Fast Detection of 5-Hydroxymethylfurfural in Dulce de Leche by SPE-LC–MS

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Abstract
This research paper investigates the use of 5-hydroxymethylfurfural (5-HMF) as marker for the heat treatment of spreadable dairy creams (dulce de leche, DL). The proposed method applies solid-phase extraction (SPE) with final analysis by liquid chromatography coupled with high-resolution mass spectrometry (SPE-LC–MS). The method was successfully applied to analyze spreadable dairy creams prepared by hot melt extrusion using different heating temperatures from 100 to 130 °C. The concentrations of 5-HMF correlated linearly with the applied temperatures, with a signal response in the range from 0.5 to 100 μM ($R^2 = 0.9997$). The limit of detection (LOD) was 1.54 ± 0.03 μM with a precision of 1.77%. The results were compared with the analysis of 5-HMF in spreadable dairy creams using reference methods for the determination of 5-HMF in milk products. These methods mainly employed acid digestion and derivatization as pre-processing steps and determined 5-HMF spectrophotometrically and via HPLC–UV. These resulted in higher LOD (2.99 and 2.01 μM) and less precision (4.44 and 2.09%) compared to the proposed method. Furthermore, the proposed pre-processing procedure was faster by omitting the acid digestion and derivatization steps and by employing SPE.

Keywords Spreadable dairy creams · LC–MS · Solid-phase extraction · Maillard reaction · Thermal processing

Spreadable dairy creams are a popular snack food, especially within children. A typical example of spreadable dairy creams is represented by dulce de leche (DL), a brown and sweet cream obtained by the extensive heating of milk, when mixed with sugars and bicarbonate. During heating, the chemical reactions occurring between amino acids and reducing sugars lead to the formation of Maillard reaction products (Maillard 1912; Jousse et al. 2002; Mundt and Wedzicha 2003; Bellarde 2005). Depending on the extent of the reaction, the heating profile, and the amount and type of ingredients used, the Maillard reaction can give rise to several chemical products, such as furosine, 5-hydroxymethylfurfural (5-HMF), pyridosine, maltosine, or maltol (van Boekel 1998; Malec et al. 1999, 2005; Francisquini et al. 2016; Francisquini et al. 2018). These can affect in different degrees the resulting colour formation (i.e. browning) and flavour development (Pinto and Wolfschoon-Pombo 1984; Pavlovic et al. 1994; van Boekel 2006; Newton et al. 2012).

Because the Maillard reaction greatly affects the sensorial characteristics of the final product, there is a certain interest to estimate the extent of the reaction developed during processing. For this purpose, several markers have been proposed. Among those, furosine (Resmini et al. 1990, 2003) and 5-HMF (Keeney and Bassette 1959) are the most studied. Especially, 5-HMF is a highly used marker to determine the extent of the Maillard reaction during heating (Troise 2018).

5-HMF is an intermediate organic compound derived from the dehydration of certain sugars during heating. The formation of 5-HMF may occur at any pH, temperature, and water activity, but its formation is favoured by high temperatures, alkaline pH, and high water activity levels. In these conditions, the Maillard reaction occurs at a much higher rate (Morales and van Boekel 1998; Labuza and Baisier 1992; Rodriguez et al. 2016; d’A Francisquini et al. 2019). For this reason, the determination of 5-HMF is an important quality index in products that may easily undergo over-processing like DL (van Boekel 1998; Pischetsrieder et al. 1999; Kowalski et al. 2013; Ritota et al. 2017).

To date, 5-HMF has been detected in several types of food, including spirits and honey (Murkovic and Bornik...
Bassette 1959; Czerwonka et al. 2020). The formation of derivatives with thiobarbituric acid (TBA) (Keeney and of samples under acidic conditions and in some cases after derivatization with thiobarbituric acid (TBA) (Keeney and Bassette 1959; Czerwonka et al. 2020). The formation of 5-HMF is typically measured using ultraviolet absorbance (UV) at 284 nm (18 000 M−1 cm−1), because of the strong absorption of furfural-like compounds (Gürkan and Altnay, 2015), or at 443 nm for the TBA derivative (Keeney and Bassette 1959). In recent years, also high-performance liquid chromatography (HPLC) gained importance for the detection of 5-HMF in food (Teixidó et al. 2006, 2008; Serra-Cayuela et al. 2013). For HPLC analysis of 5-HMF, typically liquid–liquid extraction is necessary, especially in complex matrices.

A common challenge in the analysis of products with a complex sample matrix is that many compounds naturally present in foods may also interfere with the light absorption of 5-HMF. Although acid digestion with eventual derivatization is commonly used to obtain a stronger absorption of furfural compounds, these procedures are rather time consuming and require extensive work from the operator. Furthermore, when the chromatographic resolution is poor, or the sample matrix is too complex, interferences may adversely affect the quantification of 5-HMF. Such possible lack of accuracy in the determination of 5-HMF has recently received more attention than in the past because of some cytotoxicity (Nässberger 1990), genotoxic (Severin et al. 2010), and tumoral effects (Durling et al. 2009) associated with some 5-HMF derivatives, like 5-chloromethyl and 5-sulfide-methylfurfural. Although the subject is debated (Abraham et al. 2011), the possible safety concerns related to the consumption of 5-HMF have driven the need to develop alternative methods for its quantitation. This applies even in samples containing high concentration of interferences.

The above-mentioned drawbacks can be overcome by using solid-phase extraction (SPE). Specifically, with the use of SPE, sample interferences can be reduced and the resolution of the compounds of interest can be increased. In addition, the SPE procedure can be easily automatized leading to an increase in the number of samples that can be analyzed. As an example, SPE has been used by Resmini et al. (1990) as clean-up step of dairy products to analyze furosine via HPLC–UV for the assessment of the damage induced by the heat. The application then of high-resolution mass spectrometry (MS) as detection technique can exploit its high accuracy to detect 5-HMF by using its exact mass and MS² spectrum.

For the reasons above, this study proposes a rapid method to determine the content of 5-HMF in DL samples based on liquid chromatography coupled to mass spectrometry (LC–MS) after SPE. Different DL samples subjected to increasing thermal treatments were analyzed. The results were compared with the analysis of 5-HMF in DL using previous methods for the determination of 5-HMF in milk products which employ acid digestion and derivatization as pre-processing steps (Keeney and Bassette 1959; Franci­squini et al. 2018; Czerwonka et al. 2020).

Materials and Methods

Chemicals and Reagents

5-Hydroxymethylfurfural (5-HMF) was purchased from Sigma-Aldrich (Steinheim, Germany) at purity higher than 99%. Trifluoroacetic acid (TFA) was purchased from Alfa Aesar, methanol from Honeywell, concentrated hydrochloric acid solution, trichloroacetic acid (TCA), oxalic acid, 2-thiobarbituric acid (TBA), NaCl, 8 mol L⁻¹ NaOH solution, diethyl ether, and LC–MS grade acetonitrile were purchased from Sigma-Aldrich and not purified further. For HPLC methods, ultrasonicated Milli-Q water was employed.

Heated Milk

Heated milk samples were prepared and analyzed to determine the 5-HMF content during heating. Pasteurized semi-skimmed (1.6% fat) milk was obtained from the local market. Milk samples (10 mL each) were transferred to glass vials (Pyrex vials with screwcap) and placed in a thermostatic water bath at 80 °C. After 1, 2, 3, and 4 h, the vials were removed from the water bath and immediately placed on ice. Pasteurized milk without further heat treatment was used as control.

Dulce de Leche Preparation

For the formulation of dulce de leche, pasteurized whole milk obtained from the local market, fructose, and sodium bicarbonate were used. The samples were prepared by mixing 100 mL of milk, 40 g of fructose, and 0.5 g of sodium bicarbonate. Sodium bicarbonate was added to fasten the browning reaction and to prevent protein coagulation by
increasing the pH of the mixture above the isoelectric point of the proteins, such as the casein micelles.

**Extrusion Cooking of Dulce de Leche (DL)**

DL samples were prepared by hot melt extrusion with a co-rotating twin screw extruder with a screw diameter of 11 mm and a length to diameter ratio (L/D) of 40 (Pharma 11 Twin-screw Extruder, Thermo Scientific™). To prepare the DL samples, the screw speed was set to 140 rpm while the mixture was pumped to the extruder with a peristaltic pump at constant flow rate of 0.10 mL min⁻¹. The temperature along the barrel was constant (temperatures: 100, 110, 120, and 130 °C). DL samples were produced at each of the four different temperatures. After each run, the extruder was stopped, opened, cleaned, and assembled again. Then, the temperature for the following condition was set to run for 23 h foreseen by Resmini et al. Then, 0.5 mL of sample was loaded on a preconditioned (15 mL 1:2 MeOH/H₂O v/v) C₁₈ SPE cartridge (V = 6 mL, Thermo Sci.) and eluted with 3-mL hydrochloric acid solution (3 mol L⁻¹) until dryness of the cartridge using air flow with an automated SPE system (GX-271 Aspec, Gilson Italia S.r.l., Italy). One milliliter of the eluate was dried under nitrogen flow at room temperature (MultiVap 8, LabTech S.r.l., Milano, Italy) and dissolved in 1 mL of 5% acetonitrile in deionized water and analyzed by LC-HRMS. The standard error was calculated as standard deviation divided by the square root of number of repetitions.

**Sample Preparation Procedure for Analysis by Liquid Chromatography**

The same number of samples as before were extracted in triplicate. The DL samples were extracted according to Czerwonka et al. to determine 5-HMF using HPLC–UV (Czerwonka et al., 2020). In short, 1 g of DL sample was diluted with 10 mL of water. Three millilitres of diluted sample were mixed with 1.5 mL oxalic acid solution (0.3 mol L⁻¹). After 20 min, 1.5 mL TCA solution (40%, w/v) was added, stirred, and the mixture filtered through a filter paper. Two millilitres of the filtrate was then mixed with 1 mL of saturated NaCl solution and extracted three times with diethyl ether (20, 10, 10 mL). The organic fractions were collected and combined, and the solvent was evaporated under nitrogen stream at temperature of 40 °C (Multivap 8, LabTech S.r.l., Milano, Italy). The residue was dissolved in 1 mL methanol, and 1-mL NaOH solution (1 mol L⁻¹) was added and brought to 10-mL volume with water. The mixture was filtered with a 0.45-μm syringe filter prior to injection into the HPLC system.

**Liquid Chromatography Coupled to High-Resolution Mass Spectrometry (LC-HRMS)**

The determination of 5-HMF in samples extracted by SPE and by the procedure suggested by Czerwonka et al. (2020) was performed by HPLC–UV coupled to HRMS. The system consisted of a Q-Exactive Orbitrap HRMS instrument (Thermo Fisher Scientific) coupled to an Ultimate 3000 UHPLC instrument (Thermo Fisher Scientific) with UV–Vis detector. The separation of the compounds was done at a flow rate of 1 mL min⁻¹ with a LUNA C8 LC column (150 mm × 4.6 mm i.d., 5 μm, 100 A, phenomenex) with a C8 security guard cartridge system (phenomenex). The mobile phase consisted of a combination of solvents A (water with addition of 0.1% trifluoroacetic acid v/v) and B...
(acetonitrile). The gradient was set as follows: 5% B (v/v) for 7.5 min, then from 5% B to 95% B at 8 min, hold until 8.5 min then to 5% B at 9 min followed by a re-equilibration step (5% B) from 9 to 10 min. After each sample, a wash step with a blank (phase A) was introduced with the same chromatographic set-up as before, but with a different gradient: from 5% B at 0 min to 95% B at 1 min, hold 95% until 3 min, from 95% at 3 min to 5% B at 4 min followed by a re-equilibration step (5% B) from 4 to 7 min. For Full-MS analysis, the mass spectrometer was operated in positive ionization mode using the following conditions: sheath gas at 20 (arbitrary units), aux gas at 10 (arbitrary units), aux temperature 50 °C, spray voltage at ±3.5 kV, capillary temperature at 320 °C, and RF S-lens at 65%. The mass range selected was from 50 to 750 m/z with a Full-MS set resolution of 35,000 at m/z 200, AGC target at 2×10^5, and max. injection time of 50 ms. The MS^2 measurements of the selected ions were performed with a resolution of 17,500 and AGC target at 5×10^5. Correlation of chemical compounds relative abundances and integration of the area under each peak (HPLC–MS EIC integrations) was done using Compound Discoverer 3.1 software (Thermo Scientific, Milano, Italy).

**Liquid Chromatography with Diode Array Detector (HPLC–DAD) for Acquisition of Absorption Spectra**

The determination of the absorption maxima of detected compounds was performed using a 1260 Infinity HPLC instrument coupled to diode array detector (DAD) (Agilent Technology). For chromatographic separation, the same gradient conditions were employed as for LC-HRMS. Flow rate of 1 mL/min with a LUNA C8 LC column and security guard were used. The mobile phases included solvents A (water with addition of 0.1% trifluoroacetic acid v/v) and B (acetonitrile), and the following gradient: 5% B (v/v) for 7.5 min, then from 5% B to 95% B at 8 min, hold until 8.5 min then to 5% B at 9 min followed by a re-equilibration step (5% B) from 9 to 10 min.

**Results and Discussion**

**Identification of 5-HMF**

Figure 1 shows the extracted ion chromatogram of 5-HMF obtained from the analysis of a dulce de leche sample by HPLC–MS after sample preparation using SPE. No acid digestion or derivatization of the sample was needed. The chromatogram is characterized by an intense peak with a retention time of 5.85 min. The identification was obtained by comparing its retention time, UV spectrum ($\lambda_{\text{max}} = 284$ nm obtained by HPLC–DAD, Figure 1, inset), and MS spectrum with those reported in previous studies (Keeney and Bassette 1959; Serra-Cayuela et al. 2013; Larrañaga et al. 2016). The absorption maximum and retention of the peak matched those of 5-HMF. Furthermore, a 5-HMF standard was injected to confirm the identity of the peak. The standard peak demonstrated the same properties. Final confirmation was given from the analysis of the full-MS and MS^2 spectrum (Fig. 2). The quasi-molecular ion [M+H]^+ of 5-HMF (m/z of 127.0392) was found with a mass accuracy of $\Delta m_r = -1.57$ ppm for the compound eluting at 5.85 min and the standard. Characteristic fragments like m/z [M+H]+ 109.0290, 81.0343, and 53.0395 matched those of the 5-HMF standard. The origin of such fragments is consistent with the loss of water [M+H-H_2O]^+ (m/z 109.0290) and the successive losses of CO groups [M+H-H_2O-CO]^+ (m/z 81.0343) and [M+H-H_2O-CO-CO]^+ (m/z 53.0395). A similar MS/MS spectrum was previously reported using low-resolution mass spectrometry (Serra-Cayuela et al. 2013), and the fragmentation of furan-5-(diethoxymethyl)-2-furanmethanol was described by Hu and Li (2011). The concentration of 5-HMF was proposed as index of heat treatment.

**Validation of 5-HMF for the Assessment of Heat Treatments of DL Samples**

The analytical performance of the proposed SPE-LC–MS method to determine 5-HMF was next investigated. For the
validation of the method, an analytical standard of 5-HMF was injected at increasing concentrations, measured in triplicate. The choice of mass resolution of 35,000 in Full-MS and 17,500 in MS² was a compromise between the best sensitivity and the highest resolution. Figure 3a shows the ion count signal for increasing concentrations of 5-HMF. For concentrations of 5-HMF in the range between 0.5 and 100 μM, the signal trend is linear ($R^2$ of 0.9998), with a sensitivity of $5.1 \pm 0.2 \times 10^6$ ion counts min μM$^{-1}$ and a precision of 1.8% (Table 1). The limit of detection (LOD) was $1.54 \pm 0.03$ μM and the limit of quantitation (LOQ) was $5.12 \pm 0.09$ μM.

Matrix effects of the sample were investigated by the standard addition method. When a DL sample was spiked with known concentrations of 5-HMF (from 0.2 to 27 μM, Fig. 3a), the resulting signal response gave $R^2 = 0.9995$ and slope not significantly different from the calibration with standard solutions ($p = 0.32 > 0.05$, inset Fig. 3a). The calibration curve obtained with the standard solutions

**Fig. 2** MS spectra of 5-HMF obtained by LC-HRMS. (a) Full-MS spectrum at 5.85-min retention time of DL sample processed with SPE-LC–MS. Identification of 5-HMF with a quasi-molecular ion [M + H]$^+$ m/z = 127.0392. (b) Confirmation of 5-HMF with full-MS spectrum of its analytical standard. (c) MS² spectrum of 5-HMF in DL obtained with higher-energy collisional dissociation (HCD) at 33.33 eV. Fragments of m/z 109.0290, 81.0343, and 53.0395 with proposed structures

**Fig. 3** Validating 5-HMF as a marker for thermal treatment in DL. (a) Extracted ion chromatogram of m/z 127.0392. Spiking DL with increasing concentrations of 5-HMF standard (standard addition method). Inset: calibration using 5-HMF standard solutions (▲) and standard addition method (SAM, ■); (b) UV–Vis chromatogram at 280-nm retention time of DL samples produced at different temperatures

**Table 1** Correlation between the proposed SPE-LC–MS method and the reference methods employed for the determination of 5-HMF in DL

| Method          | Linearity $R^2$ | Precision | Sensitivity | LOD [μM] | Correlation with SPE-LC–MS ($R^2$) |
|-----------------|-----------------|-----------|-------------|----------|-----------------------------------|
| SPE-LC–MS       | 0.9998          | 1.77%     | $5.1 \times 10^6$ ion counts min μM$^{-1}$ | $1.54 \pm 0.03^*$ | 1                                 |
| HPLC–UV         | 0.9997          | 2.09%     | $11.70418$ μAU min μM$^{-1}$ | $2.01 \pm 0.02^*$ | 0.9997                            |
| TBA Derivative UV | 0.9948          | 4.44%     | $0.017$ AU min μM$^{-1}$ | $2.99 \pm 0.62^*$ | 0.970                             |

*Significantly different, $p < 0.05$
could therefore be used to quantify the 5-HMF in the DL samples.

**Comparing the Proposed SPE-LC–MS Method with Reference Methods**

The proposed method for 5-HMF detection in DL with SPE-LC–MS was compared with reference methods for the determination of 5-HMF in dairy products. First, the proposed method was compared with the spectrophotometric method by Keeney and Bassette (1959), which employs acid digestion and derivatization with TBA. 5-HMF standard solutions with increasing concentrations were subjected to acidic digestion and derivatization and determined via spectrophotometer. The concentrations of 5-HMF linearly correlated between both methods ($R^2=0.970$, Table 1). Compared to the proposed SPE-LC–MS method, the spectrophotometric method has a significantly higher LOD of 2.99 ± 0.62 μM and a LOQ of 14.79 ± 2.07 μM ($p=0.0004 < 0.05$). The sensitivity was 0.017 AU min μM⁻¹ and the precision 4.44%.

Next, the proposed SPE-LC–MS method was compared with the method by Czerwonka et al. (2020), which determines 5-HMF in milk using HPLC–UV after acidic digestion and liquid–liquid extraction. To assess the analytical performance of the HPLC–UV system, a calibration curve with increasing concentrations of 5-HMF was constructed. Compared to the proposed SPE-LC–MS method, 5-HMF determination via HPLC–UV resulted in a significantly higher LOD of 2.01 ± 0.02 μM and a LOQ of 6.69 ± 0.05 μM ($p=0.014 < 0.05$). A precision of 2.09% and a sensitivity of 11.70418 μAU min μM⁻¹ were obtained. The correlation with the SPE-LC–MS method was $R^2=0.9997$ as reported in Table 1.

**Assessing the Heat Treatments of DL Samples**

Finally, the SPE-LC–MS method was used to estimate the level of heat treatment in DL produced by hot melt extrusion. Figure 3b shows the effect of the increased temperature during DL production on the resulting formation of 5-HMF. A linear increase of 5-HMF intensity with increasing temperature was observed. The concentration of the analyzed 5-HMF in all the samples was calculated against the previously obtained calibration curve using LC–MS. Table 2 reports the quantification of 5-HMF in each sample, after peak integration. The measured concentrations of 5-HMF ranged from 15.13 ± 0.12 μmol g⁻¹ with a standard error of 0.07 in DL samples prepared at 100 °C to 30.40 ± 0.30 μmol g⁻¹ with a standard error of 0.17 in the samples prepared at 130 °C.

All DL samples were also pre-processed according to Keeney and Bassette (1959) with digestion using TCA and derivatization with TBA and finally measured via spectrophotometer. For all DL samples, the concentration of 5-HMF measured with the reference method gave similar results to the proposed SPE-LC–MS method (Table 2). The concentrations of 5-HMF linearly increased in both methods with the increasing temperatures applied during DL production.

In addition, the DL samples were extracted following the procedure by Czerwonka et al. (2020) and the concentrations of 5-HMF determined using the HPLC–UV system. However, the peak of 5-HMF in the samples extracted with this procedure including acid digestion and liquid–liquid extraction was not well resolved due to the presence of other interferences making the final quantification of this compound difficult (Fig. 4). Processing the DL samples with the proposed SPE method, however, resulted in no interferences in UV.

Finally, the suitability of the method was also tested for milk heated under milder thermal conditions. 5-HMF in milk can be formed either via Amadori products of the Maillard reaction through enolization or through lactose degradation and isomerization (Czerwonka et al. 2020). When milk samples were heated at 80 °C for increasing time, even up to 4 h, the resulting 5-HMF concentration was always below the limit of detection using the SPE-LC–MS method. Thus, the proposed procedure seemed especially suitable for the analysis of dairy products extensively treated with high temperatures. This result was expected as 5-HMF is usually not used directly as marker for heated milk. In order to enable the 5-HMF detection, milk is usually subjected to acid digestion and derivatization with TBA (Keeney and Bassette 1959; Francisquini et al. 2018; Haghani-Haghighi et al., 2019), or acidic digestion followed by liquid–liquid extraction (Czerwonka et al. 2020) as described in the compared reference methods. However, in the case of DL, the

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**Table 2** 5-HMF concentrations in DL samples prepared by hot melt extrusion at different temperatures measured by the reference spectrophotometric method (with TBA-derivative by Keeney and Bassette 1959) and the proposed SPE-LC–MS method

| Temperature [°C] | 5-HMF [μmol g⁻¹ DL] | Proposed SPE-LC–MS method |
|----------------|------------------|--------------------------|
|                | Spectrophotometric method with TBA-derivative | Proposed SPE-LC–MS method |
| 100            | 14.64 ± 1.21     | 15.12 ± 0.13             |
| 110            | 16.13 ± 0.89     | 17.97 ± 0.44             |
| 120            | 29.23 ± 0.32     | 24.10 ± 2.31             |
| 130            | 37.72 ± 2.20     | 30.40 ± 0.31             |
proposed SPE-LC–MS procedure allowed a faster and better clean-up of the samples for the detection of 5-HMF. The method took less than 45 min to pre-process the samples compared to the pre-processing steps of the reference methods, which take over 60 min.

Conclusions

This study proposes a rapid method to assess the thermal treatments of spreadable dairy creams like DL using SPE-LC–MS to directly detect 5-HMF as marker. The proposed procedure allowed a faster and better clean-up of the DL samples for the detection of 5-HMF compared to reference methods, which employ spectrophotometric detection. Although using a spectrophotometer results in a faster analysis of 5-HMF, the proposed method allowed chromatographic separation of the 5-HMF peak, resulting in a better resolution. Additionally, for DL, the use of SPE allowed the removal of interferences, otherwise present when employing liquid–liquid extraction. Finally, the use of HRMS ensured the accurate identification of the marker in DL samples. The LC-HRMS system allowed to determine 5-HMF in the concentration range from 0.52 to 103 µM with a limit of detection of 1.54 ± 0.03 µM and a limit of quantitation of 5.12 ± 0.09 µM, precision of 1.77%, and sensitivity of 5.1 ± 0.2 × 10⁶ ion counts min µM⁻¹. Overall, for products produced with high contents of sugar and extensive heating, such as DL, the quantitation of 5-HMF using the proposed method represents an advantage over previous methods, thanks to the high resolution and sensitivity of LC-HRMS. This confirmed the possibility to directly use 5-HMF as index of the level of the heat treatment in DL. Future works are needed to evaluate the proposed approach with other food products especially those with high levels of 5-HMF formation.

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Declarations

Ethical Approval

This article does not contain any studies with human or animal subjects.

Informed Consent

Informed consent is not applicable.

Conflict of Interest

All authors declare no conflict of interest.

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