1. Introduction

Histamine, a common biogenic amine produced by bacterial decarboxylation from the amino acid histidine, acts to transmit signals from cell to cell in the skin, gut, and other organs of the immune system.\(^1\)\(^-\)\(^2\) It has been confirmed that the ingestion of excess histamine can trigger a wide array of symptoms including nausea, headache, diarrhea, oral numbness, palpitations, rashes, asthma and even death\(^3\)\(^-\)\(^5\) compromising health safety. In this context, histamine as the primary toxic compound existing in wine has attracted significant attention in food quality monitoring. Since conventional processing methods such as cooking and freezing cannot destroy histamine once it has formed,\(^6\) the continuous monitoring of trace levels of histamine in food products is essential.

Many methods have been developed for the detection of histamine, such as gas chromatography (GC),\(^7\) electrochemical sensing,\(^8\) capillary zone electrophoresis,\(^9\) thin layer chromatography (TLC),\(^10\) high-performance liquid chromatography (HPLC)\(^11\) and enzyme-linked immunosorbent assay (ELISA).\(^12\) Although these conventional methods have the merits of high sensitivity and accuracy, they have several inherent drawbacks. Chromatographic methods are relatively sophisticated, expensive and tedious. ELISA is strongly dependent on the use of unstable enzymes and expensive test kits, which cannot meet the high-throughput and quality control rigor required by the government and food industry. Thus, the exploitation of alternative rapid and accurate approach is of vital importance.

Spectroscopic methods are promising tools for the determination of toxic or harmful substances in food because they are inherently rapid, specific, and partially or completely computerized.\(^13\)\(^-\)\(^16\) The current fluorescence methods used in detection showed promise in the development of bioassay techniques, owing to its fast, sensitive and reproducible signal. Upconversion nanoparticles (UCNPs) doped with lanthanide have attracted tremendous attention.\(^17\)\(^-\)\(^19\) UCNPs can emit a high-energy photon at a shorter wavelength under excitation by multiple low-energy near-infrared (NIR) photons, instead of traditional photoexcitation in the UV or visible regions. As autofluorescence originating from coexisting substances in sample matrices can be greatly minimized by the use of NIR photons, the background light interference and sensitivity of the detection can be improved with UCNPs-labeled optical sensing system with NIR as the excitation source.\(^20\) In addition, UCNPs has good photochemical stability, large Stokes shifts, is nonblinking with deep penetration effects in biological samples and relatively low toxicity. All of these appealing advantages makes UCNPs a promising label compared to traditional fluorescent materials.\(^21\),\(^22\) Another promising spectroscopic technology is surface-enhanced Raman spectroscopy (SERS). SERS has become one of the most versatile quantitative approaches for detection even down to the single-molecule level.\(^23\),\(^24\) It has
been successfully used as a rapid and accurate detection method for determining many kinds of chemical compounds or bacteria in food during the past decade.\textsuperscript{25-27}

Despite the tremendous progress made in the field of fluorescence and SERS detection technologies, challenges exist. In particular, the specific recognition of target molecule (i.e., selectivity) for separation-detection is difficult to achieve using traditional host guest chemistry in the sensor design. This has become a major challenge to the analytical utility. Here, sample pretreatment is essential to eliminate interferences to improve separation and detection and obtain satisfactory results. Molecular imprinting offers a new technique for efficient separation and enrichment of specific analytes from complicated matrices. It is a kind of biomimetic technique to synthesize molecularly-imprinted polymers (MIPs) with artificially generated recognition sites that are able to specifically rebind a target molecule in preference to other closely-related compounds.\textsuperscript{28} MIPs synthesis is relatively cheap and easy, making it an excellent alternative to natural receptors. Because of these advantages, MIPs has attracted considerable attention and has been integrated with HPLC, quantum dots and graphene oxide for the determination of food chemical hazards in recent years.\textsuperscript{29-30} Feng et al.\textsuperscript{11} and Hu et al.\textsuperscript{12} had successfully combined MIPs and SERS for accurate and high-throughput detection of melamine and \textit{z}-tocopherol in milk and vegetable oils, respectively. To the best of our knowledge, very few attempts have integrated MIPs and SERS or UCNPs in a single sensor system for separation and detection of histamine in food samples.

The current study aims to develop a novel sensor material for dual mode ultrasensitive detection of histamine by integrating the selective recognition ability of MIPs and the high detection sensitivity and accuracy of UCNPs and SERS technologies.

2. Materials and methods

2.1. Chemicals and reagents

Histamine, putrescine, \( \beta \)-phenylethylamine, spermine, spermidine, tryptamine, tyramine, cadaverine, ethylene glycol dimethacrylate (EGDMA), ammonium fluoride, \( z \)-methacrylic acid (MAA), 2,2’-azobis(isobutyronitrile) (AIBN) and poly(acrylic acid) (PAA) (M.W. 2000) were obtained from Sigma-Aldrich (Saint Louis, MO, USA). Chloroform, ammonium acetate and sodium borohydride were from TCI Co., Ltd. (Portland, OR, USA). Ammonium fluoride (NH\( _4 \)F), ethanol, sodium hydroxide, methanol and sodium chloride were purchased from Fisher Scientific (Waltham, MA, USA). Glutaraldehyde, (3-amino-propyl) triethoxysilane (APTES), 1,6-hexanediamine and octadecene (ODE), tetraethyl orthosilicate (TEOS), oleic acid (OA), ytterbium(III) chloride hexahydrate (YbCl\( _3 \)), yttrium(III) chloride hexahydrate (YCl\( _3 \)) and erbium chloride hexahydrate (ErCl\( _3 \)) were all obtained from Alfa Aesar Co., Inc. (Tewksbury, MA, USA). Rice wine and red wine were purchased from Better World Market (West Lafayette, IN, USA). Canned tuna was purchased from Meijer Market (West Lafayette, IN, USA). All reagents and solvents used were analytical or HPLC grade. Deionized (DI) water (18.2 \( \text{M} \Omega \text{ cm}^{-1} \)) was prepared by the Millipore system (Billerica, MA, USA).

2.2. Reference method for histamine determination

Histamine content was determined by the method of Gao et al.\textsuperscript{4} Typically, a HPLC system (Agilent 1100 series, Germany) coupled with a quaternary pump and a diode array detector was used to determine the concentration of histamine, with the use of a HILIC column (Luna@HILIC, 3 mm, 100 \times 2 mm, Phenomenex, Torrance, CA, USA). The flow rate was 0.4 mL min\(^{-1} \) and mobile phase was consisted of ammonium acetate (5 mM) and acetonitrile (1 : 9, v/v).

2.3. Synthesis of dual signal response material

2.3.1. Preparation of UCNPs. In this work, oleic acid-capped UCNPs (OA-UCNPs) were synthesized using a previously reported solvothermal method with slight modifications.\textsuperscript{31} Briefly, 0.2366 g of YCl\( _3 \cdot 6\text{H}_2\text{O} \) (0.78 mmol), 0.0774 g of YbCl\( _3 \cdot 6\text{H}_2\text{O} \) (0.20 mmol) and 0.0076 g of ErCl\( _3 \cdot 6\text{H}_2\text{O} \) (0.02 mmol) were dissolved in 5 mL OA and 15 mL ODE in a 100 mL three-necked flask. The mixture was heated to 160 °C for 30 min to form a homogeneous yellow solution, and cooled down naturally to room temperature. Then 10 mL of methanol solution containing 0.1000 g NaOH and 0.1480 g NH\( _4 \)F was slowly added to the flask, followed by stirring at 30 °C for 1 h to consume all fluoride. Subsequently, the solution was heated to 100 °C and maintained for 15 min, ensuring that the methanol was removed completely. Then the solution was heated up to 300 °C and kept for 1 h under an argon atmosphere. After cooling to room temperature, the nanocrystals were obtained by centrifugation at 5000 rpm. The precipitate from the solution was washed with ethanol/cyclohexane mixture (1 : 1 v/v) three times. Finally, the obtained UCNPs were dried overnight in a vacuum oven at 60 °C. OA-UCNPs were thus formed and kept in a desiccator until ready to use.

The surface modification of the OA-UCNPs was performed using a ligand exchange with PAA.\textsuperscript{34} Typically, 30 mg of the OA-capped UCNPs in chloroform (4 mL) were slowly added to a water solution (8 mL) containing 200 mg PAA, and the solution was vigorously stirred for 24 h. After that, the resultant solution was transferred into the upper water layer from the chloroform layer due to the PAA attachment. The nanoparticles were separated by centrifugation and washed three times with ultrapure water to obtain PAA-stabilized UCNPs.

2.3.2. Synthesis of UCNPs@MIPs-AgNPs. UCNPs@MIPs-AgNPs was prepared as follows: 1 mmol of histamine (template), 0.25 g of UCNPs and 4 mmol of MAA (functional monomer) were dissolved in a mixture of 2 mL tetrahydrofuran and 3 mL methanol in a 50 mL flask. After the solution was mechanically stirred for 10 min before the addition of 4 mmol of EGDMA (cross-linker) and incubated in oil bath overnight at 50 °C, 500 mg AgNO\( _3 \) (AgNPs precursor) and 30 mg of AIBN (free-radical initiator) were added to the flask, followed by purging with a gentle flow of nitrogen for 15 min. Subsequently, the flask was sealed and placed in an oil bath at 70 °C for 24 h. The product was ground to powder and sieved through a 200 mesh steel sieve to obtain homogeneous fine particles. Then, excessive NaBH\( _4 \) was added to reduce UCNPs@MIPs-AgNO\( _3 \) into UCNPs@MIPs-AgNPs. Afterward, histamine was removed...
by Soxhlet extraction with methanol/acetic acid (8:2, v/v) solution until no histamine could be detected using HPLC. UCNPs@MIPs–AgNPs was then vacuum-dried at 60 °C and ground to powder with a mortar and pestle. As a reference, the non-molecular imprinted polymer (UCNPs@NIPs–AgNPs) was prepared using the same protocol without the addition of the template molecule (histamine).

2.4. Evaluation of the binding capacities of UCNPs@MIPs–AgNPs and UCNPs@NIPs–AgNPs

For the static adsorption tests, histamine was dissolved in 2 mL of ammonium acetate solution (30 mM) to form standard solutions with different concentrations (10, 25, 50, 75, 100, 150 and 200 mg L⁻¹), which were mixed with 10 mg of UCNPs@MIPs–AgNPs or UCNPs@NIPs–AgNPs. Then, the mixtures were shaken in a thermostatic water bath oscillator at 200 rpm for 45 min (25 °C) and centrifuged at 5000 × g for 15 min. The resulting supernatant was filtered through a membrane filter (pore size: 0.45 μm) and the concentrations of free histamine (i.e., unbounded histamine) were determined by the HPLC method.

The adsorption capacity (Q) was calculated by the following formula:

\[ Q = \left( C_i - C_e \right) \times \frac{V}{W} \tag{1} \]

where \( C_i \) and \( C_e \) represent the initial and final concentration of histamine, respectively; \( V \) is the volume of solution and \( W \) represents the mass of polymer.

In addition, the Scatchard model, Freundlich model and Langmuir isotherm model were used to determine the affinity of UCNPs@MIPs–AgNPs and UCNPs@NIPs–AgNPs in the study. These three models were developed according to the equations as follows:

Scatchard model:

\[ \frac{Q}{C_e} = \frac{Q_{\text{max}}}{K_d} - \frac{Q}{K_d} \tag{2} \]

Freundlich model:

\[ \log Q = m \log C_e + \log a \tag{3} \]

Langmuir isotherm model:

\[ \frac{C_e}{Q} = \frac{C_e}{Q_{\text{max}}} + \frac{1}{Q_{\text{max}} b} \tag{4} \]

where \( C_e \) (mg L⁻¹) is the free histamine concentration (mg L⁻¹) at equilibrium, \( Q_{\text{max}} \) is the saturated adsorption capacity (mg g⁻¹), \( K_d \) is the dissociation constant (mg L⁻¹), \( m \) is the adsorption intensity of surface heterogeneity, \( a \) is the adsorption capacity of histamine (mg g⁻¹) and \( b \) is the Langmuir adsorption equilibrium constant (L mg⁻¹).

For the study of kinetic adsorption of UCNPs@MIPs–AgNPs (or UCNPs@NIPs–AgNPs) towards histamine, a set of seven samples was incubated at 25 °C for different time points (5, 15, 30, 45, 60, 90 and 150 min). The mixture was treated and determined following the same procedures above mentioned for static adsorption.

The pseudo-first-order kinetic model and pseudo-second-order kinetic model were used to describe the dynamic adsorption process according to eqn (5) and (6), respectively:

\[ \ln \left( Q - Q_t \right) = \ln Q - k_1 t \tag{5} \]

\[ \frac{t}{Q_t} = \frac{1}{K_2 Q_{\text{max}}^2} + \frac{t}{Q_{\text{max}}} \tag{6} \]

where \( Q_c \) represents the adsorption capacity (mg g⁻¹) at a particular time point; \( k_1 \) means the rate constant of first-order sorption (ln[mg g⁻¹ s⁻¹]); \( k_2 \) is the rate constant of second-order sorption (mg g⁻¹ s⁻¹).

To evaluate the nature of diffusion process for histamine adsorption, the intra-particle diffusion model was introduced. The intra-particle mass transfer diffusion model proposed by Weber and Morris can be expressed as follows:

\[ Q_t = k_3 t^{1/2} + C \tag{7} \]

where \( C \) is the intercept (mg g⁻¹) and \( k_3 \) is the intra-particle diffusion rate constant (mg g⁻¹ s⁻¹), which can be obtained from the slope of the linear plots of \( Q_t \) vs. \( t^{1/2} \).

The selectivity test was performed by comparing the adsorption capacity of UCNPs@MIPs–AgNPs (or UCNPs@NIPs–AgNPs) towards histamine and that of its analogues at 75 mg L⁻¹ after 45 min incubation. Herein, histamine, β-phenylethylamine, putrescine, spermine, spermidine, tryptamine, tyramine and cadaverine were selected as competitive agents to determine histamine selectivity of the UCNPs@MIPs–AgNPs.

The distribution coefficient (\( K_d \)), selectivity coefficient (\( k \)) and the relative selectivity coefficient (\( \alpha \)) of the sorbent obtained in these comparative experiments were also used to evaluate the competitive selective capacity of UCNPs@MIPs–AgNPs and UCNPs@NIPs–AgNPs. These three parameters were calculated following the equations below:

\[ K_d = \frac{Q}{C_e} \tag{8} \]

\[ k = \frac{K_d(\text{histamine})}{K_d(\text{analogues})} \tag{9} \]

\[ \alpha = \frac{k_{\text{MIP}}}{k_{\text{NIP}}} \tag{10} \]

\( K_d \) indicates the affinity of the sorbent for a particular substance. Generally, the larger the value of \( K_d \) is, the stronger the adsorption capability of a substance would be. The value of \( k \) suggests the selectivity between the target analyte and its structural similar compounds. The value of \( \alpha \) reveals how selective a sorbent is for a particular substance compared with the selectivity of a different sorbent.

2.5. Fluorescence and SERS measurements

Sample solutions containing various concentrations (0.00 mg L⁻¹, 0.02 mg L⁻¹, 0.1 mg L⁻¹, 1 mg L⁻¹, 5 mg L⁻¹, 10 mg L⁻¹,
50 mg L\(^{-1}\), 100 mg L\(^{-1}\) and 250 mg L\(^{-1}\)) of histamine standard were mixed with 10 mg of UCNPs@MIPs–AgNPs (UCNPs@NIPs–AgNPs). After shaking at room temperature for 45 min, the fluorescence signal of the solution was scanned by using a Hitachi F-7000 fluorescence spectrometer equipped with an external 980 nm laser. For the blank (0.00 mg L\(^{-1}\)), twenty measurements were performed. For other levels (0.02 mg L\(^{-1}\), 0.1 mg L\(^{-1}\), 1 mg L\(^{-1}\), 5 mg L\(^{-1}\), 10 mg L\(^{-1}\), 50 mg L\(^{-1}\), 100 mg L\(^{-1}\) and 250 mg L\(^{-1}\)), five measurements were performed.

For SERS detection, sample solutions containing various concentrations (0.00 mg L\(^{-1}\), 0.1 mg L\(^{-1}\), 1 mg L\(^{-1}\), 5 mg L\(^{-1}\), 10 mg L\(^{-1}\), 50 mg L\(^{-1}\) and 100 mg L\(^{-1}\)) of histamine standard were mixed with 10 mg of UCNPs@MIPs–AgNPs (UCNPs@NIPs–AgNPs). After shaking at room temperature for 45 min, the mixture then centrifuged at 5000 \(\times g\) for 5 min. The supernatant was discarded and the pellet was dried before depositing onto the gold-coated microarray chip for SERS spectra collection. The SERS spectra were obtained by averaging the spectra detected at seven random different detection spots.

2.6. Characterization

The surface groups of the synthesized UCNPs and UCNPs@MIPs–AgNPs were measured using a FT-IR spectrophotometer (Nexus 670, Thermo Electron Corp., Madison, WI, USA) in the range of 650–4000 cm\(^{-1}\) with 256 scans at a resolution of 4 cm\(^{-1}\). The sample powders (5 mg) were ground with spectroscopic grade KBr powders (250 mg) and then pressed into 1 mm pellets.\(^{35}\) Morphologies of the synthesized nanoparticles were recorded by a high-resolution FEI Tecnai G2 20 transmission electron microscope equipped with a charge-coupled device (CCD) camera operating at an acceleration voltage of 200 kV. UV–vis absorption spectra were collected with a JASCO V570 UV/visible/NIR spectrophotometer (JASCO, Inc., Easton, MD, USA). A Bruker Senterra confocal Raman system (Billerica, MA, USA) coupled with 50× air objective (NA0.9, infinity and flat field corrected), a motorized microscope and a CCD array detector was employed for SERS spectra acquisition. The excited laser source was provided by a 785 nm air-cooled He–Ne laser at 10 mW and the exposure time for each SERS measurement was set at 10 s. The spot diameter of laser source with our condition is about 12 \(\mu\)m. Upconversion fluorescence spectra were recorded on a Hitachi F-7000 fluorescence spectrophotometer (Hitachi Co., Tokyo, Japan) modified with an external 0–1300 mW adjustable continuous wave 980 nm laser (Beijing Hi-Tech Optoelectronic Co., Beijing, China) instead of the internal excitation source.

2.7. Statistical analysis

All data were expressed as mean ± standard deviation (SD) of at least three determinations throughout the study, and spectra drawings were performed using Origin 8.5 (OriginLab Inc., Northampton, MA, USA). Data was analyzed using one-way analysis of variance and means were compared by Duncan’s multiple range tests using SPSS 18.0 (SPSS Inc., Chicago, IL, USA). The probability value of \(p < 0.05\) was considered significant. The OMNIC software (v. 8.0, Thermo Electron Corp., Madison, WI, USA), OPUS 6.5 software (Opus Software, Inc., San Francisco, CA, USA) and FL Solutions 2.1 software (Hitachi High-technology Co., Tokyo, Japan) were utilized for FT-IR, Raman and fluorescence spectra acquisition and instrument control, respectively. Principal component analysis (PCA) was implemented in the commercial chemometric software The Unscrambler (v 10.2; CAMO Software AS, Oslo, Norway).

3. Results and discussions

3.1. Analytical principle

The typical dual-mode system for the detection of histamine based on fluorescence and SERS is illustrated in Scheme 1. The UCNPs@MIPs–AgNPs synthesized combined MIPs with fluorescence and SERS detection modalities. When histamine molecules were bound to the material, strong fluorescence quenching and prominent electromagnetic enhancement effect could occur simultaneously. The fluorescence quenching mechanism might not be through fluorescence resonance energy transfer since no overlap existed between the histamine absorption peak (at \(\sim 210\) nm) and the UCNPs emission peak (at 541 nm). The quenching mechanism could be attributed to the photo-induced electron transfer from histamine to UCNPs. The amine groups of histamine interacted with carboxyl groups in UCNPs@MIPs–AgNPs. The lone pair of electrons in oxygen-containing groups was available for photo-induced electron transfer, leading to a decrease in emission. When analytes were absorbed on the shell of MIPs, they (histamine molecules) also located closely to SERS-active substrates (AgNPs), which were incorporated in the outer shell of MIPs during polymerization, resulting in improved reproducibility of the SERS spectra. The methodology was proposed by Liu et al.\(^{36}\) They integrated AgNPs with MIPs by adding silver nitrate as AgNPs precursor that bind to theophylline during bulk polymerization. Hu et al.\(^{37}\) successfully applied this methodology for the detection of melamine by integrating AgNPs with MIPs for melamine. Considering the fact that similar to the structure of theophylline, histamine also contains an imidazole ring, which could interact with AgNPs precursor during polymerization, this methodology was adopted in this study.

3.2. Characterization of UCNPs@MIPs–AgNPs and UCNPs@NIPs–AgNPs

3.2.1. TEM. The morphologies of UCNPs and UCNPs@MIPs–AgNPs were characterized by TEM. As shown in Fig. 1a, the synthesized UCNPs were spherical in shape with fairly uniform size of approximately 100 nm, which was in accordance with the result reported by other researchers.\(^{38}\) Fig. 1c showed the TEM image of UCNPs@MIPs–AgNPs, and it could be clearly observed that there was a shell dotted with small nanoparticles (AgNPs) uniformly coated on the surface of the UCNPs, indicating that core–shell UCNPs (UCNPs@MIPs–AgNPs) were obtained.

3.2.2. FT-IR spectra. The FT-IR spectra of the synthesized UCNPs and UCNPs@MIPs–AgNPs were shown in Fig. 1b. For
the FT-IR spectrum of UCNPs, the dominating absorption peaks observed at around 2951 cm$^{-1}$ and 2853 cm$^{-1}$, corresponding to the bending vibrations of methylene (−CH$_2$−) and methyl groups (CH$_3$−) respectively, and 1725 cm$^{-1}$, which could be attributed to the stretching vibrations of the carbonyl group (C=O). After coating MIPs on the UCNPs, due to the existence of cross-linkers and monomers used to synthesize MIPs, the peak intensities at 2951 cm$^{-1}$, 2853 cm$^{-1}$ and 1725 cm$^{-1}$ were obviously enhanced. At the same time, a new peak at around 1170 cm$^{-1}$ were observed in the spectrum of UCNPs@MIPs−AgNPs. This band was assigned to the stretching vibration adsorption of ether group (C–O–C). In addition, the peaks in the absorption band between 3200 cm$^{-1}$ and 3800 cm$^{-1}$ shifted from 3458 cm$^{-1}$ in the spectrum of UCNPs to 3441 cm$^{-1}$ in the spectrum for UCNPs@MIPs−AgNPs, which might be due to the interactions between the −NH$_2$ group of the template molecule and the C=O group of MAA. The spectra for UCNPs and UCNPs@MIPs−AgNPs had similar profiles (similar locations and appearances of the major bonds) indicated that they had same chemical nature. These results demonstrated that the polymerization on the surface of UCNPs particles had been successfully implemented.

3.2.3. Adsorption properties of UCNPs@MIPs−AgNPs and UCNPs@NIPs−AgNPs

3.2.3.1. Adsorption kinetics. To evaluate the selective recognition, binding capacity and equilibrium rate of the synthesized UCNPs@MIPs−AgNPs, static and kinetic adsorption tests were performed for both of UCNPs@MIPs−AgNPs and UCNPs@NIPs−AgNPs. The kinetic adsorption curves were shown in Fig. 2a. Here, both the absorption capacities of UCNPs@MIPs−AgNPs and UCNPs@NIPs−AgNPs increased with incubation time, reaching an adsorption equilibrium at around 45 min. Thus, 45 min was chosen as optimal extraction time. Note that at all conditions, the amount of histamine binding to UCNPs@MIPs−AgNPs was remarkably higher than that of UCNPs@NIPs−AgNPs, verifying the superior performance of UCNPs@MIPs−AgNPs versus UCNPs@NIPs−AgNPs with respect to histamine binding. The binding capacity of UCNPs@NIPs−AgNPs was contributed by nonspecific binding, which was hypothesized to be driven by the interruption of the hydrogen-bond interactions by the solvent media. UCNPs@MIPs−AgNPs contained of nonspecific binding sites as well as specific reorganization sites, which was ascribed to the host–guest shape and size recognition and the bond reaction between them. However, the number of non-specific binding sites was limited. That explained why much higher $Q$ value for UCNPs@MIPs−AgNPs was observed compared with that of UCNPs@NIPs−AgNPs.

Three different models, namely pseudo-first-order kinetic model, pseudo-second-order kinetic model and intra-particle diffusion model, were chosen to investigate dynamic adsorption mechanism of histamine. As seen in Fig. 2b and 3, the adsorption reaction of UCNPs@MIPs−AgNPs towards histamine could be well fitted by both of pseudo-first-order kinetic model and pseudo-second-order kinetic model while the fitting performance of intra-particle diffusion model was far worse. In addition, compared with pseudo-first-order kinetic model and intra-particle diffusion model, the pseudo-second-order kinetic model described the adsorption behavior of UCNPs@MIPs−AgNPs on histamine with higher $R^2$ (0.9965). Moreover, the theoretical $Q_e$ value calculated from the slope of the pseudo-second-order kinetic model ($t$/$Q_e$ versus $t$) was 1.85 mg g$^{-1}$, closed to the $Q_e$ value (1.80 mg g$^{-1}$) obtained from the experiment. These results indicated that instead of intra-particle diffusion, the adsorption process of UCNPs@MIPs−AgNPs towards histamine was controlled by chemical adsorption mechanism by sharing electrons between them.

3.2.3.2. Static binding characteristics. The static adsorption capacities of UCNPs@MIPs−AgNPs and UCNPs@NIPs−AgNPs for histamine were determined in the concentration range from...
10 mg L\(^{-1}\) to 200 mg L\(^{-1}\), and the relationship of their total adsorption capacities to the concentration of histamine was presented in Fig. 4a.

It can be seen from the static adsorption isotherm curves that the adsorption capacities for both UCNPs@MIPs–AgNPs and UCNPs@NIPs–AgNPs increased along with the increase in the initial concentration of histamine. In addition, the adsorption capacities of UCNPs@MIPs–AgNPs were constantly higher than those of UCNPs@NIPs–AgNPs, which was similar to the results of kinetic adsorption experiments. Specifically, when the concentration was 200 mg L\(^{-1}\), the adsorption capacity of MIPs (3.11 mg g\(^{-1}\)) was about 4.5 times that of NIPs (0.70 mg g\(^{-1}\)), illustrating that the chemical and physical properties of histamine had been successfully imprinted into UCNPs@MIPs–AgNPs. As a result, UCNPs@MIPs–AgNPs have higher specific recognition ability toward histamine than UCNPs@NIPs–AgNPs and were suitable for further application.

Scatchard relationship was determined to assess the affinity of UCNPs@MIPs–AgNPs toward histamine. As shown in Fig. 4b, two intersecting straight lines with different slopes corresponding to the high- and low-affinity populations of binding sites were observed in the Scatchard plot. By contrast, UCNPs@NIPs–AgNPs showed nonlinearity, indicating that there was no selective adsorption sites for histamine. To further estimate the binding properties of UCNPs@MIPs–AgNPs, the isotherm absorption data were also processed with the Freundlich and Langmuir adsorption isotherm models and shown in Fig. 5a and b. The results demonstrated that the binding process of histamine with the UCNPs@MIPs–AgNPs could be well fitted by the two isotherm models (\(R^2 = 0.9994\) and 0.9754 for Freundlich model and Langmuir model, respectively), indicating that UCNPs@MIPs–AgNPs possessed a heterogeneous binding site distribution.

3.2.3.3. Adsorption selectivity. To evaluate the specificity of the UCNPs@MIPs–AgNPs, several analogues of structural features of histamine were selected, including \(\beta\)-phenylethylamine, putrescine, spermidine, spermine, tryptamine, tyramine.
and cadaverine. The competitive adsorption experiments of these analogues with histamine on the sorbents were carried out with 75 mg L$^{-1}$ of standard solution. The molecular structures of the analytes and binding amounts of these targets on UCNPs@MIPs–AgNPs and UCNPs@NIPs–AgNPs were shown in Fig. 6. As seen from the figure, histamine owned remarkably higher binding capacity than its analogues, suggesting satisfactory selectivity of the synthesized UCNPs@MIPs–AgNPs for histamine. Table 1 and Table 2 summarized the values of $K_d$, $k$ and $a$, which were calculated to evaluate the competitive selective capabilities of UCNPs@MIPs–AgNPs and UCNPs@NIPs–AgNPs. As shown in Table 1, the $K_d$ of histamine was
clearly larger than that of the other seven structure analogues of histamine, validating the strong adsorption capacity of UCNPs@NIPs–AgNPs for histamine. In addition, the $k$ value of seven structural analogues for UCNPs@MIPs–AgNPs was several times higher than the corresponding values for UCNPs@NIPs–AgNPs (Table 2), suggesting that the UCNP-s@MIPs offering more specific recognition sites for the target molecule (i.e. histamine) than non-target substances. The relatively high $a$ values (>1) of $\beta$-phenylethylamine, putrescine, spermidine, spermine, tryptamine, tyramine and cadaverine further confirmed that UCNPs@MIPs–AgNPs for histamine was successfully synthesized and the rebinding occurred specifically and selectively for histamine rather than its structural analogues.

3.3. Analytical performance

3.3.1. Detection of histamine using fluorescence mode.

Fig. 7 showed various intensities of the fluorescence spectra of UCNPs@MIPs–AgNPs in the presence of different concentrations of histamine. In the fluorescence spectra, three dominant emission peaks were observed at 488, 541 and 665 nm, which were mainly generated by the transitions from the $2H_{9/2}$, $4S_{3/2}$, and $4F_{9/2}$ levels to the $4I_{15/2}$ ground state of the $\text{Er}^{3+}$ ion.17

As seen in Fig. 7a, within the histamine concentration range of 0.02–250 mg L$^{-1}$, the fluorescence intensities of UCNPs@MIPs–AgNPs gradually decreased as the concentration of the target (histamine) increased. Principal component analysis (PCA) was performed to further explain the fluorescence spectra. As shown in Fig. 8a, the explained variation rate for the first two principle components (PCs) was 64% (PC1) and 18% (PC2), respectively, which indicated that the first two PCs could fairly well represent the essential features of the fluorescence spectra. In addition, there was a clear separation tendency between samples containing different levels of histamine, suggesting samples containing different levels of histamine can be well discriminated using
There was a good linear relationship (correlation coefficient of 0.9956 for a linear regression fit, \( y = 1368.81 - 131.61 \lg x \) (Fig. 7b)) between the fluorescence intensities and the concentrations of histamine in samples. Moreover, the detection limit (LOD) and quantitation limit (LOQ) of the sensing system were calculated to be 0.009 mg L\(^{-1}\) and 0.02 mg L\(^{-1}\), respectively.

### 3.3.2. Detection of histamine using SERS mode

Fig. 9 displayed spectral features of histamine determined by the normal Raman and SERS spectra while Fig. 10 demonstrated the linear correlation between the SERS intensities and the concentrations of histamine. As seen in Fig. 9, some band shifts existed, which might be due to the SERS effect. When target molecules adsorbed onto the surface of substrate, some molecules interacted with the noble-metallic nanostructures, resulting in the changes in dipole of the molecules and subsequent shifts in the location of SERS bands. Four major peaks at around 1268 cm\(^{-1}\), 1331 cm\(^{-1}\), 1427 cm\(^{-1}\) and 1572 cm\(^{-1}\) were observed in similar positions in both of normal Raman spectra of histamine powder and SERS spectra of histamine solution. These four strong peaks were also reported by Gao et al.\(^4\) and Jančí et al.\(^40\) The Raman peaks at 1268 cm\(^{-1}\), 1331 cm\(^{-1}\) and 1572 cm\(^{-1}\) might be related to the imidazole ring stretching and breathing while the peak at 1427 cm\(^{-1}\) was assigned to vibrations of carboxylic group. As shown in Fig. 10a, the changes of these four strong Raman peaks depend on histamine concentrations. The SERS intensities of UCNP-s@MIPs–AgNPs increased as the concentration of histamine increased. A preliminary analysis of data was also performed by PCA to examine any possible grouping of samples according to...
the histamine contents in samples. The first two PCs explained 70% and 15% of the total variation, respectively, indicating that the cumulative reliabilities could explain most of the Raman spectra information so that they could be used to represent the variables for classification of samples. Fig. 8b showed the score plots of the first and the second principle components, with samples shaped according to different histamine levels. A clear separation was also observed, suggesting samples containing different levels of histamine could also be well discriminated with Raman spectroscopy. Within the histamine concentration range of 0.1–100 mg L\(^{-1}\), highest intensity was observed at 1427 cm\(^{-1}\), followed by 1572 cm\(^{-1}\), 1268 cm\(^{-1}\) and 1331 cm\(^{-1}\). The intensities of these four major peaks were selected for quantitative evaluation. Here, six critical levels (0.1 mg L\(^{-1}\), 1 mg L\(^{-1}\), 5 mg L\(^{-1}\), 10 mg L\(^{-1}\), 50 mg L\(^{-1}\) and 100 mg L\(^{-1}\)) were used to establish calibration curves. It appeared that the SERS intensities at 1331 cm\(^{-1}\) showed better dose-dependence effect than the other three peaks. From this observation, peak intensities at 1331 cm\(^{-1}\) was adopted and used for further assessment. The correlation between the SERS Raman intensities and the concentrations of histamine based on peak intensities at 1331 cm\(^{-1}\) was shown in Fig. 10b. A good linear relationship between the SERS Raman intensities and the concentrations of histamine was obtained. The linear regression equation fit obtained was of \(y = 28.88x + 147.03\) with the correlation coefficient of 0.9845. The LOD and LOQ value of the sensor system under SERS mode were 0.04 mg L\(^{-1}\) and 0.1 mg L\(^{-1}\) respectively.

For comparison, the analytical performances of some currently used detection methods for histamine reported in the literature are provided in Table 3. As seen from the results, the LOD obtained from the proposed method was significantly lower.

### 3.4. Analytical application

The accuracy of the proposed dual mode bioassay for determining histamine levels in real food samples was evaluated by detecting histamine in red wine, rice wine and canned tuna samples spiked with a series of known quantities of histamine (5, 15 and 30 mg L\(^{-1}\)). The results of recoveries were summarized in Table 4. All of the three spiking levels in red wine, rice wine and canned tuna samples showed good recoveries. It was worth noting that the recovery results of canned tuna ranged from 85.06% to 117.36%, which were not as good as the results from wine samples (93.30–108.30%). This could be due to the fact that the matrix in canned tuna was more complex than that of red wine and rice wine samples. In addition, food samples spiked with histamine at three levels were analyzed by HPLC-UV

![Fig. 10](image)

(a) Typical SERS response of UCNPs@MIPs–AgNPs in various histamine concentrations; (b) linear correlation between the SERS Raman intensities and the concentrations of histamine.

### Table 3  Comparison of the results for histamine detection obtained from our method and that reported by others

<table>
<thead>
<tr>
<th>Methods</th>
<th>Detection range</th>
<th>LOD(^a)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gas chromatography(^7)</td>
<td>100–1000 mg L(^{-1})</td>
<td>7 mg L(^{-1})</td>
</tr>
<tr>
<td>Capillary zone electrophoresis(^9)</td>
<td>0.0011–0.5550 mg L(^{-1})</td>
<td>0.0006 mg L(^{-1})</td>
</tr>
<tr>
<td>High-performance liquid chromatography(^11)</td>
<td>2.22–111 mg L(^{-1})</td>
<td>0.747 mg L(^{-1})</td>
</tr>
<tr>
<td>Enzyme-linked immunosorbent assay(^41)</td>
<td>0.0127–1.8531 mg L(^{-1})</td>
<td>0.0936 mg mL(^{-1})</td>
</tr>
<tr>
<td>SPR(^b)-based indirect competitive immunosensor</td>
<td>0.003–0.01 mg L(^{-1})</td>
<td>0.003 mg L(^{-1})</td>
</tr>
<tr>
<td>SPR-based molecularly imprinting(^42)</td>
<td>0.025–1 mg L(^{-1})</td>
<td>0.025 mg L(^{-1})</td>
</tr>
<tr>
<td>TLC(^c)-based surface enhanced Raman spectroscopy(^43)</td>
<td>15–100 mg kg(^{-1})</td>
<td>—</td>
</tr>
<tr>
<td>MIPs(^d)-based surface enhanced Raman spectroscopy(^4)</td>
<td>3–90 mg L(^{-1})</td>
<td>—</td>
</tr>
<tr>
<td>Electrochemical sensing(^8)</td>
<td>40–170 mg L(^{-1})</td>
<td>38 mg L(^{-1})</td>
</tr>
<tr>
<td>The developed method</td>
<td>0.1–100 mg L(^{-1})/0.02–250 mg L(^{-1})</td>
<td>0.04/0.009 mg L(^{-1})</td>
</tr>
</tbody>
</table>

\(^a\) Limit of detection. \(^b\) Surface plasmon resonance. \(^c\) Thin-layer chromatography. \(^d\) Molecularly imprinted polymers.
to verify the practicability of the developed method. No significant difference ($p > 0.05$) in the results were observed from the proposed analysis and those obtained with HPLC-UV, indicating that the proposed method had great potential for histamine detection in red wine and rice wine samples.

Compared to the commonly used quantum dots (QDs) and organic dyes, UCNPs has lower autofluorescence background, better photochemical stability, larger Stokes shifts nonblinking property and lower toxicity. These outstanding photo-physical properties make it a potential candidate for further studies. In this study, by integrating fluorescence sensing and SERS, we have established a dual mode counterpropagating-responsive method for the detection of histamine. With this dual-mode system, positive SERS response and negative fluorescence response could be achieved simultaneously, which ensured more reliable and precise quantification of histamine compared with single fluorescence sensing or SERS-based method. We believe that our method shows excellent promise in the quantification of harmful chemicals in food products.

### 4. Conclusions

A novel sensor combined the sensitivity of the upconverting and SERS technologies and the specificity of MIPs was developed. Excellent reorganization specificity, wide linear range and a low detection limit were achieved using this sensing system. In addition, the fabricated system was successfully applied to detect histamine in red wine, rice wine and canned tuna samples with good recovery. The novel strategy demonstrated great potential in sensitive and economic detection of histamine in wine samples required by both government laboratories and food industry. This easy-to-operate method could be extended to detect other hazardous substances in food samples.

### Conflicts of interest

There are no conflicts of interest to declare.

### Acknowledgements

This research was supported by funds from USDA-ARS (No. 1935-42000-049-00D) in conjunction with the Center for Food Safety Engineering at Purdue University, the Nature Science Foundation of Jiangsu Province (No. BK20160168), the National Key Technology R&D Program for the 12th Five-Year Plan (No. 2016YFD0400304), the independent research project program of the State Key Laboratory of Food Science and Technology of Jiangnan University (No. JUSRP1029), the Nature Science Foundation of Jiangsu Province (No. BK20160168) and the graduate student innovation project of Jiangsu province (KYLX15_1134). The author Wu Zhengzong is thankful for the funding from Chinese Scholarship Council (CSC).

### References