



Free radical scavenging kinetics of Maillard reaction products: A glucose-glycine model system

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ABSTRACT

This study investigated the free radical scavenging kinetics of Maillard reaction products (MRPs) generated from a glucose-glycine model system under controlled conditions (90 °C, pH 6.8) using a stopped-flow DPPH assay. Kinetic analysis revealed a time-dependent increase in antioxidant activity, highlighting the dynamic formation of reactive antioxidants during the Maillard reaction. Advanced analytical techniques, including high-performance liquid chromatography coupled with coulometric array detection (HPLC-CAD) and high-resolution mass spectrometry (HPLC-HRMS), were used to tentatively identify two key antioxidant MRPs: methylpyridin-3-ol (MP), 3,5-dihydroxy-6-methyl-2,3-dihydro-4H-pyran-4-one (DDMP) and its furanone isomer. The electrochemical properties of these compounds were evaluated, revealing their electron-donating abilities. A kinetic model was developed to simulate and predict the formation of these antioxidants, providing insights into the relationship between antioxidant reactivity and reaction progress. These findings contribute to our understanding of the formation of antioxidants during the Maillard reaction and offer a foundation for the development of natural antioxidants, which is aligned with the growing demand for clean-label food products.

1. Introduction

The Maillard Reaction (MR) plays a fundamental role in enhancing sensory attributes and preserving food quality. Maillard Reaction Products (MRPs) contribute to desirable characteristics, such as the toasted aroma of baked bread (Tang et al., 2024) and the golden-brown hue of grilled meat (Sun et al., 2023). In addition to these sensory qualities, MRPs exhibit significant antioxidant properties due to the formation of compounds, such as Amadori products and α -dicarbonyl compounds, which either possess reductone structures or can tautomerize into reductones, functioning similarly to ascorbic acid (Feng et al., 2022; Nooshkam et al., 2019; Shakoob et al., 2022). This antioxidant properties are connected to the development of flavours and aromas during the MR, in particular with umami flavour (Habinshuti et al., 2019; M. Yu et al., 2018). The antioxidant activity helps inhibit lipid oxidation and prevents the degradation of essential nutrients through various mechanisms, including radical scavenging and metal chelation (Feng et al., 2022; Kanzler et al., 2016). This dual

functionality, both sensory and antioxidant, makes MRPs valuable in preserving food quality during processing and storage (Han et al., 2022; Nooshkam et al., 2019; Sun et al., 2022). Moreover, recent interest in MRPs has grown due to their potential role in developing 'clean label' food products, which prioritize the use of natural compounds over synthetic additives for preservation (Jia et al., 2023; Kitts, 2021; Mildner-Szkudlarz et al., 2023; Monti et al., 1999). Understanding MRPs role in both sensory enhancement and antioxidant activity is therefore a key-strategy to develop natural food preservation techniques.

Despite significant progress in characterizing the sensory benefits of MRPs, the mechanisms underlying their antioxidant reactivity remain poorly understood (Cao et al., 2022; Feng et al., 2022; Sohail et al., 2022). Antioxidants prevent oxidative reactions that degrade essential nutrients, such as unsaturated fatty acids, vitamins, and pigments. However, a common misconception is that antioxidant activity is often assessed by concentration, whereas it is more accurately determined by the rate of radical scavenging reactivity (Angeli et al., 2021). This misconception likely arises from the widespread use of assays that

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express antioxidant activity in terms of equivalent concentration, such as the thiobarbituric acid (TBA) assay, thiobarbituric acid reactive substances (TBARS) assay, and the β -carotene bleaching assay, which focus on the inhibition of lipid oxidation. Additionally, scavenging capacity is also expressed in terms of equivalent concentration through assays, such as 2,2-diphenyl-1-picrylhydrazyl (DPPH) (Foti, 2015), and oxygen radical absorption capacity (ORAC) (Asma et al., 2024). While these methods are useful, they may not fully capture the reactivity of antioxidants to neutralize radicals. Thus, antioxidant reactivity, rather than concentration alone, is crucial for determining their effectiveness (Amorati et al. 2018).

This study investigated the antioxidant reactivity of MRPs formed in a glucose-glycine model system to evaluate their formation and reactivity under controlled conditions (90 °C, pH 6.8). The objectives of this study are to (1) evaluate the antioxidant efficiency of MRPs using kinetic-based DPPH assays; (2) identify the most potent antioxidants through high-performance liquid chromatography (HPLC) coupled with coulometric array detection (CAD); (3) elucidate their electrochemical properties and tentatively identify their chemical structures using high-resolution mass spectrometry (HRMS); and (4) propose a kinetic model that predicts antioxidant formation under the experimental conditions.

This research could offer new insights into the mechanisms of antioxidant reactivity in MRPs, potentially contributing to the development of food products with improved nutritional and functional properties. Furthermore, it could offer a natural alternative to synthetic additives, in line with the growing demand for clean-label food products.

2. Experimental methods

2.1. Chemicals and reagents

All chemicals and reagents used in this study were of analytical grade. Sodium hydroxide, alpha-D-glucose ($\geq 98\%$), fluorescein, Trolox, gallic acid, ascorbic acid, and potassium phosphate monobasic were obtained from Sigma-Aldrich (USA). Glycine ($\geq 98.5\%$) was supplied by Alfa Aesar (USA) and potassium phosphate dibasic trihydrate were obtained from Amresco (USA). Mass spectrometry-grade methanol, acetic acid, deuterated water (D_2O), and trimethylsilylpropanoic acid were procured from Merck (Germany). Milli-Q water was produced using a Sartorius arium® mini system (Germany).

2.2. Maillard reaction model

The Maillard reaction (MR) was studied using a glucose-glycine model system. Equimolar solutions (200 mmol/L) of glucose and glycine were prepared in 0.1 mol/L phosphate-buffered saline (PBS) at pH 6.8, following the protocol described by (Martins & Van Boekel, 2005). The solution was divided into 9 Pyrex flasks with a maximum volume of 15 mL, sealed with hermetic stoppers, and stored overnight at 4 °C for equilibration. In each vial, 10 mL of the solution was poured, and the MR was initiated by immersing the 9 flasks in a water bath at 90 °C for 4 h, with a sample (1 vial) taken every 30 min. The samples were rapidly cooled in an ice bath to prevent further reaction and were stored at -80 °C to preserve their chemical integrity prior to analysis. All experiments were performed in triplicate.

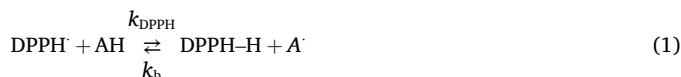
2.3. Stopped flow DPPH assay (DPPH-kin)

The antioxidant kinetics were determined using a stopped-flow system (RX2000 Applied Photophysics, UK) coupled with a Cary 60 UV-Vis spectrophotometer (Agilent Technologies, USA). A solution of 200 $\mu\text{mol/L}$ DPPH• in methanol/water (80:20) was mixed with an identical volume of an antioxidant standard (ascorbic acid or Trolox) or MRP samples, both at a concentration of 60 $\mu\text{mol/L}$ gallic acid equivalents (GAE) quantified using Folin-Cialteau method as described by Angeli (Angeli et al., 2024). The two solutions were rapidly mixed (mixing time

less than 10 ms) and transferred into the flow cell, with a final concentration of 100 $\mu\text{mol/L}$ DPPH• and 30 $\mu\text{mol/L}$ GAE in the reaction mixture. Absorbance at 515 nm was measured every 18 ms. The molar extinction coefficient (ϵ_{515}) of DPPH• in methanol was $11,200 \pm 400 \text{ mol}^{-1} \text{ L cm}^{-1}$.

Since ascorbic acid and trolox react with Folin reagent in a ratio of approximately 0.5–0.6 and gallic acid equivalent ~ 1 (Everette et al., 2010), the 30 GAE $\mu\text{mol/L}$ of MRPs corresponds to approximately 60 $\mu\text{mol/L}$ of ascorbic acid. For this reason, MRPs were diluted to a concentration of ~ 30 GAE $\mu\text{mol/L}$ prior to analysis, and reached a concentration of ~ 15 – $18 \mu\text{mol/L}$ in the reaction mixture.

The reaction kinetics were simulated using Copasi® software (version 4.34) with a three-step mechanism (eqs. (1)–(3)) (Friaa & Brault, 2006):



The three steps mechanism was developed based on literature (Angeli et al., 2021; Foti & Daquino, 2006; Foti et al., 2004) and it is suitable to study the kinetics of reaction between DPPH• and complex samples like MR samples. However, it could be simplified for single and fast standard antioxidants like ascorbic acid or Trolox, where the main reaction (eq. (1)) is enough to describe the kinetics of the oxidation, since no side or equilibrium reactions occur.

Parameters k_{DPPH} , k_{side} , and k_{term} were optimized through iterative fitting using an evolutionary strategy (SRES) program. Fitting was performed using 200 generations and 24,000 function evaluations. All analyses were performed in triplicate, and the results were averaged for data presentation.

2.4. High performance liquid chromatography

HPLC analysis was performed using a 1260 Infinity HPLC system (Agilent Technologies, USA) equipped with a Kinetex Biphenyl column (100 \times 2.1 mm, 2.6 μm particle size) and a pre-column. The mobile phases consisted of water with 0.5% acetic acid (v/v) (A) and methanol with 0.5% acetic acid (v/v) (B). The following gradient was used at a flow rate of 0.3 mL/min: 2% B from 0 to 5 min, 3% B from 5 to 9 min, 5% B from 9 to 16 min, and 2% B from 16 to 20 min. The injection volume was 10 μL , and samples were diluted 1:40 with phase A and filtered with polytetrafluoroethylene (PTFE) 0.2 nm filters prior to analysis.

Three detectors were used in parallel for sample analysis: a diode array detector (DAD), CoulArray detector (CAD), and high-resolution Q Exactive Orbitrap mass spectrometer (HRMS).

2.4.1. Diode array detector

DAD monitored absorbance from 210 to 600 nm, with a focus on the 280 nm absorption band, which corresponds to phenolic compounds and Maillard reaction products. The chromatograms were compared with the CAD and HRMS data.

2.4.2. Coulometric detector

Coulometry, an absolute electrochemical method, was employed using a 16-cell CoulArray detector (Thermo Fisher Scientific, USA). Each cell was set at increasing potentials, ranging from -200 mV to $+650$ mV in 50-mV increments to completely oxidize the analytes. Analyte concentrations were quantified based on the total charge transferred using Faraday's law (Razem et al., 2022). CAD chromatograms were collected and processed using CoulArray data station 3.10 and R software (version 4.1.2). The data were analyzed using the MALDIquant R package for

peak identification, peak height, and area calculations, as previously described (Ding et al., 2022).

2.4.3. High-resolution mass spectrometry

Tentative identification of antioxidants was performed using a Q-Exactive Orbitrap mass spectrometer (Thermo Fisher Scientific, USA). The electrospray ionization source (ESI) was operated in positive ionization mode with a capillary voltage of 4.00 kV, and a capillary temperature of 300 °C. The full MS scan was acquired from 50 to 750 m/z with a resolution of 70,000, an automatic gain control (AGC) target of 1×10^6 and maximum injection time of 100 ms. An inclusion list of targeted species was defined based on the full HRMS scan and used to run a data-dependent MS/MS analysis (MS/MS) to obtain the fragmentation patterns of each target specie. In this case, the AGC target was set at 5×10^5 , the maximum injection time was 50 ms, and the resolution was 17,500 with an isolation window of 4.0 m/z. The collision energy used for the fragmentation was 15 eV. The spectrometer was calibrated every 48-h with Pierce LTQ Velos ESI-positive calibration solution (Thermo Fisher Scientific). The data were collected with Chromeleon 7.2 and analyzed with Xcalibur 4.3, Compound Discoverer 3.3 software, and Mass Frontier 7.0 software (Thermo scientific Scientific, MA, USA).

2.5. Hydrodynamic voltammogram

Hydrodynamic voltammetry was performed using CAD to measure the redox properties of the MRPs. The half-wave potential ($E_{1/2}$) and maximum charge (Q_{max}) were derived from the HDV, providing insights into the electron-donating capacity of MRPs. $E_{1/2}$ is indicative of antioxidant potential, with lower values correlating with stronger electron-donating capacity (Razem et al., 2022).

2.6. Nuclear magnetic spectroscopy

Acetic acid and formic acid were quantified using ^1H NMR, while glucose and glycine were measured using ^{13}C NMR due to signal overlap in ^1H NMR spectra. All spectra were acquired using a Jeol JNM-ECZ600R/M3 spectrometer at 25 °C. The quantification of each compound was performed by integrating the respective peaks and comparing them with an internal standard (TSP). Calibration curves for glucose and glycine were generated using concentrations ranging from 50 to 250 mmol/L, with an R^2 of 0.997 and 0.989, respectively.

2.7. Determination of melanoidins

Melanoidins were quantified at 470 nm using a microplate reader (Tecan Infinite M Nano+, Switzerland) following the method described by (Martins & Van Boekel, 2003). The extinction coefficient ($\epsilon_{470} = 0.64 \text{ mol}^{-1} \text{ L cm}^{-1}$) and optical path length of 0.556 cm were used to calculate the molar melanoidin concentrations.

2.8. Kinetic analysis

The kinetic model for the Maillard reaction was developed based on data collected from glucose, glycine, acetic acid, formic acid, and melanoidin measurements. The model was fitted to the experimental data using Copasi® software, which minimized the residuals between the experimental and simulated data using an evolutionary programming algorithm. The model accuracy was evaluated using the R^2 value, which represents the explained variance.

2.9. Statistics

All statistical analyses (DPPH kinetic assay and MR kinetic studies) were performed using Copasi® software. Results are presented as means \pm standard deviation, and statistical significance was considered at $p < 0.05$.

3. Results and discussion

3.1. Antioxidant kinetic rate and capacity

This study investigated the formation and reactivity of antioxidants in Maillard Reaction Products (MRPs) using a stopped-flow technique, providing insights into their free radical-scavenging kinetics. Fig. 1a illustrates the stopped-flow apparatus, and Fig. 1b shows the rapid consumption of DPPH• radicals (100 $\mu\text{mol/L}$) by MRPs (30 $\mu\text{mol/L}$ GAE) after a 4-h reaction at 90 °C in phosphate buffer (pH 6.8). For comparison, the inset shows the DPPH• radicals consumption at equimolar concentrations of ascorbic acid and Trolox, two well-known water-soluble antioxidants.

3.1.1. Antioxidant kinetic rate

MRPs have high free radical-scavenging kinetics, as demonstrated by the rapid decrease in DPPH• radical concentration reported in Fig. 1b. The inset shows a comparison of fast DPPH• reduction by MRPs, Trolox, and ascorbic acid within 5 s. The rate constant k_{DPPH} quantifies the reactivity of antioxidants, providing insights into the effectiveness of MRPs in inhibiting oxidative processes. Recent studies have highlighted the utility of k_{DPPH} values for not only comparing antioxidant reactivity (Shojaee et al., 2022), but also screening for their efficacy in specific food matrices (Angeli et al., 2021), such as the anti-browning mechanism of different apple cultivars (Angeli et al., 2024).

The experiments were designed with antioxidants as the limiting reagents to ensure that only a portion of excess DPPH• was consumed. This setup accurately captures the reaction kinetics using the mechanism given in eqs. (1)–(3). The main reaction (eq. (1)) involves electron transfer between the antioxidant (AH) and the DPPH• radicals to produce the oxidized antioxidant (A•). Although theoretically reversible, the reverse reaction is typically slow for highly effective antioxidants, as noted by Foti et al. (Foti & Daquino, 2006). In addition, side reactions between A• and residual DPPH• (k_{side}) and termination processes via A• dimerization or disproportionation (k_{term}) significantly contribute to the overall mechanism, in particular when dealing with complex extracts, where the main reaction cannot describe completely the kinetics. In fact, in the reaction of DPPH• with the extracts, since these are multicomponent solutions, there are interactions and equilibria that cannot be adequately described with a simple model than include just the main reaction (k_{DPPH}). These complexities emphasize the need for quantitative kinetic studies rather than reliance solely on antioxidant capacity assays (Angeli et al., 2021; Foti et al., 2004).

The rate constant (k_{DPPH}) for the reaction between MRPs and DPPH• radicals was determined to be $6000 \pm 300 \text{ M}^{-1} \text{ s}^{-1}$ in methanol/water (80:20). Although this value is approximately half that of ascorbic acid ($k_{\text{DPPH}} = 14,450 \pm 750 \text{ mol}^{-1} \text{ L s}^{-1}$), it still indicates substantial antioxidant activity. Both MRPs and ascorbic acid had significantly higher k_{DPPH} values than Trolox ($k_{\text{DPPH}} = 1770 \pm 90 \text{ mol}^{-1} \text{ L s}^{-1}$, $p < 0.05$), indicating greater reactivity. Furthermore, MRPs exhibited higher k_{DPPH} values than other antioxidants, such as rutin, ellagic acid, and quercetin, as reported in previous studies (Angeli et al., 2021; Foti et al., 2004).

The k_{DPPH} for Trolox observed in this study is consistent with the literature values, ranging from $320 \text{ mol}^{-1} \text{ L s}^{-1}$ in pure methanol (Santosh Kumar et al., 2002) to approximately $10^4 \text{ mol}^{-1} \text{ L s}^{-1}$ in ethanol/water mixtures, with variations depending on the water content, which facilitates electron transfer (Friaa & Brault, 2006). The k_{DPPH} value of Trolox measured in this study in methanol/water (80:20) was within this range, indicating the reliability of the experimental approach.

3.1.2. Antioxidant capacity

MRPs exhibited similar antioxidant capacity to ascorbic acid when evaluated via the stoichiometry of their DPPH• reaction. The stoichiometry coefficient n was calculated as $n = \Delta(\text{Abs})/(\Delta\epsilon \times \text{L} \times [\text{AH}])$, where $\Delta(\text{Abs})$ is the absorbance change recorded at the end of the

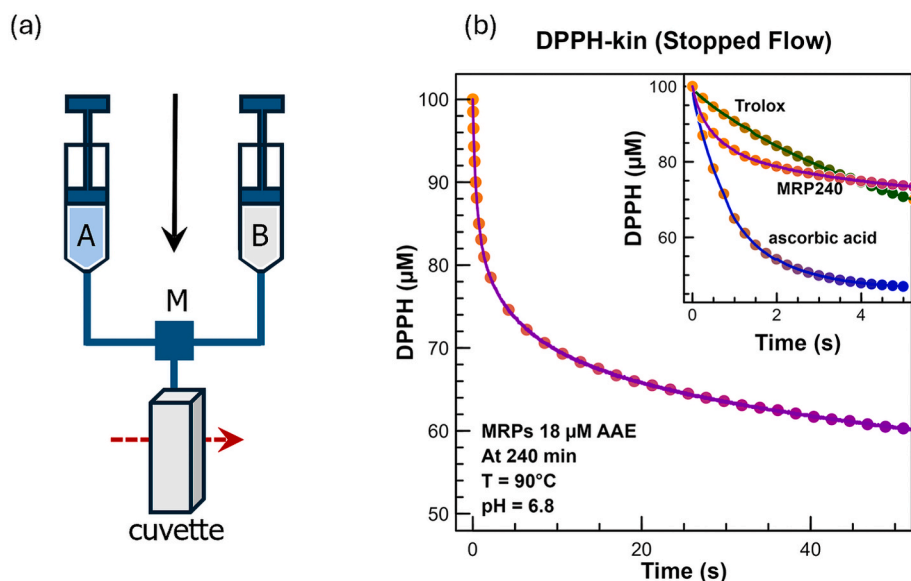


Fig. 1. Results of the kinetic DPPH assay using a stopped-flow technique. (a) Stopped-flow apparatus, including a driving motor, syringes A (filled with DPPH•) and B (filled with the antioxidant), mixer (M), and UV cell. (b) Experimental and simulated kinetic traces of DPPH• reduction by Maillard reaction products. Inset: Zoom of the DPPH• decay within 5 s for MRPs, ascorbic acid, and Trolox. AAE: ascorbic acid equivalent.

reaction, $\Delta\epsilon$ is the molar extinction coefficient of DPPH•, L is the optical path length, and $[AH]$ is the effective antioxidant concentration. At longer reaction times, when the absorbance signal of DPPH• reaches a steady state, $\sim 18 \mu\text{mol/L}$ ascorbic acid equivalent of MRPs neutralize approximately $30\text{--}40 \mu\text{mol/L}$ of DPPH• radicals, yielding $n \approx 1.8\text{--}2.0$. Similarly, $30 \mu\text{mol/L}$ of ascorbic acid or Trolox neutralize $55\text{--}60 \mu\text{mol/L}$ of DPPH• radicals, giving $n \approx 2.0$, indicating a similar antioxidant capacity of MRPs (at pH 6.8).

The mechanism behind ascorbic acid's stoichiometry involves transferring one electron to DPPH•, forming semi-dehydroascorbic acid, which further reacts with DPPH• to form dehydroascorbic acid (Kamrul et al., 2016). Similarly, Trolox, a vitamin E analog, reacts with DPPH•, forming a Trolox radical that subsequently reacts with another DPPH•

radical to complete the neutralization process. The stoichiometry coefficients of Trolox and ascorbic acid are consistent with prior research (Asma et al., 2024).

The apparent stoichiometry value of the MRPs indicates that 1 mol of MRP neutralizes approximately 2 mol of free radicals indicating that MRPs effectively scavenge free radicals based on k_{DPPH} , and they exhibit a similar antioxidant capacity to ascorbic acid or Trolox. Nevertheless, the high reactivity of MRPs confirm that they play a valuable role in broader antioxidant strategies. These results give an idea of the antioxidant behaviour of MRPs, even if it does not allow the complete characterization of the compounds present in the samples, since they were not separated or isolated. Future studies should explore the antioxidant kinetics of single compounds produced through MR and

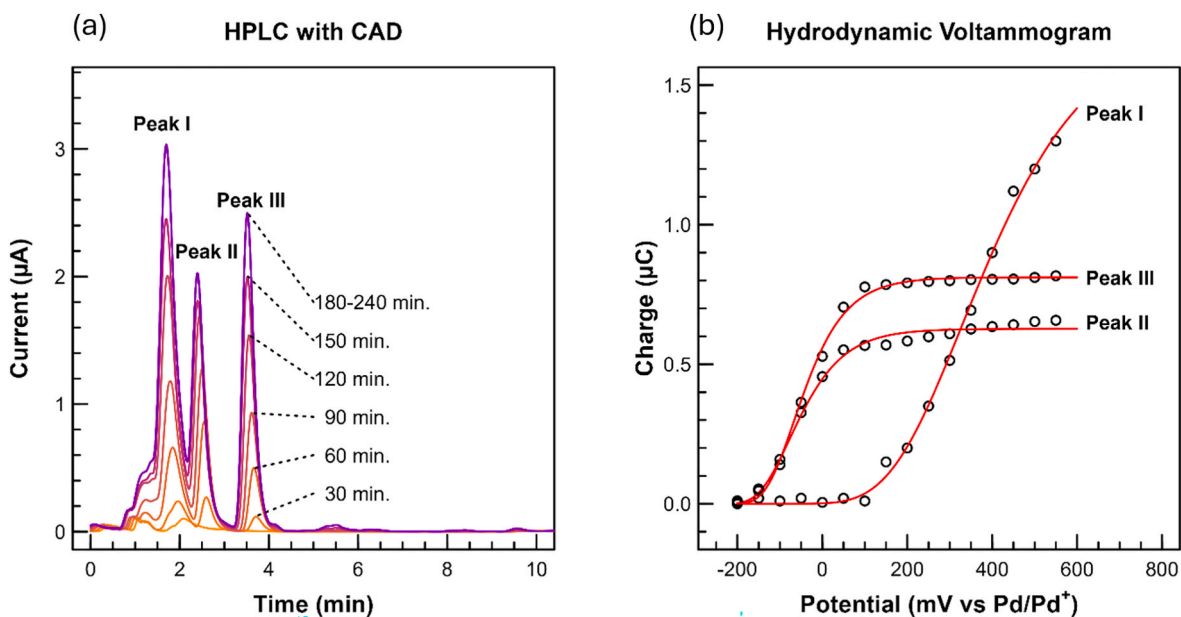


Fig. 2. (a) Total current (μA) chromatograms of MRP recorded over 0–240 min of constant heating at 90°C . Peaks I, II, and III correspond to distinct MR products. (b) Hydrodynamic voltammograms (HDVs) of Peaks I, II, and III. The HDVs depict the charge (μC) as a function of the applied potential (mV vs. Pd/Pd + pseudo-reference electrode) for the three peaks identified in the HPLC-CAD chromatograms of the MR samples.

potential synergistic action of these compounds with other antioxidants to enhance overall food stability.

3.2. Electrochemical screening of maillard reaction products

The reactivity of MRPs in scavenging free radicals, as determined by the DPPH assay in hydroalcoholic solvents (pH 6.8), is attributed to their electron-transfer capacity (Gulcin & Alwaseel, 2023). This was further investigated using high-performance liquid chromatography (HPLC) coupled with a coulometric array detector (CAD). Fig. 2a shows three main anodic peaks were detected at retention times of 1.5 min (Peak I), 2.25 min (Peak II), and 3.4 min (Peak III). The short retention times of these peaks, observed on a C18 column, suggest that the electroactive species formed in the MRPs are primarily hydrophilic, consistent with the findings of Zielińska on antioxidants in bread crusts (Zielińska et al., 2022).

Only Peaks II and III exhibited significant antioxidant activity as shown in the hydrodynamic voltammograms in Fig. 2b, recorded at pH 3.0. Peaks II and III displayed the lowest half-wave potentials ($E_{1/2}$), indicating strong electron-donating abilities. Peaks II and III had $E_{1/2}$ values of -50 mV vs. Pd reference electrode (equivalent to $+507$ mV vs. the standard hydrogen electrode (SHE)) was significantly lower than Peak I ($E_{1/2}$ of $+957$ mV vs. SHE, $p < 0.05$), demonstrating superior antioxidant potential.

In electrochemical analyses, lower $E_{1/2}$ values correlate with enhanced electron donation, a key trait of efficient antioxidants (Chevion et al., 2000). The low $E_{1/2}$ values of Peaks II and III, compared with those of ascorbic acid ($E_{1/2} = +657$ mV vs. SHE at pH 3.0) (Ding et al., 2022; Gazdik et al., 2008; Razem et al., 2022) suggest that these species are potent antioxidants.

3.3. Tentative identification of antioxidants in maillard reaction products

The mass spectra of Peaks II and III show the same molecular ion with a mass-to-charge ratio (m/z) of 145.0496, suggesting two similar molecules with different retention times. The fragmentation pattern exhibited a major peak at m/z 71.0492, indicating of a stable fragment. Other fragments (m/z 127.0390, 99.0441, and 71.0492) align with water and carbon monoxide losses. Studies on 1-deoxyglucosone (1-DG) degradation under neutral or alkaline conditions have reported the formation of 3,5-dihydroxy-6-methyl-2,3-dihydro-4H-pyran-4-one (DDMP) and its furanone isomer (Chen et al., 2021; Li et al., 2019). The chemical structure reported in Fig. 3 is consistent with the observed fragmentation patterns of Peaks II and III. DDMP is also the major product of the reaction of glucose and proline (Mori & Ito, 2004) and is found in several foods (Qiao et al., 2022).

UV spectral analysis (Supporting information S1) revealed absorption maxat 290 nm for both Peaks II and III, consistent with an enone group and suggesting a structure containing ketone and double bonds. The pyranone and furanone isomers likely contributed to the antioxidant properties observed in the DPPH-kin assays, consistent with their

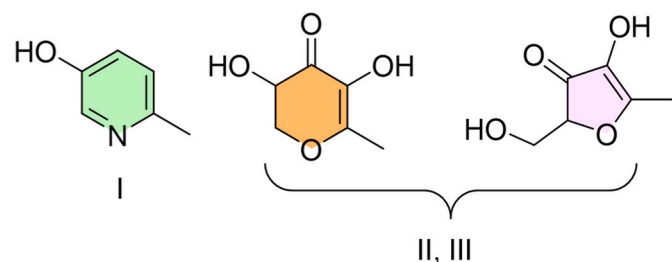


Fig. 3. Structures assigned through HRMS to the three peaks exhibiting antioxidant activity. Peak I, in green, corresponds to 6-methylpyridin-3-ol (MP), while peak II (orange) and peak III (violet) correspond to 3,5-dihydroxy-6-methyl-2,3-dihydro-4H-pyran-4-one (DDMP) and its furanone isomer.

low half-wave potential. This finding aligns with studies showing that reductone structures, like DDMP, facilitate electron donation and neutralize free radicals (Chen et al., 2021; Kanzler et al., 2016; X. Yu et al., 2013).

Peak I exhibited a molecular ion at m/z 110.0601. Fragments (m/z 94.0530, 82.0650, and 66.0420) suggested methyl groups and carbon monoxide loss. The molecular formula of the compound was identified as C_6H_7NO , and its structure, shown in Fig. 3, was proposed as 6-methylpyridin-3-ol (MP), based on mass spectral databases (m/z Cloud and ChemSpider). Peak I exhibited modest antioxidant activity, consistent with the well-known weak reactivity of pyridine derivatives with radicals (Kanzler et al., 2016, 2017; Yanagimoto et al., 2002).

3.4. Kinetic modeling of antioxidant evolution during the maillard reaction

To explain the formation of antioxidants during the Maillard reaction, a kinetic model was developed based on monitoring key intermediates using NMR spectroscopy. Fig. 4 presents the NMR traces showing the consumption of 200 mmol/L glucose and glycine, along with the formation of acetate, formate, and melanoidins over time. Notably, the pH dropped from 6.8 to 5.2, indicating increased acid formation, consistent with previous findings (Mori & Ito, 2004).

This model simplifies and builds upon previously reported mechanisms (Lund & Ray, 2017; Martins & Van Boekel, 2005). As shown in Table 1, the model captures key pathways of the Maillard reaction, including glucose isomerization to fructose, higher glucose consumption relative to glycine, and greater formation of acetic acid compared to formic acid and melanoidins. The observed increase in fructose concentration (2–5 mmol/L) is attributed to glucose isomerization (Delidovich, 2023), whereas the higher glucose consumption relative to glycine, as demonstrated in Fig. 5a, results from the regeneration of glycine from Amadori products (Martins & Van Boekel, 2005).

Melanoidins formation at 4 mmol/L as reported in Fig. 5b, was confirmed through visible spectroscopy (extinction coefficient = $0.64 \text{ mol}^{-1} \text{ L cm}^{-1}$) (Martins & Van Boekel, 2003). Acetic acid concentrations consistently exceeded those of formic acid, indicating the predominance of the 2,3-enolization pathway. The minimal formation of formic acid and the absence of significant 3-deoxyglucosone (3-DG) degradation product 5-hydroxymethylfurfural (HMF) (Lee et al., 2022), suggest that the 1,2-enolization pathway is negligible under these conditions (Yaylayan & Keyhani, 2000). The neutral pH, which disfavors 3-DG formation, further supports this conclusion.

The iterative fitting procedure described in the experimental section was employed to refine the kinetic model using nonlinear regression. The strong correlation between simulated and experimental values ($R^2 > 0.99$) confirms that the model can describe the Maillard reaction.

The key steps in the kinetic model are summarized as follow.

- 1. Glucose Degradation:** Glucose degrades into formic acid (FA) and acetic acid (AA) with a rate constant $k_1 = 0.733 \text{ s}^{-1}$, comparable to values reported by Martins and Van Boekel (2005) ($p > 0.05$).
- 2. Glucose Isomerization:** Glucose isomerizes to fructose in a reversible reaction with the forward and backward rate constants k_{2f} and k_{2b} , respectively. The high backward rate constant (88 s^{-1}) indicates limited isomerization after 4 h of heating.
- 3. Amadori Product Formation:** Glucose and glycine react to form Amadori rearrangement products (ARP) with a fixed rate constant $k_3 = 0.097 \text{ mol}^{-1} \text{ L s}^{-1}$. These ARPs eventually degrade to produce antioxidant species.
- 4. 1-DG Formation:** ARP breaks down to 1-deoxyglucosone (1-DG) while regenerating glycine. This step is crucial because 1-DG serves as a precursor for antioxidant species and melanoidins.
- 5. Further Degradation of 1-DG:** 1-DG was further degraded to acetic acid and erythrose with a rate constant $k_5 = 383 \text{ s}^{-1}$.

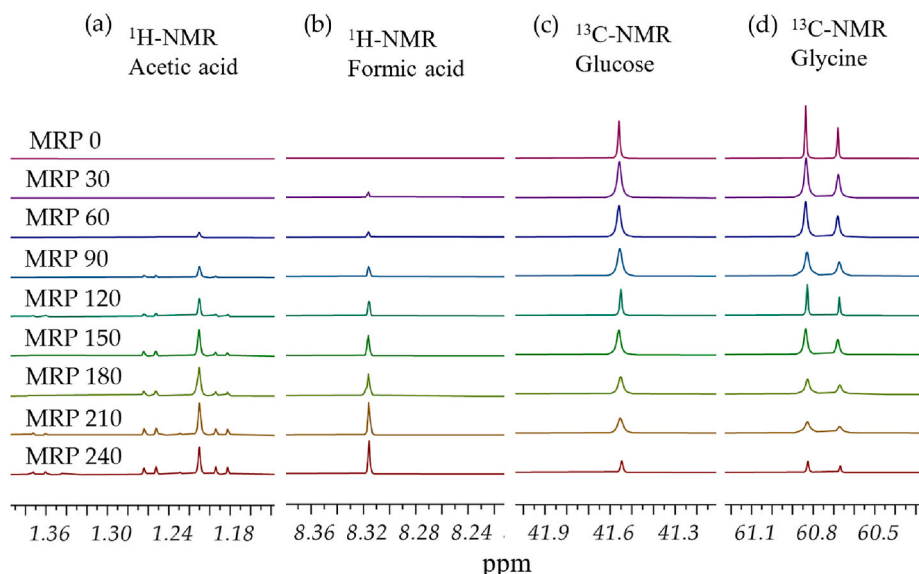


Fig. 4. (a) Stacked $^1\text{H-NMR}$ peak corresponding to acetic acid obtained from MR samples incubated for 0, 120, and 240 min at 90 °C; (b) Stacked $^1\text{H-NMR}$ peak corresponding to formic acid obtained from MR samples incubated for 0, 120, and 240 min at 90 °C; (c) Stacked $^{13}\text{C-NMR}$ peak corresponding to glucose obtained from MR samples incubated for 0, 120, and 240 min at 90 °C; (d) Stacked $^{13}\text{C-NMR}$ peak corresponding to glycine obtained from MR samples incubated for 0, 120, and 240 min at 90 °C.

Table 1

Rate constants relative to the kinetic model of MR reported by Martins & Van Boekel, 2005, and obtained fitting the same model to the data measured in this work and the %RDS. Abbreviations: formic acid (FA); acetic acid (AA); Amadori products (ARP); 1-deoxyglucosone (1-DG); browning products, such as melanoidins (B).

Rxn.	Elementary step	Symbol	Units	(Martins & Van Boekel, 2005) $\times 10^6$	This work $\times 10^6$	RSD (%)
(1)	glucose \rightarrow FA + AA	k_1	s^{-1}	0.716	0.733	2
(2)	glucose \rightleftharpoons fructose	k_{2f}	s^{-1}	6.83	0.016	ind.
(3)	glucose + glycine \rightarrow ARP	k_{2b}	s^{-1}	58.3	88	29
		k_3	$\text{mol}^{-1} \text{L s}^{-1}$	0.097	0.097	–
(4)	ARP \rightarrow 1-DG + glycine	k_4	s^{-1}	111	225	4
(5)	1-DG \rightarrow AA + erythrose	k_5	s^{-1}	383	383	–
(6)	1-DG + glycine \rightarrow brown products	k_6	$\text{mol}^{-1} \text{L s}^{-1}$	320	0.038	2
(7)	1-DG \rightarrow AH	k_7	s^{-1}		1480	6
(8)	AH \rightarrow brown products	k_8	s^{-1}		113	12

Note: when RSD is missing (–), the rate constant is fixed as in the VB calculation.

6. Melanoidin Formation: 1-DG and glycine react to form melanoidins, contributing to the characteristic browning and antioxidant activity of Maillard products. The rate constant in this study was estimated at $k_6 = 0.038 \text{ mol}^{-1} \text{L}^{-1}$, significantly lower than previously reported values (Martins & Van Boekel, 2005), possibly due to the oxidation of antioxidants (see Step 8).

Two additional key steps were included to describe the formation of antioxidants during the Maillard reaction.

7. Antioxidant Formation: 1-DG is converted into antioxidant species (AH) in reduced form ("red"), with a rate constant $k_7 = 1480 \text{ s}^{-1}$. These antioxidants play critical roles in the radical scavenging properties of MRPs, with DDMP and its furanone tautomer being the most significant contributors, as previously described.

8. Antioxidant Degradation: The degradation of antioxidant species (AH) occurs with a rate constant $k_8 = 113 \text{ s}^{-1}$. The balance between formation and degradation results in a production curve in which antioxidant levels plateau as degradation begins to dominate. This process also contributes to non-enzymatic browning.

Fig. 5c illustrates the trend in antioxidant species formation predicted by the model, which aligns with HPLC-CAD analysis results. The model's accuracy was further demonstrated by its strong correlation with experimental data ($R^2 > 0.99$). These findings indicate that optimizing the Maillard reaction during food processing can enhance antioxidant production, leading to the development of food products with improved stability.

4. Conclusions

This study investigated the formation, kinetic behavior, and antioxidant activity of Maillard reaction products (MRPs) generated from the reaction between glucose and glycine under controlled conditions (90 °C and pH 6.8). The results revealed significant consumption of glucose and glycine, leading to the formation of key MRPs, including acetic acid, formic acid, and melanoidins.

A simplified kinetic model was developed and validated using experimental data and numerical simulations.

The antioxidant capacity of MRPs, as assessed using the DPPH assay, exhibited a trend characterized by rapid reaction kinetics and high antioxidant values, underscoring their potential application in food systems where swift neutralization of reactive oxygen species is desirable. This behavior was incorporated into the kinetic model. HPLC-CAD analysis confirmed that the production of DDMP, its furanone tautomer and MPO were contributors to antioxidant activity.

The rapid reactivity of MRPs measured with DPPH assay kinetic approach suggests that MRPs could serve as effective natural antioxidants. These results underscore the importance of specific MRPs,

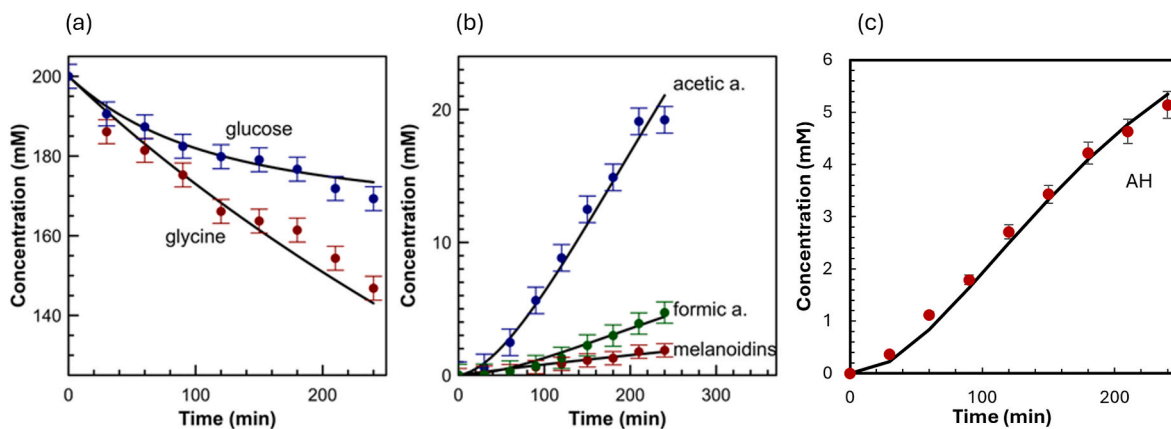


Fig. 5. Fits to experimental data (colored dots) with the updated kinetic model based on Martins & Van Boekel, 2005 (black lines) of (a) reactants (glucose, glycine) and (b) products (formic acid, acetic acid and melanoidins) using Copasi software. (c) Trend of antioxidant species formation during Maillard reaction measured (red dots) and fitted by the kinetic model (black line).

particularly DDMP and its isomer, in significantly boosting the overall antioxidant capacity of the system.

In conclusion, this study enhances our understanding of the formation, reactivity, and antioxidant potential of MRPs. These findings demonstrate that MRPs can inhibit oxidative deterioration, extend the shelf life, and improve overall food quality. While this study focused on specific temperature and pH conditions, it is reasonable to speculate that optimizing these reaction parameters could further enhance the formation of desirable antioxidants while minimizing the production of less favorable byproducts. These insights have important implications for the development of naturally derived food additives, particularly in the context of "clean label" products, where there is growing demand for natural alternatives to synthetic additives. Future research should focus on further elucidating the mechanisms governing MRP formation and degradation, and exploring their potential applications in food science and nutrition under varying processing conditions.

CRediT authorship contribution statement

Sara Bolchini: Writing – original draft, Visualization, Investigation, Formal analysis. **Lucrezia Angeli:** Methodology. **Giovanna Ferrentino:** Writing – review & editing. **M.A.J.S. Van Boekel:** Writing – review & editing. **Riccardo Amorati:** Writing – review & editing. **Matteo Scampicchio:** Writing – review & editing, Supervision, Software, Formal analysis, Conceptualization. **Ksenia Morozova:** Writing – review & editing, Supervision, Methodology, Investigation.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.lwt.2024.117316>.

[org/10.1016/j.lwt.2024.117316](https://doi.org/10.1016/j.lwt.2024.117316).

Data availability

Data are available upon request from the authors.

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