Garret, 1970; Mukerjee and Handa, 1981; Blanchard, 1990; Tseng, 1992; Oppo et al., 1999; Ekström et al., 2010). Adsorption of surfactants is even considerably increased at the sea surface during rough sea conditions, when wave breaking action causes air bubbles to be trapped beneath the water surface (O’Dowd and de Leeuw, 2007). Bubbles trapped in the liquid bulk considerably increase exchange surfaces between the sea bulk and the atmosphere. Bubbles drag surfactants along their way through the liquid bulk, reach the sea surface, to finally burst and eject aerosol droplets into the atmosphere; the principle is similar to adsorptive bubble separation in foaming systems (Backleh-Sohrt et al., 2005). Since these droplets formed from the sea surface and immediate subsurface water layers are enriched with surface active compounds, aerosols emitted into the atmosphere are also enriched with surface active materials compared to the liquid bulk. Air bubbles trapped during rough sea conditions were found to increase surfactant concentrations in aerosols by several orders of magnitude compared with those found in the liquid bulk (O’Dowd and de Leeuw, 2007). Recently, Bird et al. (2010) proved that, by rupturing, a single large bubble (with several centimeters in diameter) can fold and entrap air as it retracts, thus leading to the creation of a ring of small daughter millimetric bubbles which burst in turn (Bird et al., 2010). This phenomenon is believed to considerably increase the inner active surface and efficiency of aerosol dispersal.

From a conceptual point of view, the situation found in glasses poured with champagne or sparkling wine is very similar to that described above. As soon as a bottle of
champagne or sparkling wine is uncorked, the progressive release of CO$_2$ and other dissolved gas molecules is responsible for bubble nucleation, the so-called effervescence process (Liger-Belair, 2003). Once champagne is poured into a glass, bubbles nucleated on the glass wall drag champagne surfactants along their way through the liquid bulk (Liger-Belair et al., 2000). The formation of adsorption layers of amphiphilic macromolecules at the air/champagne interface was evidenced through ellipsometry and Brewster angle microscopy (BAM) experiments (Péron et al., 2001, 2004).

Generally speaking, bubbles bursting at a liquid surface eject two kinds of aerosols (Resch et al., 1986 and references therein): (i) droplets called film drops, formed while the film of the emerged bubble-cap disintegrates, and (ii) droplets formed by the collapse of the bottom of the bubble, called jet drops (see Fig. 1). Depending on its size, a bubble can produce up to ten jet drops, and several hundreds of film drops. Nevertheless, it was shown that bubbles with a diameter of less than about 2 mm produce no film drops as they burst (Resch et al., 1986). In the case of marine waters, bubble diameters rarely exceed about 1 mm as they reach the liquid surface and it can be concluded that only jet drops constitute the very characteristic cloud of aerosols found above the liquid surface, as revealed through high-speed micrography (Liger-Belair et al., 2001).

Accordingly, the aim of this study of marine aerosol formation was the qualitative molecular level characterization of the aerosol produced by bubbles collapsing in a specifically-designed bursting chamber and its comparison with both the original sea surface water and the organic aerosol (particle size PM$_{10}$) collected on the deck of the research vessel. The transfer of the organic signature from surface water into the burst event and the atmospheric aerosol in general was followed over all samples of the cruise and on a specific open ocean site near the Angola Basin (3.1° E, 17.7° S). The focus was on the characterization with Fourier transform ion cyclotron resonance spectrometry (FT-ICR MS) and nuclear magnetic resonance spectroscopy (NMR) of the specific molecular structures involved in the preferential transfer and the corresponding biomolecules involved leading to the formation of the marine primary organic aerosol.
2 Materials and methods

2.1 Study sites and sampling

Water samples were collected daily on the research vessel Polarstern during the expedition ANT-XXV/1 between Bremerhaven, Germany and Cape Town, South Africa from 3 to 29 November 2008 (Koch and Kattner, 2011). The selected day sample for surface water, sea spray generation and day aerosol was chosen at the Angola Basin (27 November 2008 at 3.1° E, 17.7° S) as in Hertkorn et al. (2011). During the expedition, no heavy rain showers occurred in the Intertropical Convergence Zone (ITCZ) although intense rainfall is common in this region; thus the collection of surface water with the fish sampler generated representative non diluted surface water for analysis and the generation of burst in the laboratory on board (Fig. 2). The water was filtered through pre-combusted (4 h, 450°C) GF/F filters (Whatman) as also described by Flerus et al. (2011) prior further solid phase extraction (SPE) and bursting.

2.2 Bubble bursting chamber

The sea spray was generated by bursting the surface water directly after filtration in the laboratory on the research vessel with an own build device (Fig. 2). The bubble bursting chamber was initially built to enable the bursting of continuous flow of fresh water introduced in counter flow but was used here in a static mode with around 1500 ml freshly filtered seawater each day. The number, size, and velocity of jet drops produced during bubble collapse depend on both the size of the initial bursting bubble and some liquid properties such as its surface tension, density and viscosity (Spiel, 1994, 1995, 1997); it is to be noted here that the bubble size and the foaming/bursting behavior of the sea water in the constructed vessel was very similar to the one with champagne in our previous study (Liger-Belair et al., 2009).

Briefly, the jet drops were collected on a glass surface on the top of the vessel oriented in a way to drop the liquid into a collecting vial at the corner. After 2 h of sampling
the glass surface, also containing dried jets, was washed with distilled water into the collecting vial. This final solution was desalted and concentrated using SPE as described below.

2.3 Solid phase extraction (SPE)

FT-ICR MS measurement requires the concentration and desalting of marine dissolved organic matter (DOM) prior to infusion analysis. Therefore DOM was extracted on board using reverse phase solid phase extraction (SPE). The filtered samples were acidified with HCl (hydrochloric acid, p.a. grade, Merck) to pH 2. The DOM was extracted using C18-SPE cartridges (100 mg/1 ml Bakerbond SPE columns, J. T. Baker, Netherlands). The cartridges were rinsed with 1 ml methanol (LiChrosolv; Merck), followed by 1 ml acidified ultra-pure water (1.25% formic acid) for conditioning and then the water samples were passed through each cartridge by gravity. Subsequently, remaining salt was removed with 1 ml acidified ultra-pure water. DOM was eluted with 5 ml methanol and stored in Eppendorf vials at −20°C. The surface water DOM was extracted with SPE out of 1000 ml water samples, and the burst DOM with SPE from all the volume collected in the vials (around 10 ml).

2.4 Aerosols collection and water soluble organic matter (WSOM) extraction

Particle sampling: Fine (dp < 10 µm, PM_{10}) aerosol samples were collected on circular quartz fibre filters (4.5 cm diameter, fibre glass; Whatman, pre-combusted 4 h at 450°C) with a low volume sampler (constructed by Ingenieurbüro SCHULZE, 21271 Asendorf, Germany) as described previously in Piel et al. (2006). The collector was placed at the front of the vessel at the upper deck (20 m above the sea). Approximately 60 m³ were collected every 12 h, and the day and night samples were sampled separately. Only day filters were considered in this study. The filters were immediately placed in small Petri dishes and stored at −20°C prior to their extraction at the Helmholtz Center in Munich as described in a previous work on aerosol (Schmitt-Kopplin et al., 2010).
SPE-C18. The quartz filters (1 cm² punched with the cap of 2 ml Eppendorf vials) were extracted in slight alkaline conditions (1.25 % NH₃). Subsequent SPE was made using 100 mg Bakerbond C18 cartridges (J. T. Baker, Netherlands). The cartridges were conditioned with 1 ml of methanol and 1 ml of water acidified with formic acid (1.25% formic acid). After application of 1 ml of sample (acidified with 50 µl formic acid), the columns were washed with the acidified water. The compounds were eluted with 1 ml methanol and were then ready for infusion analysis.

### 2.5 FT-ICR mass spectrometry

Ultrahigh-resolution mass spectra were acquired on a Bruker (Bremen, Germany) APEX 12Qe Fourier transform ion cyclotron resonance mass spectrometer (FT-ICR MS) equipped with a 12 Tesla superconducting magnet and an APOLLO II electrospray source. Samples were introduced into the microelectrospray source (Agilent sprayer) at a flow rate of 120 µl h⁻¹ with a nebulizer gas pressure of 20 psi and a drying gas pressure of 15 psi (250 °C). Spectra were externally calibrated on clusters of arginine (2.87 × 10⁻⁵ mol l⁻¹) and accuracy reached values below 0.1 ppm in day-to-day measurements. The spectra were acquired with a time domain of 4 megawords in a mass range of 100–2000 m/z. Before Fourier transformation of the time-domain transient, a sine apodization was performed. 1024 scans were accumulated for each sample.

FT-ICR mass spectra with m/z from 150 to 2000 were exported to peak lists at a signal to noise of 2. Possible elemental formulas were calculated for each peak in batch mode by the FormCalc software tool written in-house and with the NetCalc network approach described previously (Tziotis et al., 2011). The NetCalc software uses the mass difference information obtained from the FT-ICR mass spectra and a list of pre-chosen molecular transformations in order to reconstruct three compositional networks corresponding to surface, aerosol, and burst samples, respectively. Those networks, which are rich in structural information, are then combined into a single network with the purpose of detecting the common masses between the three systems and visualising their interactions. The NetCalc algorithm performs a network-based formula calculation.
on each of those networks. Out of the newly-found elemental compositions, further network analysis is illustrated via the compositional frequency histograms. The generated formulas were validated by setting sensible chemical constraints (N rule, positive double bond equivalent, O/C ratio ≤ 1, H/C ratio ≤ 2n + 2 element counts: C ≤ 20, H ≤ 30, O ≤ 6, N ≤ 5, S ≤ 1). The software uses theoretical isotope scoring in conjunction with an elemental composition calculator to score comparison of experimental and theoretical isotope patterns. From the lists of m/z, lists of final formulas (from 1500 to 3000 hits) could be generated and classified in containing CHO (blue), CHNO (orange), CHOS (green) or CHNOS (red) molecules.

The Masstrix web-interface (http://www.MassTRIX.org) described by Suhre and Schmitt-Kopplin (2008) was used for the annotation of selected exact masses to organism metabolites based on the KEGG and LipidMap databases for the characterization of the biosignatures.

2.6 Nuclear magnetic resonance spectroscopy (NMR)

NMR spectra of methanolic aerosol extracts were acquired with a Bruker Avance NMR spectrometer at 800.35 MHz ($B_0 = 18.8$ T) at 283 K from a few µg of solid obtained by evaporation of original methanol-h$_4$ solution. Proton NMR spectra were acquired in approx. 70 µl CD$_3$OD (Merck, 99.95% ²H) solution with a 5 mm z-gradient $^1$H/$^13$C/$^15$N/$^31$P QCI cryogenic probe in sealed 2 mm Bruker MATCH tubes. 1-D $^1$H NMR spectra were recorded with standard presaturation to attenuate residual water present (90° excitation pulses $^1$H: 10 µs, 5 s acquisition time, 5 s relaxation delay, typically 256–512 scans, 1 Hz exponential line broadening). A phase sensitive, gradient enhanced COSY NMR spectrum with solvent suppression (cosydfgpph19) was acquired with a Bruker Avance NMR spectrometer at 500.13 MHz ($B_0 = 11.7$ T) at 283 K with an acquisition time of 1 s, a relaxation delay of 0.5 s, a spectral width of 6493 Hz (12.5 ppm) with time proportional phase increment (TPPI). 400 scans and 569 increments were acquired and computed in absolute value mode to a 16k × 1024 matrix with 2.5 Hz exponential multiplication in F2 and an unshifted sine bell in F1.
3 Results and discussion

The sea spray transport of DOM to the atmosphere was analyzed in the laboratory on the ship with fresh marine water samples, and we present the qualitative description of the organic fractionation during the passage from the surface water into the formed burst droplets toward the aerosols. Transfer of pollutants into the oceanic air and surface water was already followed in this journal issue with hexachlorocyclohexanes showing deposition and volatilization processes (Xie et al., 2011). Here we focused on chemical structural changes, depletions and enrichments in the process of primary marine aerosol formation via the bubble bursting process. The structural description of the organic material transferred to the atmosphere is based on FT-ICR MS and NMR.

The analysis of all FT-ICR mass signals within surface water, burst and daily aerosol samples is presented in Fig. 3. The van Krevelen diagram for the surface water mainly showed CHO and CHNO type compounds up to an m/z range of 800. At higher O/C and H/C values carbohydrate type compounds were also ionized. The cumulated burst samples had lower mass values up to 600 with a lower aromatic content, a relative increase in CHOS and a depletion of CHNO compounds. The cumulated data of all day aerosols reflected a typical aerosol fingerprint with mainly low mass values up to 500; aerosols showed an increased proportion of aromatic, nitrogen and sulfur compounds representative of anthropogenic input by biomass burning or ship exhaust as well as sulfonated secondary organic aerosols as described in Schmitt-Kopplin et al. (2010).

To extract the m/z values which commonly occurred in surface and burst samples and those which showed an average increase/depletion in the burst relative to the surface samples we followed the same data evaluation procedure as described in the chemical description of Champagne aerosol in Liger-Belair et al. (2009). We observed a clear relative increase in intensity of mainly CHO-compounds of higher aliphaticity and lower oxygen content (Fig. 4). This is in accordance with the previous study in which surface active compounds and fatty acids were preferentially concentrated in the
formed jets. In addition, many other compounds with a high binding affinity to those surfactants are co-fractionated and concentrated as well.

To verify this natural fractionation process, we selected a particular sampling point in the Angola Basin (3.1°E; −17.7°S) also studied with NMR in detail by Hertkorn et al. (2011). The surface water layer and the aerosols of this sampling site were specific of pristine open ocean (Fig. 5), with no anthropogenic organic signatures as found in the van Krevelen diagram of all transect data in Fig. 3. The surface water showed the typical distribution of marine DOM around the carboxylic rich alicyclic (CRAM) organic fraction (Hertkorn et al., 2006) which represents highly transformed organic matter; the corresponding structural network is very compact, representative of an extensive connectivity in involved mass differences. The burst concentrates mainly CHO and few CHOS compounds and lesser CHNO compounds; this is also visible in the looser structural network representation. Finally, the aerosol fraction showed a relative increase in surface active compounds of CHO and CHOS type in comparison with the surface water. The corresponding compositional network is much looser showing fewer connectivity corresponding to a less transformed type of organic materials probably of biological origin. The mass range followed the same trend as in Fig. 3 with lower m/z values in the order surface/burst/aerosol (not shown in figure).

Within all the stations of the cruise in compositional networks showing the same trends of looser network in the order surface/burst/aerosol, corresponding to lower overall chemical diversity. This is not written in caption of Fig. 6. In addition, we used this approach to integrate the datasets from these three fractions and highlighted the common masses in all three types of samples.

The three Angola Basin samples were analyzed with NMR spectroscopy (Fig. 7). All NMR spectra were characterized by large and distinct NMR resonances in the aliphatic section $\delta_H \sim 0.5–4.3$ ppm, indicating a suite of several abundant small spin systems containing aliphatic and oxygenated units with nicely resolved couplings. Polymethylene NMR resonances ($\delta_H \sim 1.28$ ppm) were abundant in all samples, especially in the aerosol sample where they contributed almost 50% of the entire proton NMR integral.
(Table 1). The ratio of methyl to polymethylene, an indicator of “pure” aliphatic branching ($\text{H}_3\text{C-C-C-Z}$, with any heteroatom Z at least four bonds away from methyl) grew in the order surface water $<$ aerosol $<$ burst; i.e. aliphatics appeared most branched in the burst sample. This was corroborated by a parallel increase of functionalized aliphatics with $\delta_H \sim 1.35$–1.9 ppm, which again followed the order surface water $<$ aerosol $<$ burst. Oxygenation, i.e. the occurrence of HCO substructures, were of equal abundance in aerosols and surface water and declined in burst (Fig. 7).

When comparing the spectra, total aliphatics ($\delta_H \sim 0.5$–1.9 ppm) grew in the order surface water $<$ burst $<$ aerosols. However, polymethylene contributed most to aerosols, whereas a large diversity of “pure” aliphatics occurred in burst. The water sample showed a specific underlying broad hump characteristic of CRAM which contributed to most of the section integral from $\delta_H \sim 1.9$–3.1 ppm.

Several of the aliphatic NMR resonances occurred in all three samples (Fig. 8), and a single COSY of sample burst could reveal major connectivity patterns. Intra-aliphatic cross peaks ($\delta_H \sim 0.8$–1.8 ppm) were common and bound to carbonyl derivatives and other functionalized aliphatics. Aliphatics connected to oxygenated units (C-HC-HC-O-) were less common than molecules with connectivities within oxygenated systems, i.e. (Y-O-C-HC-HC-O-Z, with arbitrary substitution for Y and Z). Burst sample showed conspicuous olefinic NMR peaks which were not present in the aerosol filter sample, but visible in the surface water. Based on J-couplings and chemical shift characteristics this spin system was assigned as acrylic acid. Acrylic acid is a bacterial cleavage product to DMS (dimethylsulfide) of DMSP (dimethylsulfoniopropionate), a ubiquitous osmoprotectant in marine biota (Yoch, 2002). It was shown having bio-signalling effect in the marine food web with antibacterial activities at higher concentrations (Zimmer and Butman, 2000) and has a moderate photostability leading to polyethylene type of photoproducts (Bajt et al., 1997). Acrylic acid was also found in this study to be co-concentrated into the sea spray relative to sea water and thus could undergo further photodegradation in the aerosol aqueous phase potentially leading to oxidized aliphatic type structures in primary and secondary organic aerosols.
The masses that increased in intensity from surface via bursts to aerosols were determined and visualized in Fig. 9. These compounds that were continuously concentrated from the marine surface to the atmosphere had especially very low masses ($m/z < 300$) involving mainly CHO and CHNO compounds. These masses were checked as possible biosignatures using the MassTRIX annotation webserver available for metabolomics (Suhre and Schmitt-Kopplin, 2008). The annotated $m/z$ mainly corresponded to the pathway of alpha-linolenic acid metabolism, fatty acid biosynthesis and metabolism, biosynthesis of unsaturated fatty acids, galactose, fructose and mannose metabolism and amino acids related biosynthesis and metabolism (Fig. 9).

About 120 masses were concentrated gradually from the surface into the atmosphere system and represented over 1000 isomers in MassTRIX commonly with fatty acid type of structures within the mass range 150 to 490 $m/z$. This group included homologous series of oxo-fatty acids, hydroxyl-fatty acids, mono-, di- and tricarboxylic acids, epoxy fatty acids, unsaturated fatty acids, primary and secondary alcohols, sugars, monoterpenes, methyl-branched fatty acids, straight chain fatty acids, methoxy-fatty acids, and glycerols. Recently, ultrahigh resolution mass spectrometry (FT-ICR MS) enabled us to discriminate hundreds of biochemical components (based on accurate exact mass analysis and database search) that are preferentially partitioning in the droplets ejected out of champagne; in comparison with the bulk liquid, the droplets following bubble collapse were definitely over-concentrated with various surface active compounds, many of them showing organoleptic interest (Liger-Belair et al., 2009). We also recently hypothesized that the organic fingerprint in marine water creates a specific smell which serves the olfactory orientation of marine animals (Koch et al., 2011).

The biomolecules of the surface water leave a specific molecular fingerprint in the aerosol. Our recent results show that the regional qualitative dissolved organic carbon (DOC) variations as found also in the surface water (Koch et al., 2011) and corresponding to Longhurst-ecological provinces of the sea (Longhurst, 1998) are partially reflected in the consequent aerosols (work in progress, Schmitt-Kopplin, 2011). We showed here experimentally on one specific site as well as on average over all the
data of the cruise that these compounds are preferentially concentrated in aerosols with a specific N- and S-partitioning and contribute largely to the formation of primary organic aerosols (POA) in the atmosphere. The biozone regional water biogeochemistry may thus have a direct impact on regional to global scales within possible lateral ocean-ocean or ocean-land transport of organic compounds in POA. Up to now, the transport and the effect of inorganic salt and less of organic compounds on the global climate have been intensively studied (Facchini et al., 2008a,b). Studies related to the effect of marine POA in atmospheric chemistry are very scarce (Eckström et al., 2010) and limited to low molecular organic compounds as most of the water soluble higher molecular weight and colloidal organic carbon remains uncharacterized (Rinaldi et al., 2010).

4 Conclusions

We described the structure discriminating transfer of DOM from marine surface water into the atmosphere via bubble bursting processes. This involves adsorptive bubble separation processes leading to a preferential concentration of surface active compounds in the bubble jets drop. These jet drops are conveyed into the atmosphere and form the main part of marine primary organic aerosols. Many surface active biomolecules were found to be selectively concentrated from the surface water through the simulated sea spray into the aerosol phase. Acrylic acid is also concentrated into the sea spray and probably further photodegraded into oxidized aliphatic compounds as found in the aerosol SOA. Ongoing studies integrate this information into a global description of the transfer processes leading to representative marine primary aerosols with structural fingerprints indicative of the biogeography and biogeochemistry of the corresponding water masses.
References


Péron, N., Cagna, A., Valade, M., Bliard, C., Aguíé-Béghin, V., and Douillard, R.: Layers of


Table 1. NMR section integrals (in percent of total NMR integral) according to coarse substructures given.

<table>
<thead>
<tr>
<th>δ(H) [ppm]</th>
<th>10–7.5</th>
<th>7.0–5.1</th>
<th>4.9–3.1</th>
<th>3.1–1.9</th>
<th>1.9–1.35</th>
<th>1.35–1.25</th>
<th>1.25–0.5</th>
<th>ratio</th>
<th>ratio</th>
<th>sum: 1.9–0.5 ppm</th>
</tr>
</thead>
<tbody>
<tr>
<td>Key substructures</td>
<td>H$_{3}$, HCOOH$^*$</td>
<td>C–CH$<em>{2}$O$</em>{2}$CH</td>
<td>HCO$^{**}$</td>
<td>HC–N, HC–C–O</td>
<td>HC–C–O– (CH$<em>{2}$)$</em>{n}$</td>
<td>H$<em>{3}$C/(CH$</em>{2}$)$_{n}$</td>
<td>H$<em>{3}$C/H$</em>{2}$C–C–O–</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Surface water</td>
<td>8.6</td>
<td>0.7</td>
<td>15.0</td>
<td>17.1</td>
<td>14.4</td>
<td>31.3</td>
<td>13.0</td>
<td>0.42</td>
<td>0.90</td>
<td>58.7</td>
</tr>
<tr>
<td>Burst</td>
<td>11.6</td>
<td>1.3</td>
<td>13.9</td>
<td>13.4</td>
<td>13.6</td>
<td>27.7</td>
<td>18.5</td>
<td>0.67</td>
<td>1.36</td>
<td>59.8</td>
</tr>
<tr>
<td>Day aerosol</td>
<td>6.4</td>
<td>0.2</td>
<td>15.2</td>
<td>10.0</td>
<td>8.9</td>
<td>48.5</td>
<td>10.8</td>
<td>0.22</td>
<td>1.21</td>
<td>68.2</td>
</tr>
</tbody>
</table>

$^*$ formic acid HCOOH ($\delta_H = 8.52$ ppm) is a major contributor in this section.

$^{**}$ section integral excluded HD$_{2}$COD NMR resonance at $\delta_H = 3.30$ ppm.
**Fig. 1.** Scheme of the two production ways of droplets from a bursting bubble; redrawn from Resch et al. (1986).
Fig. 2. Apparatus used for the generation and collection of the burst generated freshly from the surface water. The water was collected with the “fish” sampler in the surface of the water layer (left); bursting bubbles at the surface of the chamber (middle) and on sea water surface of the ocean (right).
Fig. 3. Representation of the ICR-FT MS converted data van Krevelen diagrams (right panels) of all the surface, all burst perhaps some cutoff/inclusion criteria should be defined, like and all day aerosol cumulated mass data over the entire cruise ANT-XXV/1. Left panels represent the m/z dependant H/C values and the relative frequency histograms of CHO (blue), CHOS (green), CHNO (orange) and CHNOS (red) compounds.
Fig. 4. Analysis of all burst (B) data versus all surface (S) data based on the intensity ratios \( \frac{R_{iB/S}}{} \) of \( m/z \) within 1 ppm similarity windows. Differentiation of the \( m/z \) intensity enhanced \( \left( \frac{R_{iB/S}}{} > 2 \right) \) or depleted \( \left( \frac{R_{iB/S}}{} < 2 \right) \) in the burst relative to the surface as illustrated in the van Krevelen diagrams (bubble size was proportional to the intensity ratios \( R_{iB/S} \)). A gradient in increasing ratio is found from higher oxygen content towards lower H/C values (unsaturated and aromatic compounds) and towards higher H/C (saturated).
Fig. 5. Sampling day 27 (Angola Basin) represented as van Krevelen diagrams of the surface, burst and day aerosol samples and corresponding compositional frequency histograms and corresponding compositional networks (color code as in previous figures).
Fig. 6. Compositional network representation of the mass differences connected with m/z of the surface, burst and day aerosol samples and their integration in the final network enabling to filter the common and unique signals for process description.
Fig. 7. 800 MHz $^1$H NMR spectra of aerosol samples in CD$_3$OD ($\delta_H = 3.30$ ppm). Center panel: entire spectrum with (top panel): vertical expansion with identical chemical shift scale. Bottom panel: expansions of aliphatic ($\delta_H = 0–5$ ppm) and unsaturated ($\delta_H = 5–10$ ppm) proton chemical environments with individual intensity adjustments to reveal also molecules of little abundance. Note the intensity ratio of the formic acid $\text{HCOOH}$ ($\delta_H = 8.52$ ppm) NMR peak compared with the other signals; the two adjacent signals are the $^{13}$C satellites [$^{1}$J($^{13}$C-$^1$H) = 191 Hz] with 0.5% abundance of the central signal.
Fig. 8. Panel (A): section of unsaturated protons with an olefinic spin system tentatively assigned as acrylic acid (red: chemical shifts [ppm], blue: J couplings [Hz]); a few neutral aromatics resonate from $\delta_H \sim 7.1$–7.4 ppm; the large NMR peaks at $\delta_H \sim 7.6$–7.8 ppm represent exchangeable protons, likely from phenolics or peptides. Panel (B): section of unsaturated protons in ocean water, indicating small amounts of acrylic acid (asterisks); small pH variations cause minor chemical shift displacement with respect to those shown in panel (A). Panel (C): aliphatic section of $^1$H, $^1$H COSY NMR spectrum of marine burst sample together with $^1$H NMR projection spectra of burst, surface water and aerosol and key substructures indicated. Section a: intra aliphatic HC-HC-HC-CH cross peaks with methyl ($\delta_H < 1$ ppm) and within branched aliphatics; section b: functionalized aliphatics connected with carbonyl derivatives C-HC-HC-COX (COX = COOH, COOR, CONH); note the absence for cross peaks further downfield from ($\delta_H > 2.4$ ppm) which excludes oxygenated spin systems (HCO-); section c: functionalized molecule with several remote heteroatoms Z-C-HC-CH-C-O [Z = O (N)]; section d: -O-HC-CH-O- cross peaks, esters, ethers and alcohols; section e: cross peaks HC-HC-HC-O- connecting aliphatics with oxygenated carbon.
Fig. 9. Top: extracted masses common to the surface/burst/aerosol system and increased from surface towards the atmosphere represented as \( m/z \) dependant H/C (left) and van Krevelen diagrams (right) with corresponding CHNOS-frequency diagram. Bottom: number of metabolites found in the various metabolic pathways.