

# Graphene Oxide-Reinforced Hollow Fiber Solid-Phase Microextraction Coupled with High-Performance Liquid Chromatography for the Determination of Cephalosporins in Milk Samples

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**Abstract** An effective and sensitive method to determine six cephalosporins (cefadroxil, cephapirin sodium, cefixime, cefuroxime sodium, cefoxitin sodium, and ceftiofur hydrochloride) in milk samples was developed using graphene oxide-reinforced hollow fiber solid-phase microextraction (GO-HF-SPME). After extraction, analytes were desorbed and analyzed using high-performance liquid chromatography-photodiode array detection (HPLC-PDA). The graphene oxide (GO) was dispersed in *N,N*-dimethyl formamide by ultrasonication and then immobilized into the pores of a hollow fiber (HF). Several factors of GO-HF-SPME experiment, such as the pH of the sample solution, type of organic desorption solvent, pH of the desorption solvent, extraction time, and desorption time, were optimized to achieve the highest extraction efficiency. Under the optimized extraction conditions, the method showed good linear ranges (0.05–10 µg/mL) with correlation coefficients higher than 0.9950, low limits of determination (LODs, 0.01–0.02 µg/mL), and good recoveries (71–108 %) at three different spiked levels. The results demonstrated that GO-HF-SPME would be a promising method for the enrichment of cephalosporins in dairy products.

**Keywords** Graphene oxide · Hollow fiber · Cephalosporins · High-performance liquid chromatography · Solid-phase microextraction

## Introduction

People have recently been paying increasing attention to noxious substances in their foods, surroundings, and medicines. However, most of these substances exist in intricate sample matrices at trace levels. Therefore, appropriate sample preconcentration techniques are urgently needed to effectively detect them. Globally, milk is one of the most consumed foods and is important for human health. Because it contains essential nutritional components such as proteins, calcium, and saturated fat, it is considered a necessary and inexpensive food. However, its quality can be impaired by the prevalent use of cephalosporin antimicrobials as feed additives in veterinary medicine (Samanidou and Nisyrinou 2008). Therefore, the monitoring of antibiotic residues is very important in controlling the safety of milk for human consumption.

In recent years,  $\beta$ -lactam antibiotics have been rapidly developed, where most attention has been focused on cephalosporin antibiotics. Cephalosporins were used for the treatment of infections caused by Gram-positive and Gram-negative bacteria. They function by inhibiting the synthesis of essential structural components of the bacterial cell wall. They are among the safest and most effective broad-spectrum bactericidal antimicrobial agents; therefore, they are the most prescribed (Oliveira et al. 2007). Cephalosporins are divided into four generations—first-, second-, third-, and, recently, fourth-generation compounds—according to the extent of their effect against Gram-negative activity. Third-generation cephalosporins are characterized by a broader antibacterial spectrum in comparison with the first- and second-generation agents because they are resistant to  $\beta$ -lactamase. Fourth-generation agents include molecules with a spectrum similar to those of the third-generation agents but with a markedly increased stability toward hydrolysis by  $\beta$ -lactamase (Pehourcq and Jarry 1998). Each generation exhibits a broader spectrum of activity

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than the preceding generation. These antibiotics currently hold a large share of the global market and can be considered one of the most important and most frequently used groups of antibiotics (El-Shaboury et al. 2007). Researchers have noted that method development for these antibiotics is substantially more difficult than that for other antibiotics because of the chemical instability of the common  $\beta$ -lactam nucleus and the minor differences in chemical structures among the analogues. Cephalosporin residues constitute a risk to human health because they can cause allergic reactions in hypersensitive individuals and can lead to the emergence of drug-resistant bacteria. Analysis of these residues plays a key role in ensuring food safety (Song et al. 2014).

Some methods have been successfully developed for the determination of cephalosporin antibiotics in biological samples using high-performance liquid chromatography (HPLC) with ultraviolet (UV) detection (Jin et al. 2014; Quesada-Molina et al. 2012; El-Shaboury et al. 2007). At the same time, a solid-phase extraction method was developed for the extraction of several cephalosporin antibiotics from aqueous solutions using carbon nanotubes as the adsorbent (Niu et al. 2007). Moreover, solid-phase microextraction, which allows a large sample volume to be concentrated and purified in a single step, has been widely applied in the analysis of trace levels of analytes in water samples. The sample pretreatment techniques used for the isolation of cephalosporins from various matrices are usually solid-phase microextraction (SPME) and liquid–liquid extraction (Ghidini et al. 2003; Holstege et al. 2002; Moats and Romanowski 1998; Choma et al. 2008; Liu et al. 2007). Since being discovered in 2004 (Novoselov et al. 2004), graphene, a new two-dimensional material comprising a single layer of  $sp^2$ -hybridized carbon atoms (Geim and Novoselov 2007), has become a topic of intensive research and has gained extensive attention in the materials sciences because of its various unique properties (Lee et al. 2008; Bolotin et al. 2008). Graphene has been reported to possess a high theoretical specific surface area ( $2630 \text{ m}^2/\text{g}$ ) (Stoller et al. 2008), which suggests a high sorption capacity. In addition, because of its large delocalized  $\pi$ -electron system, graphene can form strong  $\pi$ - $\pi$  stacking interactions with benzene rings (Cai et al. 2003). These unique properties make graphene superior for the adsorption of benzenoid compounds (Chen et al. 2010; Zhang and Lee 2011; Zhang et al. 2011). Graphene oxide (GO), a precursor to graphene after reduction, consists of a hexagonal carbon network bearing hydroxyl and epoxide functional groups on its “basal” plane, whereas the edges are mostly decorated by carboxyl and carbonyl groups (Boukhvalov and Katsnelson 2008; Kudin et al. 2008). These high-specific-surface-area functional groups can adsorb cephalosporins in milk samples. Over the past few years, GO has been used as an adsorbent in SPME (Ponnusamy and Jen 2011; Wang et al. 2014), solid-phase extraction (SPE) (Ning et al. 2014; Liu et al. 2011; Luo

et al. 2013), liquid-phase microextraction (LPME) (Payán et al. 2010), magnetic solid-phase extraction (MSPE) (Wang et al. 2012; Wu et al. 2011), and dispersive solid-phase extraction (DSPE) (Wen et al. 2014). Because milk is a complex matrix with high protein and fat contents, which often interfere with analytical procedures, emphasis is placed on sample preparation for extracting and determining cephalosporins in milk.

Excellent cleanup efficiency can be achieved with the hollow fiber (HF) playing the role of a filter. The large molecules (proteins and saturated fat) cannot permeate through the wall pores of the HF (Li et al. 2011), which makes it very applicable to complex matrix samples. Moreover, the HF is disposable after each use because of its low cost, which can overcome carry-over problems and enhance the reproducibility.

In view of the above, GO-reinforced HFs were prepared according to previous reports (Song et al. 2013; Ye et al. 2014). The aim of the present study was to develop and validate a simple, rapid, and cost-effective method for the simultaneous determination of the selected cephalosporin antibiotics (cefadroxil, cephapirin sodium, cefixime, cefuroxime sodium, cefoxitin sodium, and ceftiofur hydrochloride), which are widely used antibiotics, in milk samples using a method involving GO-HF-SPME coupled with HPLC. What is more, chromatographic conditions were successfully optimized by gradient elution method (Sun et al. 2014; Hu et al. 2015). The accuracy and precision of the HPLC analysis and method detection limits are described. The SPME optimization and stability of the analytes were also discussed in this study. The method was validated using fresh milk products from a supermarket.

## Materials and Methods

### Chemicals and Reagents

Cefadroxil (FAD), cephapirin sodium (HAP), cefixime (CFM), cefuroxime sodium (CXM), cefoxitin sodium (FOX), and ceftiofur hydrochloride (EFT) were purchased from Sigma-Aldrich (Germany). *N,N*-Dimethyl formamide (DMF), acetone, and acetic acid were obtained from the Beijing Chemical Plant (Beijing, China). Chromatographic-grade acetone, acetonitrile, and methanol were purchased from Merck Co. (Darmstadt, Germany). All other reagents were of analytical grade and were used as received, without further purification.

GO was supplied by the XFNano Materials Tech. Co., Ltd. (Nanjing, China). Accurel Q3/2 polypropylene HF membranes (200- $\mu\text{m}$  wall thickness, 600- $\mu\text{m}$  internal diameter (i.d.), 0.2- $\mu\text{m}$  average pore size) were provided by Membrane (Wuppertal, Germany). Ultrapure water ( $18.2 \text{ M}\Omega \text{ cm}^{-1}$ ) was obtained from a Milli-Q system (Billerica, MA, USA). All solutions were filtered through 0.22- $\mu\text{m}$  filters (Tianjin, China).

## Instrumental and Chromatographic Conditions

All separations were performed on a HPLC (Waters 2695, Waters Technologies, USA) equipped with a photodiode array detector (PDA, Waters 2998, Waters Technologies, USA). A reversed-phase  $C_{18}$  column (5  $\mu\text{m}$ , 150 mm  $\times$  4.6 mm i.d.) was used as the separation column. The data were acquired using the Empower software (Waters Technologies, USA). The gradient elution was performed with acetonitrile (solvent A) and a sodium acetate–acetic acid (NaAC–HAC) buffer solution (pH=3.4, solvent B). The elution program was as follows: 0.00–5 min, solvent A/solvent B (5/95, v/v); 5.00–15.00 min, solvent A/solvent B (25/75, v/v); and 15.00–20.00 min, solvent A/solvent B (5/95, v/v). The flow rate, injection volume, and UV wavelength were 0.8 mL/min, 10  $\mu\text{L}$ , and 254 nm, respectively. All of the mobile phases were filtered using a 0.22- $\mu\text{m}$  filter membrane and degassed for 30 min before use. The centrifuge was purchased from Sigma (3 K15, Germany), and the centrifugations were performed at 25  $^{\circ}\text{C}$ .

Scanning electron micrographs were obtained using an S4800 field emission scanning electron microscope (Hitachi, Japan) operated at 15 kV. Raman spectra (Renishaw, England) were obtained under appropriate conditions with excitation at 633 nm. The laser power was set to 50 %, and backscattered light was collected using a 50-fold objective lens following a 10-s exposure time. The other conditions are listed in the corresponding figures.

## Dispersion of GO

To prepare surfactant-stabilized GO dispersions, 60.0 mg of GO was sonicated in 20 mL of DMF for 2 h, after which GO was evenly dispersed in the DMF at a concentration of 3.0 mg/mL GO.

## Preparation of GO-Reinforced HF

The HF was cut manually into segments of 2 cm, ultrasonically cleaned in acetonitrile for 15 min to remove any possible

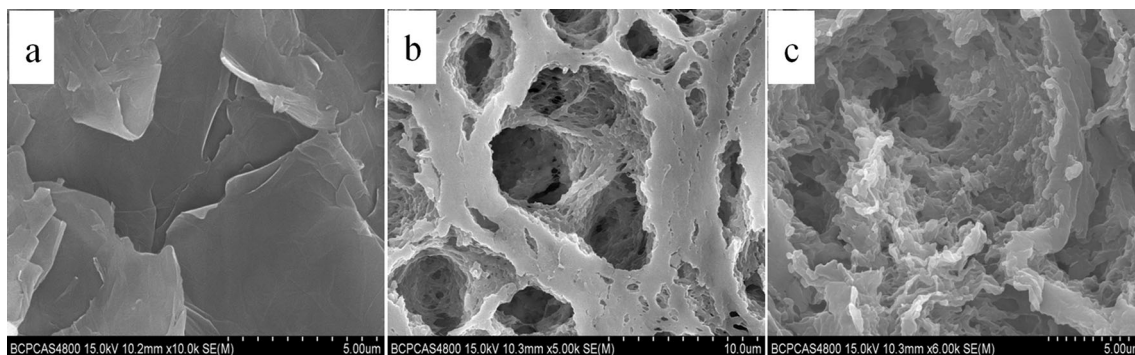
impurities, and then dried in air. To prepare completely immobilize GO onto the HF wall pores, HF segments were immersed in the homogeneous 3.0 mg/mL GO dispersion. GO was then adsorbed onto the wall pores of HF via sonication at room temperature for 2 h. Afterward, ultrapure water was used to remove excess GO from the surface and inner lumen of the HF. Finally, the prepared GO-HF was dried on a filter paper in air.

## Preparation of Standard Solutions and Milk Samples

The stock solutions of FAD, HAP, CFM, CXM, FOX, and EFT dissolved in acetonitrile were stored at 4  $^{\circ}\text{C}$ . Working solutions at different concentrations were freshly prepared daily by diluting the stock solutions with ultrapure water. Fresh milk product was purchased from a supermarket in Beijing (China) and was stored at 4  $^{\circ}\text{C}$ . Approximately 0.1 mL of milk, 0.2 mL of HAC, and a buffer solution of NaAC–HAC were added to a total volume of 1.5 mL, and the resulting solution was extracted under the optimized conditions.

## GO-HF-SPME Procedure

The HF was first cleaned with ultrapure water, and the treated GO-HF was then placed into 1.5 mL of standard solution (pH was adjusted to 4.0 using acetic acid) of the corresponding concentration. Next, the solution extraction process was performed for 30 min (the optimum extraction time) at an agitation rate of 1500 rpm. The GO-HF was subsequently removed, and the redundant solvents were absorbed by a filter paper. Finally, the GO-HF was placed in an end-sealed pipette tip with 50  $\mu\text{L}$  of methanol (pH was adjusted to 8.0 using sodium hydroxide) to desorb analytes for 50 min at an agitation rate of 1500 rpm. Fifty microliters of the desorbed solution was drawn out, and the resulting desorption solution was rapidly transferred into a sample bottle for analysis of the target by HPLC. To erase any possible carry-over effect, fresh GO-HF was used in each experiment. Common, unmodified HF was used to extract analytes under identical conditions using the aforementioned procedure and subsequently compared the enrichment performances of HF and GO-HF.



**Fig. 1** SEM images of GO (a), HF (b), and GO in the wall pores of the HF (c). Conditions: acceleration voltage, 15 kV

## Results and Discussion

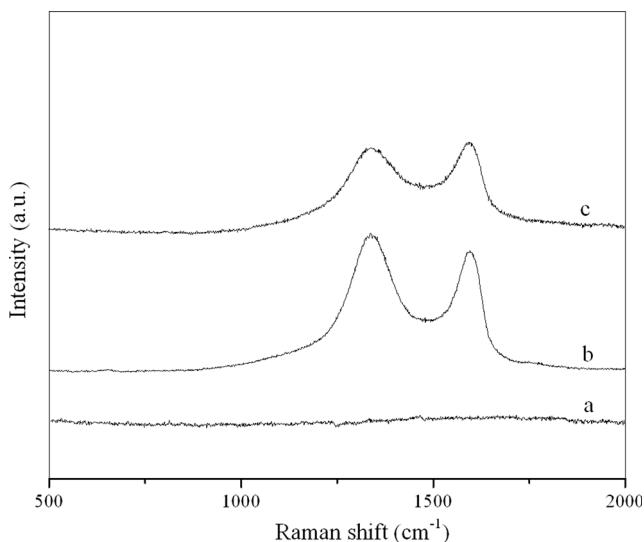
### Characterization of GO and GO-HF

GO, HF, and GO-HF were characterized by SEM. Figure 1a shows that the GO sheets assume a layered structure and exhibit wrinkled, thin, and smooth surfaces. As shown in Fig. 1b, the surface of the common HF is multi-aperture. Moreover, the holes were filled with GO after ultrasonication, as shown in Fig. 1c. A comparison of these micrographs indicates that GO was adsorbed onto the wall pores of the HF.

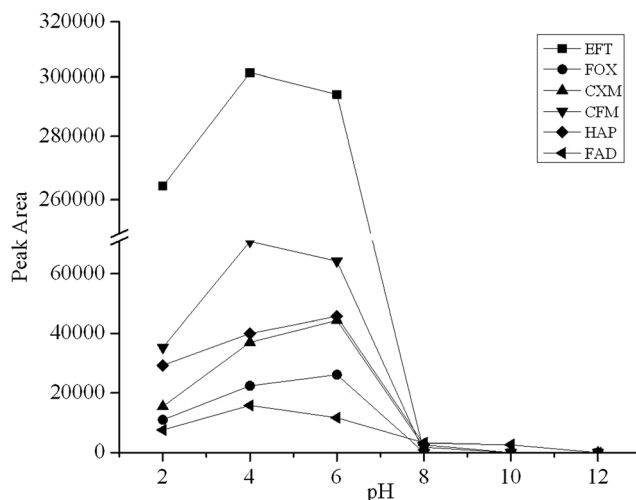
Raman spectroscopy is a perfect measure used to characterize the electronic states and structure of GO. The D and G bands represent disordered  $sp^3$  carbon and ordered  $sp^2$  crystalline graphite-like structures, respectively. Figure 2 shows that the commercial HF had no absorption peaks, and GO produced a D band (approximately  $1336\text{ cm}^{-1}$ ) and G band (approximately  $1596\text{ cm}^{-1}$ ). After HF was modified, GO-HF spectra show D band (approximately  $1342\text{ cm}^{-1}$ ) and G band (approximately  $1587\text{ cm}^{-1}$ ), which agreed with the usual features of carbon materials in the Raman spectra. Therefore, the Raman spectra indicate that GO was attached to the HF.

### Optimization of the Extraction Procedure

To obtain a high extraction efficiency and enrichment factor, a series of sample solutions spiked with  $1.0\text{ }\mu\text{g/mL}$  standard mixture was used to study the extraction performance under different experimental conditions. All of the experiments were performed in triplicate, and the means of the results were used for optimization evaluation.



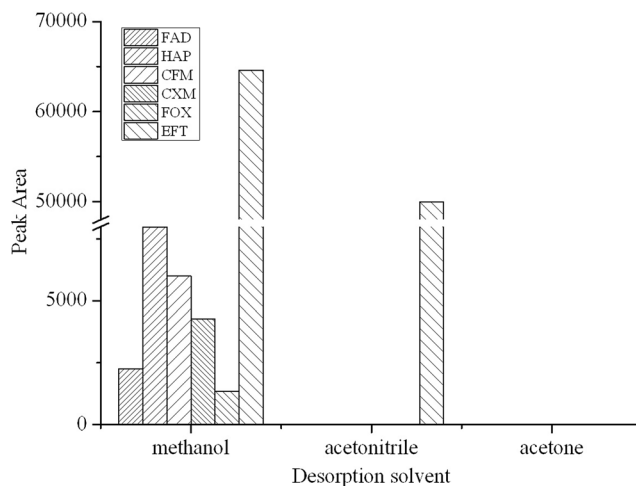
**Fig. 2** Raman spectra of HF (a), GO (b), and GO-HF (c)



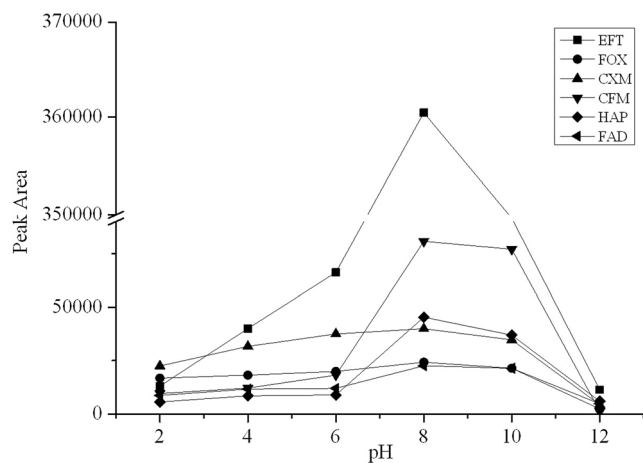
**Fig. 3** Effect of sample pH on the peak areas of cephalosporins extracted with GO-HF-SPME

### Effect of the Cephalosporin Extract Solution pH

A suitable pH of the sample solution can improve the extraction efficiency and reduce matrix interferences. Hydrophobic and  $\pi$ - $\pi$  interactions are the two main interactions between the GO and the analytes, and the pH should be adjusted such that the majority of the analytes remain in their molecular form to enhance the extraction efficiency. Therefore, the influence of the pH of the sample solution on the extraction efficiency was studied in the pH range from 2.0 to 12.0. As the results illustrate in Fig. 3, when the pH was greater than 6.0, the amounts of analytes extracted were substantially reduced because of the ionization of the target compounds. However, when the pH was 4.0, more target objects exist in the form of electric neutral molecules and cephalosporin was absorbed more easily by GO, making its peak area the largest. Therefore, the optimum pH of the sample solution was 4.0.



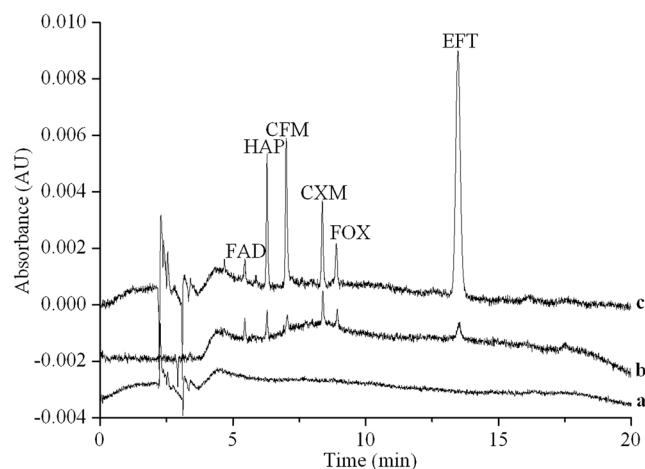
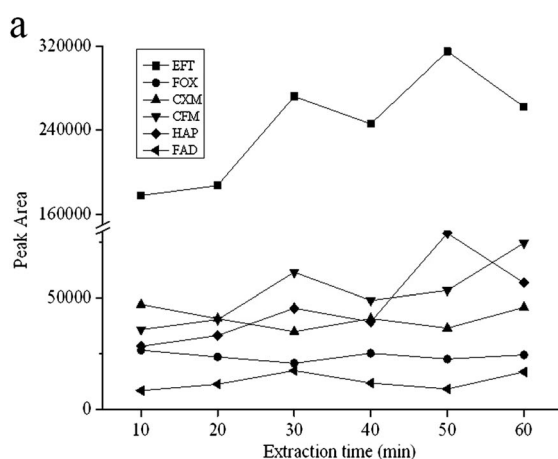
**Fig. 4** Effect of various desorption reagents on the peak areas of cephalosporins



**Fig. 5** Effect of the pH of the desorption solvent on the peak areas of cephalosporins

### Selection of the Organic Desorption Solvent

The polypropylene membrane and GO are hydrophobic, and low wettability was observed when GO-HF was directly exposed to a sample solution, which led to slow extraction rates and low extraction efficiencies. Solving these deficiencies required improvement of the wettability of the GO-HF. In this study, the wettability of GO-HF was enhanced by conditioning with an organic solvent that was suited to the fiber. Methanol, acetonitrile, and acetone, as three common organic solvents used for desorption, were investigated in the GO-HF-SPME procedure. The experiments demonstrated that methanol produced better results than acetonitrile and acetone, as shown in Fig. 4. The excellent wettability of methanol was demonstrated by its higher extraction efficiency. Thus, methanol was selected for subsequent experiments.



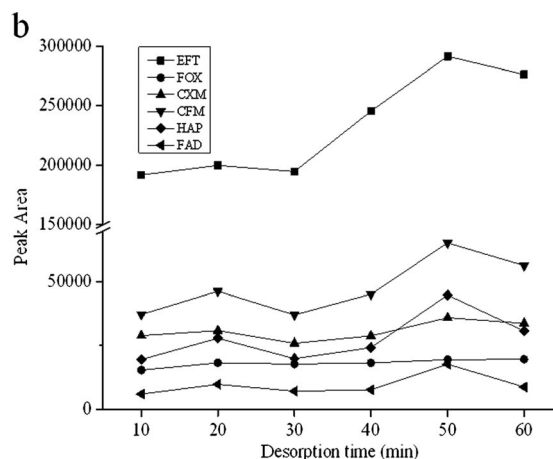
**Fig. 7** Chromatogram of a 0.2 µg/mL standard solution injected directly (b), treated by HF (a), and treated by the GO-HF-SPME method (c)

### Effect of the pH of the Desorption Solvent

In SPME, mass transfer was promoted by optimal pH conditions in the sample solution. In the present study, desorption was performed under different pH levels ranging from 2.0 to 12.0. As indicated by the results illustrated in Fig. 5, an increase in the peak area was observed in the pH range from 2.0 to 8.0. When the pH was adjusted to 8.0, the peak area reached its maximum and then decreased in the range of 8.0 to 12.0. Therefore, the optimum pH value of desorption solution was selected as 8.0.

### Effect of the Extraction Time

The extraction time is a very important parameter of the extraction procedure. The function of the extraction time was examined in the present study. The extraction and desorption of analytes on GO-HF are a dynamic balance; thus, the



**Fig. 6** Effects of different extraction times (a) and desorption times (b) on the peak areas of cephalosporins extracted by the GO-HF-SPME method

**Table 1** Performance parameters of the GO-HF-SPME method

Analytes	Linear range (µg/mL)	Linear equations	Regression coefficient	LOD (µg/mL)	LOQ (µg/mL)	RSD (%; <i>n</i> = 6) (intra-day)	RSD (%; <i>n</i> = 9) (inter-day)
FAD	0.1–10	$y = 9765x + 13391$	0.9995	0.02	0.06	5.5	8.3
HAP	0.1–10	$y = 48571x + 91719$	0.9998	0.02	0.06	5.8	6.9
CFM	0.05–10	$y = 39662x + 146558$	0.9951	0.01	0.03	4.3	5.9
CXM	0.05–5.0	$y = 28539x + 51176$	0.9979	0.01	0.03	5.7	6.7
FOX	0.1–10	$y = 13747x + 47517$	0.9978	0.02	0.06	3.8	5.7
EFT	0.05–10	$y = 428113x + 263997$	0.9960	0.01	0.03	4.7	6.4

extraction efficiency usually varies with the extraction time. Therefore, the extraction time was investigated over the range from 10 to 60 min, as shown in Fig. 6a. The peak areas of the analytes were increased as the extraction time was increased from 10 to 50 min. Given the comprehensive effects of the six cephalosporins and the high extraction efficiency, 30 min was used as the extraction time in subsequent experiments. Although this extraction time is relatively long, a large number of samples can be extracted simultaneously because of the simple experimental setup and the low cost of the GO-HF.

#### Effect of the Desorption Time

After extraction, the HF was transferred to a centrifuge tube containing methanol for desorption via vortex mixing. A series of desorption times was investigated in the range of 10–60 min at room temperature to ensure that all of the cephalosporins were completely desorbed from the GO-HF. As shown in Fig. 6b, the peak areas increased as the desorption time was increased from 10 to 50 min, after which they decreased slightly. This trend may result from the re-absorption of the desorbed analytes during the desorption period. The largest peak areas of the cephalosporins were obtained at a desorption time of 50 min. Therefore, 50 min was chosen as the optimal desorption time.

**Table 2** Analysis results of cephalosporins from real samples using the presented method

Analytes	Real sample (µg/mL)	Level 1 <sup>a</sup>		Level 2 <sup>b</sup>		Level 3 <sup>c</sup>	
		<i>R</i> (%)	RSD (%)	<i>R</i> (%)	RSD (%)	<i>R</i> (%)	RSD (%)
FAD	ND	92	6.8	89	8.6	83	7.2
HAP	ND	98	2.6	81	7.3	71	6.8
CFM	ND	108	4.2	98	5.8	94	4.3
CXM	ND	89	3.8	82	6.4	80	8.2
FOX	ND	96	4.8	92	4.6	87	3.6
EFT	ND	98	5.6	96	5.6	85	3.8

*R* recovery, *ND* not detectable

<sup>a</sup> Spiked 0.2 µg/mL six cephalosporins

<sup>b</sup> Spiked 2 µg/mL six cephalosporins

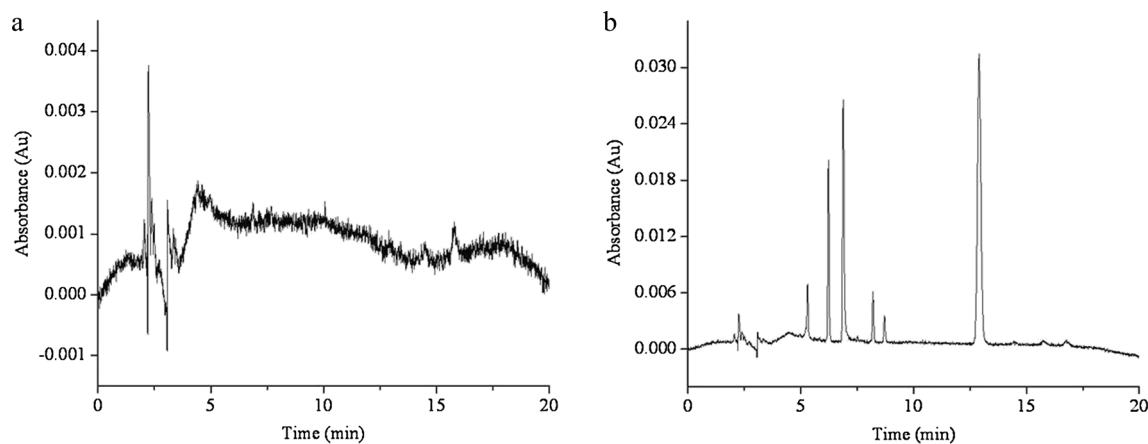
<sup>c</sup> Spiked 4 µg/mL six cephalosporins

#### Comparison Between HF and GO-HF

In this research, the unmodified HF and GO-HF were used separately to extract a 0.2 µg/mL standard solution mixture via the proposed SPME laboratory procedure; the results are compared in Fig. 7. As evident in this figure, the GO-HF resulted in a substantially greater peak area compared to that achieved with the common HF. GO-HF exhibited advantages in the extraction process (curve c), which are attributable to GO, whose high adsorption capacity and efficient extraction or desorption capability resulted in increases in both the amount of analyte extracted and the extraction efficiency. The six target analytes were only slightly detected when no preparation procedure was used (curve b), but they could not be detected using the proposed SPME procedure with unmodified HF (curve a). This result was likely a consequence of the HF lacking any special adsorption ability toward the six target analytes.

#### Evaluation of the SPME Method

The spiked samples were extracted using an established procedure and analyzed via HPLC-PDA. The performance parameters of the GO-HF-SPME technique, such as the linearity, low limits of determination (LODs), and limits



**Fig. 8** Chromatograms of **a** blank milk and **b** a milk sample spiked with 1.0 µg/mL of six cephalosporins extracted by the GO-HF-SPME method

of quantification (LOQs), for the cephalosporins were evaluated under the optimum extraction conditions. As shown in Table 1, the concentration of each of the cephalosporin samples was analyzed three times. The target analytes had satisfactory linear ranges and correlation coefficients. Moreover, the linear regression equations for FAD, HAP, CFM, CXM, FOX, and EFT were  $y=9765x+13391$ ,  $y=48571x+91719$ ,  $y=39662x+146558$ ,  $y=28539x+51176$ ,  $y=13747x+47517$ , and  $y=428113x+263997$ , respectively, where  $y$  represents the peak area of the cephalosporin and  $x$  denotes the concentration of the cephalosporin in the standard solution. The LODs and LOQs of the cephalosporins, which were investigated in standard solutions, were calculated at signal to noise ratios of 3 ( $S/N=3$ ) and 10 ( $S/N=10$ ), respectively. The LODs were 0.02 µg/mL for FAD, HAP, and FOX and 0.01 µg/mL for CFM, CXM, and EFT. Meanwhile, LOQs of 0.06 µg/mL for FAD, HAP, and FOX and 0.03 µg/mL for CFM, CXM, and EFT were obtained.

The intra-day and inter-day precisions of the instrument were investigated using the GO-HF-SPME technique via the repeated injection of a standard solution. The intra-assay precision was measured for six continuous injections during the same day, whereas the inter-assay precision was measured on

three consecutive days. The results are shown in Table 1. The relative standard deviations (RSDs) of the peak areas in the intra-assay precision were 5.5, 5.8, 4.3, 5.7, 3.8, and 4.7 % for FAD, HAP, CFM, CXM, FOX, and EFT, respectively, whereas the RSDs of the peak areas were 8.3, 6.9, 5.9, 6.7, 5.7, and 6.4 % for FAD, HAP, CFM, CXM, FOX, and EFT, respectively, in the inter-assay precision. The RSD values for all of the compounds ranged from 3.8 to 8.3 %, demonstrating that the GO-HF preparation method was quite stable and acceptable.

#### Analysis of the Cephalosporins in Milk Samples Using GO-HF-SPME

To assess the accuracy of the established analytical method, recoveries at three spiked concentration levels were tested. Three different concentrations of cephalosporin were spiked into the original sample, respectively. Then, three sets of milk samples were extracted using the same method as that previously optimized and analyzed. A summary of the recovery data obtained by HPLC is shown in Table 2. The recoveries ranged from 71 to 108 %, and the RSDs ( $n=3$ ) ranged from 2.6 to 8.6 %.

**Table 3** Analytical results of cephalosporins from the seven milk products ( $n=3$ )

Manufacture	Batch no.	Fat content (mg/mL)	Protein content (mg/mL)	Carbohydrate content (mg/mL)	Cephalosporin content (µg/mL)
A	A1	2.5	2.0	3.1	Unfound
	A2	2.3	2.0	3.3	Unfound
B	B1	2.4	2.0	3.0	Found <sup>a</sup>
C	C1	2.5	2.0	3.3	Unfound
	C2	2.5	2.0	3.2	Unfound
	C3	2.0	1.6	4.3	Found <sup>a</sup>
D	D1	2.7	2.3	3.3	Unfound

<sup>a</sup> Found: EFT could be found but could not be quantified

**Table 4** Comparison of several methods used for the detection of the target analytes

No.	Matrix	Analytes	Detection method	Extraction method	Volume of elution solvent (mL)	Linear range ( $\mu\text{g/mL}$ , $\mu\text{g/g}$ )	$R^2$	Recovery (%)	References
1	Raw cow milk	HAP EFT	UPLC-MS	DSPE	1	0.001–0.061 0.0006–0.102	0.994 0.981	96.0–104.5	Dorival-Garcia et al. 2016
2	Wastewater	CXM	HPLC-MS	SPE	0.5	0.0001–0.1	0.9990	63–91	Rossmann et al. 2014
3	Water	HAP	HPLC-PDA	LLE	1	0.0194–0.2 0.0282–0.4	0.9920 0.9930	80.6–105.5 67.5–99.7	Quesada-Molina et al. 2013
4	Meat	HAP	UPLC-MS	SPE	3	0.023–0.14	0.9986	98.2–103.4	Hou et al. 2013
5	Milk	EFT	UPLC-MS	SPE	6	0.001–0.05	0.9985	95.7–104.2	Gros et al. 2013
6	Wastewater	CXM EFT	UPLC-MS	SPE	6	0.001–0.05	0.9990 0.9971	71–133 40–77	Gros et al. 2013
7	Urine	HAP	HPLC-UV	MISPE	1	0.1–50	0.9989	70–129	Panahi et al. 2013
8	Milk	FAD EFT	HPLC-UV	MISPE	1	–	0.9998	92.3–95.7	Quesada-Molina et al. 2012
9	Milk	HAP FAD EFT	HPLC-PDA	MSPD	1	0.0192–0.24 0.0357–0.24 0.0331–0.24	0.9984 0.9944 0.9952	93.3–103.9 96.8–105.5 87.7–100.6	Karageorgou et al. 2012
10	Milk	CXM FAD EFT	HPLC-UV	MSPD	2	0.0462–0.2 0.005–0.2	0.9916 0.9999	93.8–101.9 95.5–103.6	Karageorgou and Samanidou 2010
11	Plasma	CXM	HPLC-PDA	SPE	2	0.017–0.16	0.9907	97.4–108.6	Nemutlu et al. 2009
12	Amniotic fluid	CFX	HPLC-PDA	SPE	2	0.035–50	0.9994	94.2–95.8	This study
13	Milk	FAD HAP CFX CXM FOX EFT	HPLC-PDA	SPME	0.05	0.05–10	0.9965 0.9995; 0.9998 0.9951; 0.9979 0.9978; 0.9960	98.3–99.7 71–108	This study

LLE liquid–liquid extraction, MISPE molecularly imprinted solid-phase extraction, MSPD matrix solid-phase dispersion extraction, UPLC-MS ultra-performance liquid chromatography–mass spectrometry, HPLC-MS high-performance liquid chromatography–mass spectrometry, HPLC-PDA high-performance liquid chromatography–photodiode array, HPLC-UV high-performance liquid chromatography–ultraviolet



In addition, real blank milk was extracted and analyzed to investigate the specificity of the method. Figure 8 shows typical chromatograms obtained from the analysis of the real blank milk and milk samples containing 1.0 µg/mL cephalosporins after extraction via the proposed procedure. No peaks were observed at the retention times of the corresponding analytes in the blank sample, which indicates that other coexisting species and the sample–matrix components did not interfere with precise analysis of the analytes. Then, the developed method was applied for the detection of cephalosporins in the different batches of milk sample from four manufactures. Contents of fat, protein, and carbohydrate in these milk samples were in the range of 2.0–2.7, 1.6–2.3, and 3.0–4.3 mg mL<sup>-1</sup>, respectively. Results shown in Table 3 indicated that EFT was found in two batches of milk but could not be quantified.

### Comparison of Some Methods Used for Determination of Cephalosporins

The extraction efficiency of the proposed method was compared with other reported sample preconcentration techniques from the angle of detection method, extraction method, the volume of elution solvent, linearity range, relative coefficient, and recovery. As listed in Table 4, the analytical property of the presented method is comparable with most of the listed methods. The GO-HF-SPME required relatively long extraction time and desorption time to obtain the highest extraction performance. Due to that preparation of dispersion of GO, the cleanup function of HF, and dilution of standard solutions take the place of the other, complicated steps were needed ahead of this microextraction. However, the procedure is simple and the extraction and desorption are completed using a vortex mixer. Of elution solvent, 0.05 mL was used in the developed method, making it green and cost-effective. The linear dynamic ranges and correlation coefficients were analogous to those reported sample preconcentration techniques. The large specific surface area and effective adsorption of GO-HF improve the extraction efficiency of this developed method. Thus, the novel mode using GO-HF as a new microextraction medium was useful and high stability.

### Conclusions

A new method based on GO-HF-SPME combined with HPLC was successfully developed for the determination of cephalosporins in milk samples. In this research, GO was well dispersed in DMF and adhered to the wall pores of HF via ultrasonic-assisted dispersion. The prepared GO-HF was then applied to SPME. In the present experiments, HF can prevent large molecules, such as proteins, from entering the small pores. The target cephalosporins were first adsorbed by GO-

HF and then desorbed by methanol via vortex mixing. Most importantly, the cleanup function of HF and selective adsorption of GO resulted in negligible matrix effects, and the sample required only simple homogenization and dilution with ultrapure water prior to the GO-HF-SPME procedure. Therefore, this method can be directly applied in relatively complicated samples. After the optimization of the extraction conditions for cephalosporins, relatively low limits of detection, effective sample cleanup, and acceptable precisions of repeatability as well as reproducibility were obtained. Therefore, this method is suitable as an alternative to other microextraction methods proposed for the enrichment of cephalosporins. In conclusion, the results of the present experiments showed that GO-HF-SPME coupled with HPLC is a rapid, simple, and cost-effective technique with a low consumption of organic solvent and is suitable for monitoring six types of cephalosporins in milk samples. In addition, the GO-reinforced HF-SMPE method would have potential applications for the concentration of other surrounding pollutants.

### Compliance with Ethical Standards

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**Conflict of Interest** Xinwei Chen declares that he has no conflict of interest, and Nengsheng Ye declares that he has no conflict of interest.

**Ethical Approval** This article does not contain any studies with human participants or animals performed by any of the authors.

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