MOLECULAR RECOGNITION OF SULFAQUINOXALINE AND SULFAPYRIDINE WITH MOLECULARLY IMPRINTED POLYMER

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ABSTRACT

The selective separation of sulfapyridine (SPD) from sulfaquinoxaline (SQX) is investigated by applying high performance liquid chromatography (HPLC) with molecularly imprinted polymer (MIP) as the stationary phase. Herein we report the synthesis of a molecularly imprinting polymer, SPD-MIP, by free radical polymerization process. Because of this powerful method synthetic polymers with specific binding sites to template molecule are provided. In addition, the separation performances were represented by using buffer/acetonitrile (3/2, v/v) as mobile phase under 272 nm UV detection. In order to compare the chromatographic data from the stationary phase, capacity factor (k') and separation factors (α) were given. The value of 2.71 (α) revealed that the MIP was able to recognize structurally subtle differences from the template molecule. Our results are discussed with regard to the amount of template, the composition of the chromatographic mobile phase and adsorption capacity.

Keywords: selective separation, sulfapyridine (SPD), sulfaquinoxaline (SQX), high performance liquid chromatography (HPLC), molecularly imprinted polymer (MIP)

INTRODUCTION

Determination of sulfonamides residues in food, especially in animal liver and kidney or milk, has received much attention in recent years. More than ten different kinds of sulfonamides are known to be used for domestic animals in Taiwan. Therefore, inspection of the residual sulfonamides is considered to be the most important duty for public health agencies. For years there has been a continuous interest for the development of analytical methods for the determination of sulfonamides in varying samples. The current methods for sulfonamides determination have been analyzed successfully by the liquid chromatography (LC) with ultra-violet detection1-4, fluorescence detection5-8, mass spectrometry9-10 or other11. These studies have been applied to pharmaceutical assays as well as in food sample. However, all of these methods involve tedious analytical procedures such as extraction complicated steps that are time consuming and give low resolution; sometimes recoveries are low and variable or must be expensive reagents and sophisticated instruments.

The aim of this work is to develop a simple, selective and more sensitive method for the identification and quantification of the sulfonamides (SQX and SPD, Table 1) by using molecular imprinting technique. In the molecular imprinting process by which functional and cross-linking monomers are co-polymerized in the presence of a molecular template12-14. After removal of the templates, polymers generate selective binding sites which are complementary to the template. Therefore, MIP is able to rebind the same template or its analogue. Besides, MIP can offer potential advantages in many fields, such as drug separation in analytical chemistry15-19 sensitive determination used as sensors20-22 and extraction. Of the various application areas of MIPS, separations are by far the broadest and most rapidly expanding.

Table 1.- Sulfonamide Structures, CAS numbers and molecular weights under this study.

<table>
<thead>
<tr>
<th>Structure</th>
<th>Name</th>
<th>CAS number (Molecular weight)</th>
</tr>
</thead>
<tbody>
<tr>
<td><img src="image1" alt="Structure 1" /></td>
<td>Sulfaquinoxaline (SQX)</td>
<td>59-40-45 (300.2)</td>
</tr>
<tr>
<td><img src="image2" alt="Structure 2" /></td>
<td>Sulfapyridine (SPD)</td>
<td>144-83-2 (249.3)</td>
</tr>
</tbody>
</table>

In our previous studies, we described the preparation of a stationary phase of HPLC based on ketoconazole-imprinted polymer23. This method proved to be fast and reproducible, showing linearity over a broad range of sample concentration. To further increase the utility of this technique, we have prepared sulfapyridine-imprinted polymer with methacrylic acid as the functional monomer molecule. The result polymers were packed in an analytical column and evaluated as the stationary phase of HPLC. Liquid chromatographic conditions (mobile phase composition, sample loading, flow rate) were altered in order to achieve the highest resolution. The MIP was successfully applied to the direct detection of SQX and SPD in analysis.

EXPERIMENTAL

Material

Sulfaquinoxaline (SPD), sulfaquinoxaline sodium (SQX) were purchased from Sigma (St. Louis, MO, USA). Ethylene glycol dimethacrylate (EGDMA, 98%) and methacrylic acid (MAA, 99%) which were distilled to remove the inhibitors in previous using were obtained from Merck (Germany). Acetonitrile, acetone, ethanol, methanol, sodium phosphate and phosphoric acid were purchased from TEDIA (Fairfield, OH, USA). These were all HPLC grade.

Chloroform and acetic acid were from R.D.H. (GC grade) while 2.2'-Azoobisobutyronitrile (AIBN) was obtained from TCI (Tokyo, Japan). Water was double de-ionized.

Preparation of the molecularly imprinted polymer

A conventional imprinted polymer with SPD as the template were prepared, referred as SPD-MIP. In a conical Erlenmeyer flask with a screw top, SPD (template, 1 mol%) and AIBN (initiator, 0.56 mmol) were dissolved in acetone (10 mL) and then EGDMA (93 mol%), MAA (6 mol %) were added. The flask was placed in an ultrasonic water bath until clear solutions were obtained, then cooled on ice and the solutions sparged with nitrogen. The flask was placed under a UV-lamp (365 nm) at 4 °C for 2 h. The resulting polymers were dried in a vacuum oven for 24 h at room temperature. The hard polymers were ground in a laboratory mortar grinder.

The particles were sieved to collect particles 25–45 μm in size. Grinding and sieving were repeated until all material passed the 45 μm sieve. Non-imprinted reference polymers were prepared using the same conditions with the exception that template was omitted. The particles were suspended in methanol (30 mL) by sonication for 3 min, placed in a slurry reservoir, and was then packed in a stainless steel column (150 x 4.6 mm I.D.) using an air-driven fluid pump. The particle weight in each column was approximately 3.32 g. The packing was carried out under a pressure of 300bar with acetone (300 mL) as the packing solvent. All polymers were washed on-line with methanol-acetic acid (9:1, v/v) mixture at a flow rate of 0.5 mL/min until no further template
bleed could be detected using HPLC.

**HPLC-analysis**

The HPLC system consisted of a JASCO PU-2080 chromatograph with a JASCO UV-2075 variable wavelength detector and a Rhodyne 7725 syringe loading sample (20 μL) injector. For data analysis peak integration was performed using a Peak ABC Chromatography Workstation Ver.2.10 integrator. The apparatus provide assurance that all the UV-absorbing components are detected, if present in sufficient quantity. Analyses were run at flow-rates of 1.0 mL/min with detection at 272 nm. Capacity factors (k') were determined from k' = (t - t₀)/t₀, where t is the retention time of a given species and t₀ is the elution time of the void marker (determined by injection of toluene). Effective separation factors (α) were calculated from the relationship α = K'_{SPD}/K'_{SQX}, where K'_{SPD} and K'_{SQX} were the capacity factors of the SPD and SQX, respectively.

**Binding Experiment**

The concentration of standard solutions was 20 μmol/ L–1.0 μmol/ L substrate (SPD and SQX). A calibration graph was found between concentration of substrate and the absorbance. An accurately weighed 5mg portion of the polymer particles was transferred into a 50 mL centrifuge tube, 15 mL of vary concentration standard solution was added and the tube rotary. The SPD imprinted particles were equilibrated in the aqueous solution of SPD or SQX analogs at 4 °C for 12 hours. This solution was centrifuged at 2000 rpm for 10 min at 4 °C. The centrifugate was transferred into 50 mL volumetric flask. The absorbance of the solution was measured by HPLC with UV detector after the substrate bound. The concentration of substrate was estimated based on the standard curve. The value of substrate bound to the imprinted particle, [C] (μmol/g), was calculated according to the equation:

\[ [C] \text{ (μmol/g)} = (C_0 - C)V/g \]  

where C₀ and C were the molar concentrations (μM) of SPD or other substrates after and before equilibration. V and g were volume of substrate solution and weight of imprinted particles, respectively.

**RESULTS AND DISCUSSION**

**Characterization of the SPD-imprinted polymer**

The SPD-MIP was prepared utilizing MAA as the functional monomer, which acts as both a hydrogen bond donor and acceptor with the imprint species. The bulk polymer was ground into powder, packed into HPLC columns and rendered it suitable for a stationary phase. The template was extracted from the polymer using MeOH/AcOH (9:1; v/v) solution.

The solvent plays an important role in formation of the porous structure in MIP. It is also clear that the polarity of solvents used in the imprinting analyses affects the specificity of the polymers. Non-polar porogens such as toluene or polar porogen such as acetonitrile did not sufficiently dissolve SPD at low temperature. However, acetonitrile is a non-protonic solvent that may not make the hydrogen bonding formation in the polymerization system. Thus, using acetone as the polymerization solvent is advantageous for the present imprinting, rebinding to the template. In this study, acetonitrile is successfully used as solvent for preparing polymer-introduced recognition sites for SPD.

The results of this study show that better selectivity is obtained at the lower temperature (4 °C) polymerization versus the identical polymers thermally polymerized. To polymerize at lower temperatures, it is necessary lower temperature (4 ℃) polymerization versus the identical polymers thermally polymerized. It was found that the value [C]_{ksp} of SPD binding the imprinted particles decreased from 21.52 μmol/g to 14.96 μmol/g, when the temperature of polymerization increased from 4 °C to 65 °C. Meanwhile, the value [C]_{ksp} of SQX decreased from 15.12 μmol/g to 13.56 μmol/g. It is clearly showed that the binding amounts of SPD to the imprinted particles were much higher than those of SQX. This indicated that in the MIP prepared with low temperature, the binding ability of SPD to the imprinted polymer increased because recognition site formation made homogeneity of the template in the particle. Therefore, the increased numbers of the imprinted sites led to the increased capacity of SPD binding when the temperature was 4 °C in the polymerization process. Furthermore, the binding amount of blank polymers (non-MIP) was clearly lower than that of MIP, showing that the absorbivities of blank polymer were physical adsorption (non-selective). Non-MIP lacked template molecules during polymerization, thus polymers would not be imprinted. Evidence of the binding results was the fact that this polymerization process was capable of imprinting the SPD shape into the polymer and at lower temperature was efficient to fix the shape of SPD template. In addition, the extraction manner left SPD imprinting sites in the polymer matrix. As a result, the recognized SPD was shown efficiently in the imprint particles. Overall, it can be seen that the MIP prepared in this research showed significant selectivity to the print molecule alone.
indicated by the imprinted The ΔG on the imprinted particles and completely was evaluated with separation factor as described in shown in 20 mixture of these two compounds was injected for analysis in a total volume of 50 μL. As described above we can estimate the selectivity of MIPs by recovery measurements of an analyte and other structurally related analogues.

The retention time for SPD and SQX were 0.96 min and 1.02 min when the flow rate was 1 mL/min.

Table 2.- For SPD on the imprinted polymers. Fitted parameters for the Langmuir plots to estimate the binding parameters of SPD-MIP.

<table>
<thead>
<tr>
<th>MIP system</th>
<th>K (L/mol)</th>
<th>Qmax (μmol/g)</th>
<th>ΔG° (kJ/mol)</th>
</tr>
</thead>
<tbody>
<tr>
<td>4 □</td>
<td>5.55</td>
<td>22.30</td>
<td>-4.24</td>
</tr>
<tr>
<td>65 □</td>
<td>4.54</td>
<td>18.23</td>
<td>-3.74</td>
</tr>
</tbody>
</table>

From the slope and intercept of the plot, the equilibrium constant K and the apparent maximum number Qmax of the higher affinity binding sites can be calculated to be 5.55 L/mol and 22.30 μmol/g for MIP prepared at 4 □, respectively. In the same way, K and Qmax of the lower affinity bonding sites were calculated to be 4.54 L/mol and 18.23 μmol/g for MIP prepared at 65 □, respectively. Table 2 shows the values of ΔG° calculated from K by using Eq.(3), indicating that SPD interacts most strongly with the polymers. Therefore, the MIP synthesized with MAA is expected to give the highest selectivity to SPD. As described, above we can estimate the selectivity of MIPs by recovery measurements of an analyte and other structurally related analogues.

**Molecular recognition of SPD and SQX with molecularly imprinted polymer as the stationary phase of HPLC**

In single or binary substrate solution, 1 g/L of SPD and SQX solution or mixture of these two compounds was injected for analysis in a total volume of 20 μL and eluted isocratically at a flow-rate of 1.0 mL/min. Subsequently, the imprinting effect was evaluated by HPLC with UV detection. The results were shown in Figure 3. Molecular recognition effect of SPD from the binary mixture solution was evaluated with separation factor as described in experimental section. Validation revealed that the retention time of SPD and SQX were 7.96 min and 3.63 min when the flow rate was 1 mL/min.

![Figure 2](https://via.placeholder.com/150)

**Figure 2.** Langmuir plots to estimate the binding parameters of SPD-MIP.

![Figure 3](https://via.placeholder.com/150)

**Figure 3.** Chromatograms at 272 nm of SPD and SQX by HPLC using molecularly imprinted polymer as the stationary phase.

Table 3 shows the retention time, retention factor and separation factor (α) values obtained after separation by HPLC. The SPD and SQX are readily and rapidly separated when 34 mmol/mL NaH2PO4 buffer solution/ acetonitrile (3:2, v/v) is used as the mobile phase. From Table 3, the retention times for SPD and SQX were 7.77~7.95 min and 3.55~3.64 min, respectively. The difference of retention time (tR) between SPD and SQX were 4.14~4.41 min. The chromatographic run was completed in less than 10 min and completely separated from the other peaks under a flow rate of 1 mL/min. It was found that the values of α obtained for the imprinted particles prepared at 4 □ were 2.55~2.71 for SPD/SQX, which was a nearly constant. The SPD-imprinted polymer prepared at 4 □ showed higher selectivity to SPD than that of 65 □, further evaluation was carried in binary mixture with a total of 1g/L.

The retention time for SPD and SQX were 0.96 min and 1.02 min when used blank polymer (non-MIP) as the stationary phase, and the difference of retention time (tR) between SPD and SQX was 0.06 min. It can be seen that the blank polymer had little selectivity for SPD and SQX, while the retentivity and selectivity of the imprinted polymer for the two molecules were greatly strengthened by molecular imprinting as indicated by the imprinted particles. The good imprinting effect was likely due to the carboxylic group on the monomer and the electrostatic interaction or hydrogen bonding between imprinted polymer and template molecules in the polymerization. Apparently, the SPD-imprinted particles selectively bound SPD rather than SQX. Although SQX was structurally close to the SPD template, the retentions of SPD were higher than SQX on the imprinted polymers. These results indicated that imprinted polymers formed recognition sites for the SPD compound as the template molecules. The significant difference in the retention was due to the fact that SPD molecules were removed by washing the polymer matrix with an acetic acid/methanol solution, and therefore the polymers left cavities of a complementary size and shape, providing a great deal of free amino residue groups in the copolymer matrix. The higher population of free amino residue group in the polymer particles seems to increase the affinity of the solutes.
reason is that the removal of SPD leaves a clear cavity whose morphology is similar to SPD on the particles. Thus the stationary phase utilized the cavity and the free amino residue groups caught SPD. Therefore, these data strongly suggested that the SPD imprinted polymer had effective selectivity by hydrogen bonding to separately bind in the binary components.

Table 3.- Composition of solution with a total of 1g/L and resolution of SPD and SQX used MIP as the stationary phase of HPLC.

<table>
<thead>
<tr>
<th>Solution No.</th>
<th>Concentration in sample</th>
<th>Retention time, min (capacity factor)</th>
<th>α (separation factors)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>SQX SPD (g/L)</td>
<td>SPD</td>
<td></td>
</tr>
<tr>
<td>1</td>
<td>1.0 0.0</td>
<td>3.55</td>
<td>2.55</td>
</tr>
<tr>
<td>2</td>
<td>0.9 0.1</td>
<td>3.63 (2.78) 7.77 (7.09)</td>
<td>2.71</td>
</tr>
<tr>
<td>3</td>
<td>0.7 0.3</td>
<td>3.55 (2.69) 7.96 (7.29)</td>
<td>2.60</td>
</tr>
<tr>
<td>4</td>
<td>0.5 0.5</td>
<td>3.64 (2.79) 7.94 (7.27)</td>
<td>2.61</td>
</tr>
<tr>
<td>5</td>
<td>0.3 0.7</td>
<td>3.63 (2.78) 7.94 (7.27)</td>
<td>2.66</td>
</tr>
<tr>
<td>6</td>
<td>0.1 0.9</td>
<td>3.59 (2.73) 7.95 (7.28)</td>
<td></td>
</tr>
<tr>
<td>7</td>
<td>0.0 1.0</td>
<td>------</td>
<td></td>
</tr>
</tbody>
</table>

*the retention time of toluene was 0.96 min.

CONCLUSION

SPD-imprinted polymers were successfully prepared by bulk polymerization at 4℃ in acetone solvent. It was efficient to fix the shape of SPD template into the particle through hydrogen bonding with low temperature. However, under high temperature treatment it showed lower effect on imprinting. Results showed that SPD imprinted particle with 4℃ preparation provided high recognition and selectivity to SPD rather than SQX. The SPD-imprinted particle had effective separation property of the binary components. Therefore, it was meaningful to develop SPD recognition materials applying in the various fields such as chromatographic separation, biosensors and drug therapy.

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REFERENCES