



Quantification of 21 sugars in tropospheric particulate matter by ultra-high-performance liquid chromatography tandem mass spectrometry

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Abstract. Sugars compose an important class of compounds by mass in atmospheric particulate matter (PM), often
with biogenic and anthropogenic sources, many of them still poorly characterized. These sugars are mainly
15 analysed by gas-chromatography coupled to mass spectrometry (GC-MS) or ion chromatography coupled with
pulsed amperometric detection (IC-PAD). However, these techniques present several disadvantages such as a
complex preparation for GC-MS, or a limited range of possible analytes and elevated limits of quantification for
IC-PAD. This hinders our capability to perform analyses of extensive time series, in order to develop our
knowledge of the phenomenology of these species. In this paper, we present the validation of an ultra-high-
20 performance liquid chromatography tandem mass spectrometry (UHPLC-MS/MS) method for the simultaneous
quantification of 21 sugars in atmospheric PM. The sample preparation is simple, fast and safe, consisting of an
aqueous extraction. The method was validated in terms of linearity, accuracy, repeatability, reproducibility, and
recovery. This technique showed excellent linearity ($r^2 > 0.99$), precision (relative standard deviation $RSD < 25\%$)
and extraction yields (results between 70 and 120%). The suitability of the method for analyses of samples from
25 sites with very low PM concentrations was demonstrated with samples from the High-Altitude Research Station
Jungfraujoch (JFJ), Switzerland. A series of samples spanning a 6-year period is presented. Results for arabitol,
levoglucosan and 2-methyl-tetrols display strong seasonality, due to seasonal variation in chemical production and
boundary layer dynamics, with atmospheric convection and transport from the valleys to high altitudes mostly in
summer. This simple and fast method facilitates the analysis of large sets of PM samples and sugar compounds,
30 and opens the door to a better understanding and attribution of their sources.

1 Introduction

Particulate matter (PM) is involved in many aspects of Human environments, including large impacts on climate
change, health, dispersion of pollutants, or damages to all compartments of the critical zone. Some impacts can be
positive (the fertilization of Oceans), but most are negative, particularly when it comes to anthropogenic emissions.
35 Assessing these impacts requires a proper understanding of some properties of PM (mass, size distribution,
chemistry, physical and optical properties, etc.), and also a good parametrization of the fluxes in and out of the
atmosphere, in order to develop Chemical Transport Models (CTM). CTM are a prerequisite for the global
assessment of the impacts of the PM in Air Quality, Human health, or climate. However, the main component by



mass of PM, the organic matter (OM), is still poorly understood, even after decades of research on its chemical composition, emissions and transformation processes, and observations worldwide of its variability. CTM are generally off by up to an order of magnitude, compared to measurements (Ciarelli et al., 2016), meaning that sources and/or processes of OM components are not fully captured. Indeed, the current state of the art concerning molecular characterization of OM cannot allow a better mass apportionment than about 25 to 50% at best of the total OM mass (Michoud et al., 2021; Wang et al., 2020). Obviously, several OM sources or formation processes *in situ* are still not understood nor their emission fluxes properly quantified.

While the formation of Secondary Organic Aerosol (SOA) is largely studied by a score of papers every year, the Primary Biogenic Organic Aerosol (PBOA) is largely disregarded, while studies point out its large contribution to the OM fraction (up to 25-35 % of the total OM mass on seasonal average for France) (Samaké et al., 2019). As opposed to Secondary Biogenic Organic Aerosol (SBOA), that is formed in the atmosphere from the biogenic volatile organic compounds (VOCs) emitted by biota, PBOA is a subset of organic PM that comprises all particulate material of biogenic origin directly entering the atmosphere. PBOA includes components like plant debris, fungi, pollen, and viruses (Amato et al., 2017; Elbert et al., 2006; Hummel et al., 2015). The current estimate of the global PBOA flux to the atmosphere is between 10 and 310 Tg per year (Hummel et al., 2015), compared to values of 21 (MACCCity) or 30 (CMIP6) Tg per year for particulate organic components from anthropogenic origins (Rémy et al., 2019). The wide uncertainty on the flux of PBOA indicates by itself the extent of our poor knowledge on this fraction. Moreover, PBOA is currently very rarely included in CTM's (Rémy et al., 2019). One attempt to model PBOA on the European scale using the COSMO-ART CTM (Hummel et al., 2015) showed that PM mass associated with solely fungal spore emissions could represent on yearly average 15% of the total PM (about 25 - 30 % of organic PM) mass over Europe. A couple of papers very recently described the modelling of PBOA from fungal spore emission in CHIMERE (Vida et al., 2024) and EMEP models (Felix-Lange et al., in progress). A better assessment of this PBOA fraction is all the more important, since they also have a significant impact on the oxidative potential of PM, which can be a prevalent source in OP contribution at spring (Samake et al., 2017; Dominutti et al., 2023).

A large part of the PBOA fraction is related to the fungi emission sources (Hummel et al., 2015; Marynowski et al., 2020; Wang et al., 2020). Therefore, the seasonal cycles of some sugars and sugar alcohols (S and SA) concentrations observed in atmospheric PM can be directly related to the microbiology of large biomes. However, this link has only been characterized for a limited number of cases and S and SA (Marynowski et al., 2020; Samaké et al., 2020), and the main sources and drivers of many other S and SA in PM have not been identified yet. Currently, the mechanisms of S and SA emissions (along with their full organic cortege) to the atmosphere, and their fate and transfer to other ecosystem compartments remain poorly understood. Anhydrosugars, another form of sugars in atmospheric PM can be natural but also anthropogenic, such as levoglucosan and its isomers mannosan and galactosan, which are commonly used as a tracer for the assessment of PM from biomass burning (natural wildfires, domestic wood heating or cooking). These anhydrosugars – carbohydrate derivatives formed by the elimination of a water molecule - have been of primary importance for source apportionment studies since at least 15 years (Herich et al., 2014; Weber et al., 2019), and are now a basic measurement for PM analysis. The main suspected sources of other sugars measured in atmospheric PM are summarized in Table S1. Many of the links between chemical species and sources proposed in the literature Table S1 are still tentative, with only few comprehensive measurements for many of the species cited.



Analyses of sugars are mostly performed by gas chromatography coupled with mass spectrometry (GC-MS) (Vincenti et al., 2022). This method, with high selectivity and specificity, is generally performed with large sample volume and a sample preparation using derivatization of the hydroxyl groups (Graham et al., 2002). The introduction of ion chromatography coupled with pulsed amperometric detection (IC-PAD) in the mid 2000's (Engling et al., 2006; Yttri et al., 2007) made the determination of sugars in PM easier, at the expense of a narrowing of the range of the accessible sugars due to potential interferences with lack of column separation. Nevertheless, IC-PAD allows for the determination of 8 - 12 S and SA without too many interferences, including the three anhydrosugars (AS) (levoglucosan, mannosan, galactosan) (e.g., Oduber et al., 2021). In recent years, liquid chromatography tandem mass spectrometry (LC-MS/MS) has become more common in laboratories due to its greater efficiency and resolution, speed, simplicity, and low cost compared to more conventional chromatographic techniques (López-Ruiz et al., 2019). Indeed, LC-MS/MS can offer a much larger chemical speciation of S and SA and much lower limits of quantification than IC-PAD, together with much simpler sample preparation than GC-MS.

This work was focused on the validation of a UHPLC-MS/MS method for the simultaneous quantification of 21 sugars in PM, with a very simple and fast sample preparation. It presents the overall analytical method and its performances, including limits of quantification, linearity, intermediate precision studies, accuracy, and recovery. As a proof of concept, applicability, and to showcase challenging real-world samples, a timeseries of measurements of some species are presented for samples from Jungfraujoch (JFJ) site, Switzerland (3580 m a.s.l.), an alpine high-altitude international research station (Bukowiecki et al., 2016). Ultimately, the high sensitivity and ease of this method could pave the way for numerous additional studies on these families of chemical species, enhancing our understanding of atmospheric sugar cycles and their connection to biological processes.

2 Material and methods

2.1 Standards and reagents

A large literature survey was conducted to make an assessment of the sugars, sugar alcohols and anhydrosugars that have previously been cited in the literature concerning atmospheric particulate matter, as synthesized in Table S1. A large selection of compounds was initially tested during the method development and 28 of them were kept for the development processes and method validation. Finally, only 21 of them were properly quantified and monitored for routine analysis. The chemical structures and physicochemical properties of these target compounds are shown in Table S2 and their suppliers are presented below.

Adonitol (BioXtra, $\geq 99.0\%$ (HPLC)), L-(+)-Arabinose (BioUltra, $\geq 99.5\%$ (sum of enantiomers, HPLC)), L-(-)-Arabitol ($\geq 98\%$ (GC)), Erythritol (CRM), L-(+)-Erythrulose ($\geq 85\%$ (HPLC)), D-(-)-fructose ($\geq 99\%$ (HPLC)), D-(+)-galactose ($\geq 99\%$ (HPLC)), D-(+)-Glucose (BioUltra, anhydrous, $\geq 99.5\%$ (sum of enantiomers, HPLC)), Glycerol ($\geq 99.5\%$), Inositol (CRM), D-Lactose monohydrate (BioUltra, $\geq 99.5\%$), Levoglucosan (1,6-Anhydro- β -D-glucose (99%)), Maltitol ($\geq 98\%$ (HPLC)), D-(+)-Maltose monohydrate (BioUltra, $\geq 99.0\%$), D-Mannitol (BioUltra, $\geq 99.0\%$ (sum of enantiomers, HPLC)), D-(+)-Mannose (synthetic, $\geq 99\%$ (GC)), D-(+)-melezitose hydrate ($\geq 97\%$ (HPLC)), L-Rhamnose monohydrate ($\geq 99\%$), D-(-)-Ribose ($\geq 99\%$ (GC)), Sedoheptulosan, D-sorbitol (BioUltra, $\geq 99.0\%$ (HPLC)), Sucrose (Saccharose $\geq 99.5\%$ (GC), BioXtra), D-Threitol (99%), D-(+)-Trehalose dihydrate (CRM), D-(+)-Xylose ($\geq 99\%$ (GC)) and D-(+)-Xylose ($\geq 99\%$ (GC)) were obtained from



Merck-Sigma (France). Galactosan (1,6-Anhydro-beta-d-galactopyranose 97%) and Mannosan (1,6-Anhydro-beta-d-mannopyranose 97%) were purchased from Combi-Blocks (USA). 2-methyl-D-Erythritol (2S, 3R) and its
120 3 isomers (2-methyl-L-Erythritol (2R, 3S); 2-methyl-D-treitol (2S, 3S) et 2-methyl-L-treitol (2R, 3R)) were synthesized by Plateau Synthèse Organique, Département de Chimie Moléculaire (University of Grenoble Alpes, France) according to (Ghosh et al., 2012). Isotopically labelled standards, myo-Inositol (1,2,3,4,5,6-D₆, 98%), D-Glucose (1,2,3,4,5,6,6-D₇, 97-98%) and 1,6-Anhydro-beta-d-glucose (Levoglucosan) (U-13C₆, 98%) were obtained from Eurisotop. Ammonia solution (25%, LC-MS grade), Water (Rotisolv®, LC-MS grade) and
125 Acetonitrile (Rotisolv® ≥99.95%, LC-MS grade) were obtained from Roth.

2.2 Calibrators preparation

Individual stock solution of each target analyte and each internal standard are prepared in MilliQ® water (18.2 MΩ. hm.cm) at 1000 ppm and stored at 4°C. A stock solution of all target analytes (MIX-28) is prepared by mixing each target analytes in MilliQ® water. In parallel, a stock solution of the three internal standards, inositol-D6,
130 glucose-D7 and levoglucosan-13C6 (MIX-SI) is prepared in the same manner at 10 ppm. MIX-28 and MIX-SI water solutions are used to prepare an eight-point calibration curve in solvent with final concentrations of 90% acetonitrile, 0.005 % NH₄OH and 8 ppb for internal standards. Calibration range for each target analyte was adapted from a wide variety of analysed real samples, and is displayed in Table 2.

2.3 UHPLC-MS/MS conditions

135 All measurements were performed with ultra-high-performance liquid chromatography (ExionLC – AD binary pump, Shimadzu) coupled to a tandem mass spectrometer (AB SCIEX 5500 QTRAP). The column oven was set at 30°C. The injection volume was 30 µL. The chromatographic separation was performed on a Luna Omega Sugar (150 mm x 2.1 mm x 3 µm) column from Phenomenex (France) with a compatible guard column (SecurityGuard Cartridges, Sugar, 4 x 2.0 mm ID, Phenomenex). This column is of hydrophilic interaction chromatography
140 (HILIC) type and is adapted to hydrophilic interaction and separation of hydrophilic compounds such as sugars, in aqueous matrix. One overall goal of this development was to get simplified water extraction of the samples, but it made direct injection impossible since HILIC mode supports only comparable sample and eluants initial conditions (10% water, 90% acetonitrile). Other columns with such specifications (BEH Amide 100 mm x 2.1 mm x 1.7 µm from Waters, and HPLC Ultra Amino 150 mm x 3 mm x 3 µm from Restek) were tested but did not
145 provide such good results in terms of separation (peak shapes, return to baseline) after optimization, compared to the Luna Omega Sugar column.

Mobile phase (A) was 0.002% ammonia in H₂O (LC-MS grade) and mobile phase (B) was pure acetonitrile (LC-MS grade). The gradient elution was programmed as follows (2 isocratic steps followed by the rinsing of the column and equilibration): 0.00 – 9.00 min, 10 % mobile phase (A), 0.45 mL/min; 10.00 – 23.00 min, 20% mobile
150 phase (A), 0.40 mL/min; 23.01 – 28.00 min, 40 % mobile phase (A), 0.45 mL/min; 28.01 – 29.01 min, 10% mobile phase A, 0.45 mL/min; 29.01 – 36.00 min, 10% mobile phase A, 0.45 mL/min. Total run time was 36 min. In order to limit soiling of the MS system, the divert valve was programmed to MS analysis only between 1 and 25 min. The autosampler temperature was set to 5°C in order to limit potential evolution of the samples during the analytical batch.



155 The mass spectrometer was operated in negative electrospray ionization (ESI) mode using a scheduled multiple reaction monitoring (MRM). The ESI parameters were optimised and finally set to the following ones: spray voltage -4500 V, source temperature 400°C, Ion Source Gas 1 (GS1) 40.0 psi, Ion Source Gas 2 (GS2) 60.0 psi. The MRM transition per compounds were determined by infusion of individual standards at 100 ppb in 10/90% water/acetonitrile, in presence of 0.005 % NH₄OH. For each target analyte, infusion of monospecific standard

160 solution was made: a full scan mass spectrum was acquired to determine its precursor ion. Then, product ions scans were acquired on the selected precursor ion to select a product ion of interest. One MRM transition is chosen to characterize each compound. All MRM parameters for each analyte are summarized in Table 1.

Given the fact that some analytes are isomers, the identification of each analyte is also performed using its elution order and its retention time. Given the nature of the HILIC phase, retention time may slightly change between each

165 batch, and a standard solution is used to recalibrate the retention time of each analyte before batch launches. The system is controlled with the Analyst® software version 1.7 (AB SCIEX). All data processing (integration peak and quantification) was conducted on SCIEX OS v1.7, using individual internal standards for each compound with the external calibration to correct potential ionisation variations in the source.

170 **Table 1 : Multiple Reaction Monitoring (MRM) parameters for all analytes and internal standards. DP: Declustering Potential; CE: Collision Energy; CXP: Collision Cell Exit Potential; *: retention time can be adjusted according to the ageing of the column.**

Analyte	Q1 (m/z)	Q3 (m/z)	Retention time (min)*	MRM window (sec)	DP (volts)	CE (volts)	CXP (volts)	IS
Levoglucozan-13C6 (L-IS)	166.906	105.000	2.38	60.0	-95	-14	-11	-
Glucose-D7 (G-IS)	185.991	60.900	11.21	120.0	-40	-22	-9	-
Inositol-D6 (I-IS)	184.899	89.000	15.20	60.0	-110	-24	-11	-
Adonitol	150.941	89.000	5.75	60.0	-60	-16	-11	L-IS
Arabinose	148.932	89.000	6.00	120.0	-30	-10	-11	L-IS
Arabitol	150.920	89.000	6.41	120.0	-75	-16	-11	L-IS
Erythritol	120.946	89.000	3.83	60.0	-45	-14	-11	L-IS
Erythrulose	119.132	70.900	2.03	60.0	-30	-16	-9	L-IS
Fructose	178.918	89.000	7.78	180.0	-40	-12	-9	G-IS
Galactosan	160.947	100.900	2.09	60.0	-75	-16	-11	L-IS
Galactose	178.903	89.000	12.00	90.0	-65	-12	-11	G-IS
Glucose	178.984	89.000	11.14	200.0	-55	-10	-11	G-IS
Glycerol	90.916	59.000	2.41	60.0	-55	-14	-7	L-IS
Inositol	178.921	86.900	15.07	60.0	-90	-22	-13	I-IS
Lactose	341.007	161.000	17.85	120.0	-95	-10	-17	I-IS
Levoglucozan	160.900	100.900	2.39	60.0	-70	-14	-11	L-IS
Maltitol	343.017	179.000	16.55	90.0	-135	-20	-11	I-IS
Maltose	341.005	161.000	16.29	120.0	-90	-10	-15	I-IS
Mannitol	180.923	89.000	11.36	120.0	-100	-18	-11	G-IS
Mannosan	160.901	100.900	2.09	60.0	-65	-18	-11	L-IS
Mannose	178.919	89.000	10.25	60.0	-45	-12	-11	G-IS
Melezitose	503.061	323.100	21.90	90.0	-180	-28	-15	I-IS
Rhamnose	162.894	59.000	3.47	60.0	-60	-18	-9	L-IS
Ribose	148.927	88.900	3.50	90.0	-45	-10	-11	L-IS
Sedoheptulosan	190.897	116.900	6.66	90.0	-90	-20	-13	L-IS



Analyte	Q1 (m/z)	Q3 (m/z)	Retention time (min)*	MRM window (sec)	DP (volts)	CE (volts)	CXP (volts)	IS
Sorbitol	180.913	88.900	10.71	120.0	-80	-20	-11	G-IS
Sucrose	341.013	59.000	14.61	60.0	-130	-54	-7	I-IS
Threitol	120.942	88.900	3.83	120.0	-55	-14	-11	L-IS
Trehalose	341.006	58.900	17.48	60.0	-150	-52	-7	I-IS
Xylose	148.922	59.000	4.67	60.0	-30	-18	-7	L-IS
2-methyl-tetrols	134.941	85.000	2.76	60.0	-60	-20	-9	L-IS

2.4 Sample collection and preparation

Ambient aerosol samples were collected at the High Altitude Research Station Jungfraujoch (JFJ) (3580 m a.s.l., Switzerland) as part of the Swiss National Air Pollution Monitoring Network (NABEL). The JFJ station is located in the free troposphere with regular intrusion of planetary boundary layer air masses. It generally offers very low background aerosol concentrations. Samples were collected over 24h on quartz fiber filters (150 mm diameter discs, Pallflex 2500 QAT-UP) using DHA-80 high volume samplers (DIGITEL) with a PM10 inlet, operating at 45 m³ h⁻¹. After sampling, the filter samples are transported back to the laboratory for gravimetric determination of the PM10 mass concentration. Subsequently, the filters are put inside a glassine envelope (PAWI Packaging Schweiz AG), sealed in polyethylene bags and stored at -18°C until further analysis (Hueglin et al., 2005). The samples studied here were composites of 4 daily filters distributed over 2 weeks, for a period of several years from 2011-2016. A total of 136 composite samples and 11 field blanks were analysed.

For the extraction of analytes of interest, a portion of the quartz filter (typically a total of 10.2 cm² for each composite sample from the JFJ site) was used. The filter was extracted in 6 mL of MilliQ® water with vortex shaking for 20 min. This sample preparation was adjusted for JFJ samples for which very low concentrations of sugars were expected. In cases of daily samples from most low altitude sites, 5 cm² of quartz filter are generally extracted in 7 mL of water (Aas et al., 2025). Then, the supernatant is filtered with 0.2 µm Ion Chromatography Acrodisc®13 pre-washed with ultrapure water. If necessary, extracts are stored in a freezer at -20°C for later analysis. Finally, 100 µL of the extract is diluted with 900 µL of the dilution mixture (99% acetonitrile, 0.005% NH₄OH and internal standard adjusted at 9 ppb from MIX-SI) to reach chromatographic equilibration conditions, and then analysed by UHPLC-MS/MS. Samples stay 69 hours at most in the refrigerated autosampler before analysis, in order to avoid potential aging processes.

2.5 Method performance evaluation

2.5.1 Method validation

The performance of the method was evaluated by studying linearity, limit of quantification (LOQ), limit of detection (LOD), precision, bias, and recovery. The linearity was evaluated with 8 points of calibration per compound. The calibration curves were determined using least-squares linear regressions. The LOD was determined as the minimal detected standard with a signal/noise ratio larger than 3. Precision studies including intra-day and inter-day evaluations, and bias were determined with the injection of a standard every 8 injections during each batch. Accuracy was evaluated for levoglucosan by comparison to standard reference material NIST®



SRM® 2786 (Fine Atmospheric Particulate Matter). Recovery was evaluated on summer and winter filters collected on the roof of the laboratory in Grenoble (France), to test for a potential seasonal effect.

2.5.2 Calculation and analytical batch

205 The quantification is based on the isotope dilution with the addition of isotope labelled sugars (inositol-D6, glucose-D7 and levoglucosan-13C6) into the injected sample to correct analytical variation of the source of the mass spectrometer. Calibration curves are drawn by plotting the area ratio (Compound area/IS area) against the concentration ratio (Compound concentration/IS concentration). Thus, the sample concentration, in ppb, can be determined with the area ratio of the sample. Finally, results were converted in ng/m^3 using the extracted surface
210 of filter, the volume of air collected, the initial volume of water for the extraction, the dilution factor with the eluant, and taking into account the average of the sample field blanks of the corresponding field campaign. In daily-routine, full set of calibration solutions are injected every 40 samples (about every 24 hours of analysis), and the standard 3 of the calibration range (STD3) is injected every 8 injections as a quality control. These repetitions are performed to ensure that the LC conditions, in particular column ageing, are compatible with an
215 accurate quantification of samples.

3 Results and discussion

3.1 Sample preparation

The sample preparation exclusively in water and without derivatization is intended to be simple, limiting the preparation stages, the handling, the possibilities of contamination, and the use of solvents and other chemicals.
220 All of these convey major advantages for this preparation in terms of time consumption, consumables costs, and required manpower. Moreover, these same aqueous extracts can be directly used for other analyses such as major ions and organic acids by ion chromatography coupled to mass spectrometer on anion canal (IC-MS) (Glojek et al., 2024), sugars analysis by IC-PAD (Samaké et al., 2019), or HUmic LIke Substances analysis using a TOC analyser after separation by HPLC-Fluorescent (Baduel et al., 2009). Therefore, these extracts, the associated costs
225 and the consumption of the sample filter surface can be shared between these different analyses, which is another big advantage.

3.2 Method validation

3.2.1 Retention times and co-elutions

230 The 28 compounds eluted along the 25.0 min MS analysis period (Figure 1) and retention times were repeatable with a relative standard deviation (RSD) below 2.0% for all compounds for 146 successive injections over a period of 61 hours. However, some chemical species cannot be separated in this run. First, the 2-methyl-erythritol and 2-methyl-threitol four stereoisomers are co-eluted and present identical MRM transition, but they also provide the same response factor. 2-methyl-D-Erythritol is then used as the quantification
235 standard and represents therefore the four isomers, named commonly as 2-methyl-tetrols. Second, the two compounds in each pair galactosan/mannosan and erythritol/threitol present the same retention time and the same



MRM transitions, but with different response factors in terms of intensity for identical concentrations. Their quantification cannot be achieved with this method.

Finally, the Luna Omega Sugar column was globally suitable for the analysis of the other sugar compounds from Table 1, but it must be replaced around every 300-600 injections due to the column ageing. Indeed, a drift of retention time was observed with ageing, with (for the example of figure S1) an average over all 28 compounds of 0.5 min after 512 injections, which can go up to 2.2 min for the case of melezitose at the end of the run (Figure S1). This ageing may be due to anion (such as Cl^- from extract samples) progressive occupation of active separation sites (like amine), leading to less effective separation of sugars and retention time decrease through numerous and charged sample injections (personal communication J. Lacouchie-Payen, Phenomenex). The pairs adonitol/arabitol and sorbitol/mannitol tend to co-elute with increased column ageing (Figure S1). This cannot be reversed with any cleaning procedure. A criterion to discard an aged column is the difficulty to separate completely those 2 pairs of compounds.

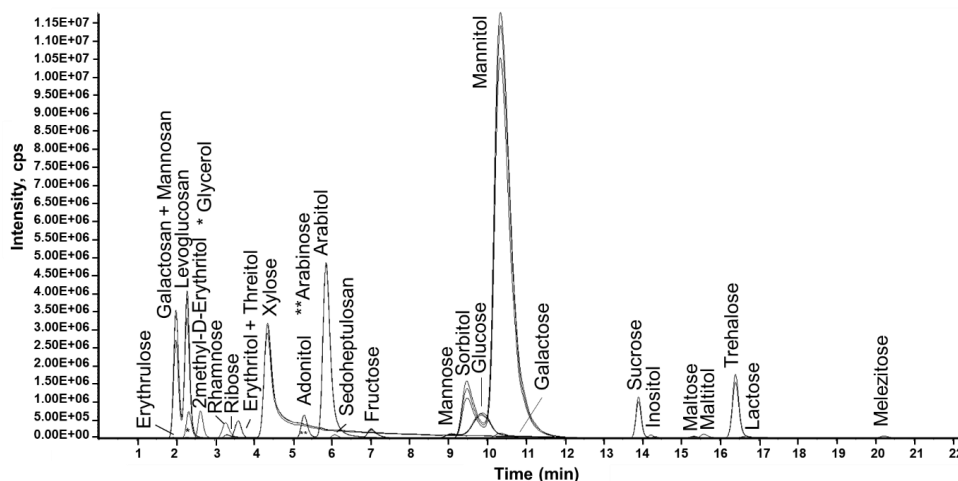


Figure 1: Chromatogram of standard 1 (STD1), the highest concentrated standard of calibration range, showing the elution of the 28 sugars, sugar alcohols and anhydro-sugars on the Luna Omega Sugar column.

3.2.2 Cases of tested but not analyzable compounds

Several compounds were tested and monitored but no result can be returned. This is the case for arabinose and galactose which were not automatically detected in most of the analytical batches due to their weak intensity and their elution at the end of the trailing peak of xylose and glucose, respectively. Therefore, linearity cannot be evaluated, neither proper quantification. However, our experience with many series of samples in various atmospheric conditions is that they are not present in the atmospheric PM10 with the LOD mentioned in Table 2. Finally, glycerol is also a specific case. It has been reported in plants in high concentrations (Gerber et al., 1988) and has an influence on plant flowering (Lazare et al., 2019). Moreover, Kang et al. (2017) found higher level of glycerol in PM2.5 filter samples during winter and autumn when vegetation decays and fungal population increases. Several studies worldwide confirmed the presence of glycerol in PM and associated it with biomass burning (Zangrando et al., 2016). Our results of the method performance evaluation showed that glycerol can be easily contaminated. This can take place at any time from the sampling to the LC-MS/MS analysis, as currently



glycerol is present in everyday life (Bagnato et al., 2017) as it is included in a large variety of products such as cosmetics, food (E422, (EFSA Panel on Food Additives and Flavourings (FAF) et al., 2022), pharmaceutical and tobacco products in particular e-cigarettes (Kubica, 2023). We choose not to report glycerol concentrations since results obtained are often (but not always) erratic. More work must be done for understanding the contamination processes, limiting them, and obtain trustable results.

3.2.3 Linearity and Limit of Detection (LOD)

The calibration range defined for this method was determined for each compound in order to represent the full range of the concentrations generally expected from atmospheric samples prepared with our usual sampling and extraction conditions. This was determined following our experience for the compounds measured from a large literature review. The linearity was evaluated for the full 8 points calibration range of each compound. A very good correlation between prescribed and observed concentrations were observed with r^2 above 0.99 (Pearson's criteria), except for maltitol and sorbitol (Table 2).

The analytical Limit of Detection (LOD) is the smallest standard detected in the analytical run, with a signal/noise ratio > 3 . Values are displayed in ppb in the extract. The analytical LOD is compound-dependent and varies from 0.001 – 2.5 ppb, and after conversion in ng/m^3 using our standard sampling and extraction conditions, from 0.001 – 3.6 ng/m^3 (Table 2). However, the actual field LOD will vary according to the field blank value, for each compound and each field campaign.

3.2.4 Intermediate precision studies

In our classical analytical batch, the standard 3 (STD3) is injected every 8 injections as a quality control. Bias is calculated for STD3 as follows (Eq. 1):

$$\text{Bias} = \left| 100 - \frac{\text{Experimental concentration of STD3}}{\text{Theoretical concentration of STD3}} \times 100 \right| \quad (1)$$

94% of intra- and inter-day bias results obtained over a period of 5 days were below a 25% bias, meaning that the method quantification is robust (Table 2). 83% of intra- and inter-day RSD results obtained over a period of 5 days were also below 25%, meaning that this UHPLC-MS/MS method is repeatable and reproducible (Table 2). We note that for intra-day, more RSD results ($n=6$) were larger than 25%, compared to inter-day RSD results ($n=3$). However, for the compounds with particular interest (like the ones used in source apportionment studies such as arabitol, fructose, glucose, inositol, levoglucosan, mannitol, sucrose, trehalose and 2-methyl-tetrols), RSD results were similar between intra-, and inter- day. Conversely, the species with the worst results happen to be of lower interest for atmospheric samples, with maltitol never being detected with a LOD of 0.05 ng/m^3 , and sorbitol rarely seen with a LOD of 0.005 ng/m^3 .

Table 2 : Performance of the sugars UHPLC-MS/MS method. RSD: Relative Standard Deviation; LOD: Limit of Detection; STD3: standard 3 of the calibration range

Compounds	Calibration range (ppb)	Linearity (r^2)	STD3 Concentration (ppb)	Intra-day (n=13)		Inter-day (n=5)		Median LOD (ppb) (n=5)	Median LOD (ng/m^3) (n=5)
				Bias (%)	RSD (%)	Bias (%)	RSD (%)		
Adonitol	0.001 - 10	0.9992	1.000	7.6	17.3	6.7	16.6	0.001	0.011



Compounds	Calibration range (ppb)	Linearity (r ²)	STD3 Concentration (ppb)	Intra-day (n=13)		Inter-day (n=5)		Median LOD (ppb) (n=5)	Median LOD (ng/m ³) (n=5)
				Bias (%)	RSD (%)	Bias (%)	RSD (%)		
Arabinose	0.005 - 50	N/A	4.993	N/A	N/A	N/A	N/A	2.497	3.614
Arabitol	0.010 -100	0.9997	10.000	8.9	21.9	11.0	18.5	0.010	0.011
Erythrulose	0.001 - 10	0.9985	1.001	10.2	16.6	0.5	17.2	0.010	0.108
Fructose	0.001 -10	0.9959	0.994	38.8	15.2	10.6	15.5	0.001	0.001
Galactose	0.001 -10	0.8904	1.000	N/A	N/A	N/A	N/A	0.010	0.050
Glucose	0.010 -100	0.9995	9.938	3.5	3.2	1.7	7.9	0.010	0.042
Glycerol	0.050 -500	0.9986	50.000	17.7	9.4	12.9	17.5	0.050	0.054
Inositol	0.001 -10	0.9995	1.010	0.9	6.2	4.1	5.7	0.051	0.055
Lactose	0.001 -10	0.9998	0.998	8.0	20.0	9.2	17.0	0.050	0.054
Levoglucosan	0.100 -1000	0.9998	100.380	3.6	5.8	1.3	4.1	0.100	1.084
Maltitol	0.001 -10	0.9684	1.004	49.7	57.4	23.9	23.2	0.050	0.054
Maltose	0.001 -10	0.9981	0.996	7.5	17.0	6.2	18.2	0.050	0.042
Mannitol	0.050 - 500	0.9988	49.600	12.5	26.5	19.2	21.5	0.050	0.054
Mannose	0.010 - 100	0.9991	10.000	2.1	26.9	1.4	27.8	0.100	0.540
Melezitose	0.001 -10	0.9989	1.006	7.6	45.4	7.8	28.5	0.010	0.011
Rhamnose	0.010 - 100	0.9996	10.060	7.6	5.6	3.9	10.0	0.010	0.109
Ribose	0.005 -50	0.9936	5.000	12.1	28.5	15.5	21.4	0.250	0.270
Sedoheptulosan	0.001 -10	0.9991	1.008	3.9	6.0	2.5	10.8	0.050	0.109
Sorbitol	0.005 - 50	0.9466	4.959	63.7	110.8	14.2	54.5	0.005	0.005
Sucrose	0.005 -50	0.9941	5.040	19.5	10.0	14.5	9.9	0.005	0.005
Trehalose	0.010 - 100	0.9996	9.920	8.0	9.9	0.6	8.4	0.100	0.107
Xylose	0.100 - 1000	0.9996	99.800	10.2	8.0	5.3	12.3	0.100	0.108
2-methyl-tetrols	0.005 - 50	0.9997	4.494	7.4	7.5	3.5	6.4	0.004	0.007

3.2.5 Accuracy

Accuracy was evaluated for levoglucosan by aqueous extraction and UHPLC-MS/MS analysis of fine particulate matter reference material of Standard Reference Material (SRM®) 2786 from the National Institute of Standards and Technology (NIST®) (NIST® SRM® 2786) (n=6). Precision was 4.2% and bias was 10.0% which were compliant with RSD <15% and bias ±15%. In this SRM, concentrations for galactosan and mannosan are also available, but due to coelution and identical MRM transitions with different response factors in terms of intensity for identical concentration, their quantification is not guaranteed with this method. Currently, to our knowledge, no SRM is available for the others sugars monitored in this UHPLC-MS/MS method. However, for information purposes only, results of the concentrations (mg/kg) for 21 sugars of the NIST® SRM® 2786 are provided in Table S3.

3.2.6 Recovery

Extraction yield was assessed by comparing results from samples spiked at the STD3 level before and after extraction. The experiment was performed on summer (n=3) and winter filters (n=3) collected in the urban background atmosphere of Grenoble. The average chemical composition of the PM10 for a large range of compounds is described in Borlaza et al., (2021). It should be noted that several of the compounds presented in



Table 3 present very low concentrations (sub-ng/m³ or lower), explaining some of the very high results (like sedoheptulosan). However, mean extraction yields were satisfying with most of the results between 70 and 120% (Table 3).

Table 3: Extraction yield evaluated on summer and winter filters.

Compounds	Extraction yield (%)		
	Summer filter (n=3)	Winter filter (n=3)	Mean
Adonitol	82.4	88.2	85.3
Arabitol	83.7	89.3	86.5
Erythrulose	N/A	N/A	N/A
Fructose	83.9	90.8	87.4
Glucose	97.8	122	110
Glycerol	80.5	90.3	85.4
Inositol	81.1	88.1	84.6
Lactose	84.0	84.2	84.1
Levogluconan	77.2	94.5	85.8
Maltitol	53.9	71.5	62.7
Maltose	91.9	68.5	80.2
Mannitol	77.0	84.4	80.7
Mannose	81.4	87.6	84.5
Melezitose	88.0	101	94.6
Rhamnose	81.5	97.9	89.7
Sedoheptulosan	702	93.0	82.2
Sorbitol	57.6	78.8	68.2
Sucrose	201	110	155.4
Trehalose	80.7	100	90.4
Xylose	79.5	81.0	80.3
2-methyl-tetrols	89.0	90.2	89.6

In summary, this UHPLC-MS/MS method provides reliable and low-level quantification for 21 sugars for atmospheric fine particles including sugars (erythrulose, fructose, glucose, lactose, maltose, mannose, melezitose, rhamnose, ribose, sucrose, trehalose, xylose), sugar alcohols (adonitol, arabitol, inositol, maltitol, mannitol, sorbitol, 2-methyl-tetrols) and anhydro-sugars (levogluconan, sedoheptulosan). New sugars (adonitol, erythrulose, fructose, maltitol, maltose, melezitose, sedoheptulosan, sucrose, trehalose, xylose and 2-methyl-tetrols) not quantified with traditional IC-PAD method (Engling et al., 2006; Yttri et al., 2007; Samaké et al., 2019) are provided and will allow further exploration of their sources and the processes they undergo in the atmosphere.

3.3 Jungfrauojoch 6-year time series of samples

3.3.1 Atmospheric concentration calculations

The interest of this UHPLC-MS/MS method is demonstrated with the analysis of a long time series of samples collected in the low concentration levels environment of the High Altitude Research Station Jungfrauojoch (Switzerland). These samples can be regarded as challenging, insofar as they usually contain very low PM values



and hence sugar concentrations. With the simple and fast sample preparation method, the 136 composite samples and 11 fields blanks were prepared in two days. After analysis, the results of composites and field blanks were processed in the same manner in SCIEX OS. The field limit of quantification (Field LOQ) was calculated for each compound (Table S4), based on the field blanks results, as follows: (Eq. 2)

$$\text{Field LOQ} = \text{Mean}_{\text{field blanks}} + 2 \times \text{Standard Deviation}_{\text{field blanks}} \quad (2)$$

When results for composite samples were above this Field LOQ, the value of mean field blank was subtracted from the initial sample value in ppb, subsequently converted in ng/m^3 .

3.3.2 Evolution of the concentrations

Median and seasonal concentrations of the 21 quantifiable sugars are presented in Table S4. The time series of 3 sugars of interest (arabitol, 2-methyl-tetrols and levoglucosan) are plotted on Figure 2 over a 6-year period, with a time resolution of 20 composite samples per year.

Arabitol displays values between LOD (0.011 ng/m^3) and 2 ng/m^3 , with seasonal variation patterns (May–November occurrence) and maximal values in high summer. Interannual variability of the maxima summer concentrations are between 0.6 and 2 ng/m^3 . Following the same pattern as arabitol, the concentrations of other primary biogenic (PBOA) sugars such as mannitol, glucose, fructose, trehalose, and melezitose also display strong seasonal variations with much larger concentration in summer that are below LOD in winter. All concentrations (Table S4) are much lower (by a factor of about 20) than usual concentrations in low-altitude environments in France, and also in Alpine valleys (e.g. in Grenoble (Samaké et al., 2019) or Slovenian valleys (Glojek et al., 2024)).

In comparison, the secondary organic aerosol (SOA) tracer 2-methyl-tetrols, associated with the oxidation products from isoprene emissions, also displays strong seasonality with inter-annual variability of concentrations, but with a shorter occurrence of this compound compared to arabitol, restricted to June-to-early September (Figure 2). Contrary to primary biogenic sugars, concentrations reach much higher values (between 5 and 25 ng/m^3 for each summer maximum), and are only 4 to 12 times lower than in Alpine lower altitudes (Glojek et al., 2024).

Finally, the levoglucosan timeseries is really noisy, without any particular seasonal pattern (Figure 2). The concentrations are low ($< 6 \text{ ng/m}^3$), except for three specific episodes in July months, reaching up to 77 ng/m^3 . Those low background concentrations in summer are equivalent to those in lowlands, but in winter are 500–1000 times lower than winter concentrations in alpine valleys that can reach up to $8 \mu\text{g/m}^3$ in cases of temperature inversion layers in narrow Alpine valleys (Bonvalot et al., 2016; Herich et al., 2014).

These results align well with biogeochemical understanding in light of the specific environmental context of the JFJ site, where the atmospheric signal can only be captured when both production in the surrounding lowland areas and efficient transport to high altitudes occur concurrently. The levoglucosan time series is consistent with findings from a 14-months Aerosol Chemical Speciation Monitor (ACSM) study at JFJ (Fröhlich et al., 2015), which highlighted the absence of local planetary boundary layer (PBL) sources during summer, and the lack of vertical PBL transport to the site in winter, despite elevated winter PBL concentrations primarily driven by domestic wood burning. The few observed peaks are most likely associated with long-range transported biomass burning plumes, such as those originating from Canadian wildfires in late June and July 2013, which were confirmed to have reached JFJ based on Copernicus Atmosphere Monitoring Service (CAMS) data.



Similarly, the distinct seasonality observed for arabitol and 2-methyl-tetrols (representing PBOA and SOA, respectively) reflects the influence of local sources within the PBL combined with efficient upward transport to JFJ during summer, when convective mixing is favored by warmer conditions. In contrast, the absence of both significant sources and vertical transport during winter explains the minimal concentrations observed during this season. Furthermore, the observed concentration differences between arabitol and 2-methyl-tetrols may reflect variations in local PBL composition as well as differential chemical aging during transport to JFJ. The full time series of sugars concentrations will be discussed in a dedicated publication, including a larger array of chemical species.

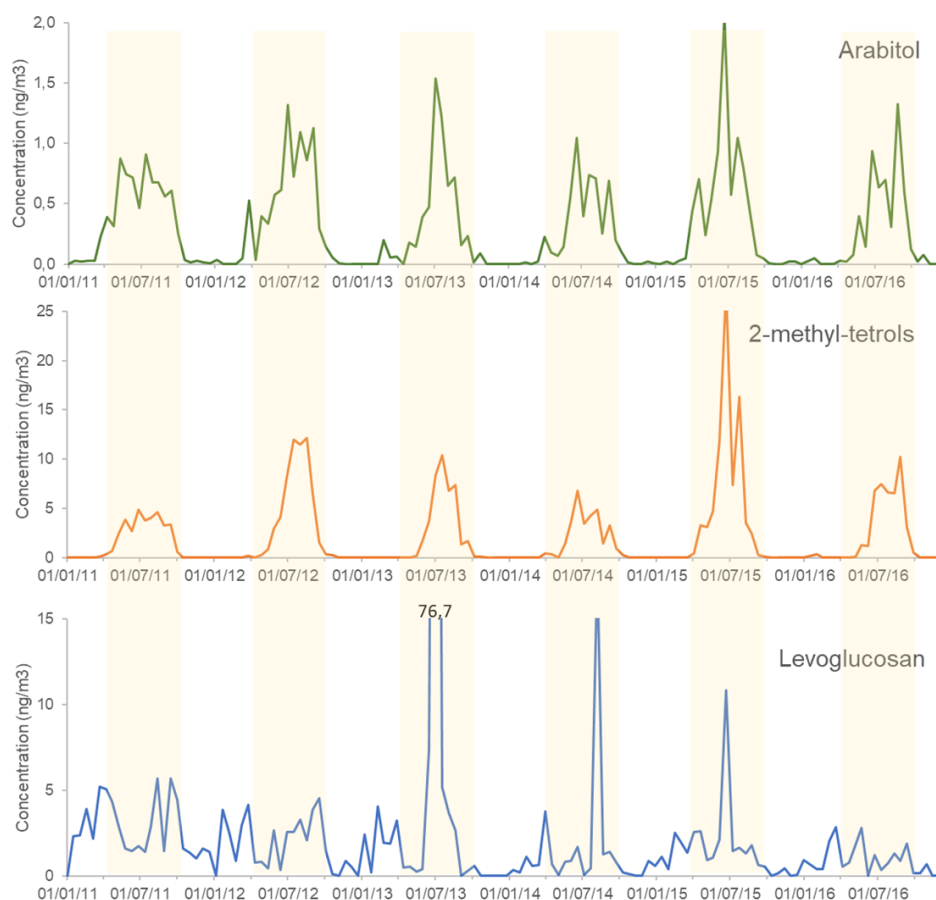


Figure 2 : Evolution of sugars arabitol, 2-methyl-tetrols and levoglucosan concentrations in the air (ng/m^3) over years 2011-2016 at Jungfraujoch.

4 Conclusion

This UHPLC-MS/MS method for molecular-resolution quantification of sugars in atmospheric PM₁₀ samples offers several advantages. First, the sample preparation is intended to be simple, with an extraction in water,



without any other steps like solvent reduction or derivatization. This conveys a major advantage of this preparation in terms of time consumption, consumables costs and required manpower. The preparation of the series of samples (n = 147) taken as an example in this work was performed in only two days. Moreover, this sample preparation can be adjusted according to the type of samples, including low concentrations as in this work.

Analytical results were very good in terms of linearity ($r^2 > 0.99$), precision ($RSD < 25\%$), extraction yields (results between 70 and 120%). Accuracy was compliant by comparison with the NIST® SRM® 2786. This method offers a larger number of sugar compounds properly quantified than with traditional IC-PAD method), which will allow enhanced exploration of sources and fate of a larger array of sugars in the atmosphere.

This highly sensitive method was successfully applied to 136 composite samples and 11 field blanks from Jungfraujoch (JFJ), Switzerland, a high latitude alpine station. These JFJ results provide concentrations for the European free troposphere, and indicate large seasonal variations over a 6-year period for many compounds, including arabitol, 2-methyl-tetrols and levoglucosan.

The high-resolution molecular-level chemical speciation achieved with this approach enables to further explore sugars sources and processes. This method has already proven effective across a range of sites and sample types: PM10 samples in Alpine valley sites, like in Kanal (Slovenia) (Glojek et al., 2024) and Grenoble (France) (Cruaud et al., in prep), but also in cloud water (Bianco et al., 2025), snow samples (Schivalocchi et al., in prep) and ice cores samples (Piot et al., in prep). The wide applicability across matrix—from atmospheric aerosols to snow, cloud water, and ice cores—demonstrates its potential to greatly enhance our understanding of the dynamics of sugar compounds in various environmental compartments. As such, this method opens new opportunities for in-depth investigations into the sources, transformations, and transport of sugars in the atmosphere, as well as their interactions with climatic and biological processes.

Author contributions

Pauline Bros: Investigation, Writing -Original Draft, Visualization, Writing - Review & Editing; **Sophie Darfeuille:** Methodology, Investigation, Writing - Review & Editing, Visualization, Supervision; **Véronique Jacob:** Methodology ; **Rhabira Elazzouzi:** Resources, Data Curation ; **Dielleza Tusha:** Investigation ; **Tristan Rousseau:** Investigation ; **Julian Weng :** Investigation, Writing - Review & Editing; **Patrik Winiger:** Resources, Writing - Review & Editing, Funding acquisition; **Imad El Haddad:** Resources, Writing - Review & Editing, Funding acquisition; **Christoph Hueglin :** Resources, Writing - Review & Editing, Data Curation; **Gaëlle Uzu :** Funding acquisition; **Jean-Luc Jaffrezo :** Conceptualisation, Writing - Review & Editing, Supervision, Funding acquisition, Project administration

Competing interests

The authors declare that they have no competing interests.

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