Mechanisms of P-Glycoprotein Modulation by Semen Strychni Combined with Radix Paeoniae Alba

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Abstract
Semen Strychni has been extensively used as a Chinese herb, but its therapeutic window is narrowed by the strong toxicity of the compound, which limits its effectiveness. Radix Paeoniae Alba has been reported to reduce the toxic effects and increase the therapeutic effects of Semen Strychni, but the underlying mechanism remains unknown. This research aimed to explore the mechanism through which P-glycoprotein (P-gp) is modulated by Semen Strychni combined with Radix Paeoniae Alba in vitro. An MTT assay was used to study cytotoxicity in an MDCK-MDR1 cell model. Rh123 efflux and accumulation were measured to assess P-gp function. The expression levels of MDR1 mRNA and P-gp protein in MDCK-MDR1 cells were investigated. A P-gp ATPase activity assay kit was applied to detect the effect on P-gp ATPase activity. Semen Strychni combined with Radix Paeoniae Alba could induce P-gp-mediated drug transport by inhibiting brucine and strychnine transport in MDCK-MDR1 cells, enhancing the P-gp efflux function, upregulating the P-gp expression and MDR1 mRNA levels, and stimulating P-gp ATPase activity.

1. Introduction

Semen Strychni has been used in the treatment of joint pain, arthritis, and rheumatic diseases in China. Its main bioactive components, Strychnos total alkaloids, which comprise more than 70% of the alkaloids isolated from Semen Strychni, have been shown to exert antitumour, analgesic, anti-inflammatory, antioxidant, and antiangiogenesis effects as well as other practical effects [1]. Although it is widely used as a folk medicine, the use of Semen Strychni is limited by its narrow therapeutic window due to its strong excitatory effects on the central nervous system [2]. Some studies have demonstrated the appearance of persistent or transient neurovirulence in humans who excessively use Semen Strychni or its representative constituents, brucine, and strychnine. Therefore, it is very important to find a method to protect against the toxicity caused by Semen Strychni.

Protective herbal therapy is considered a reliable method that can reduce the toxicity of Semen Strychni and increase its use in clinical practice. Radix Paeoniae Alba has been reported to have wide-ranging pharmacological activities on the nervous system [3] and has been combined with Semen Strychni in the clinic and in the prescription of Chinese patent drugs, such as Gù Cí tablets, Shu Luo Yang Gan pills, and Qi Wei Tang. Radix Paeoniae Alba reportedly decreases the toxic effects and increases the therapeutic effects of Semen Strychni [4–6]. Additionally, Radix Paeoniae Alba and its active ingredients have shown protective effects in previous cytotoxicology experiments [7, 8].

P-gp is an energy-dependent transporter that relies on the energy released by ATP hydrolysis to transport the substrate to the extracellular domain. It can affect drug absorption in the intestinal tract, drug distribution to the brain, and drug elimination by the liver and kidneys [9]. Researchers have identified several Chinese herbal medicinal ingredients that can mediate P-gp efflux, including tetrandrine [10], dauricine [11], and quercetin [12]. Brucine and strychnine have been shown to be potential P-gp substrates [13]. Some research has indicated that P-gp takes part in the transport of brucine at the blood-brain barrier (BBB) [14]. As a result, the induction of P-gp activity can promote the removal of toxic drugs from the cell and meanwhile may cause potential herb-herb interactions.

Keywords: Semen Strychni; Radix Paeoniae Alba; P-glycoprotein; efflux; toxicity
2. Materials and Methods

2.1. Reagents. The RNA extraction reagent TRIzol was obtained from Ambion (USA). The PrimeScript RT reagent kit and Pgp-Glo® assay system were purchased from Promega (USA). The PowerSYBR Green PCR Mix was obtained from Life Technologies, Inc. (USA).

2.2. Preparation of RPAE and SSE. We obtained Radix Paeoniae Alba extract (RPAE, containing 57.18% paeoniflorin, as determined by HPLC) as reported in the literature [17]. We also obtained Semen Strychni extract (SSE, containing 17.70% brucine and 35.95% strychnine, as determined by HPLC) as reported in the literature [18].

2.3. Cells. MDCK-MDRI cells were bought from the Zhongya Biological Gene Institute (China) and cultured in DMEM containing 100 U/mL penicilllin, 1% NEAA, 100 μg/mL streptomycin, and 10% FBS at 37°C in an atmosphere of 5% CO₂ at 95% relative humidity [19, 20]. The cells were passaged using a trypsin-EDTA solution every three to four days when they had grown to 80%–90% confluence.

2.4. MTT Assay. The cells (5 × 10⁴ cells/well) were seeded into a 96-well culture plate and cultured for 24h. Fresh medium containing different test drugs was added to the relevant wells. After 4h, the culture medium was removed, and the cells were washed twice with HBSS. Next, 20 μL of MTT was put into each well and removed after 4h. Then, add 200 μL of DMSO to every well. The plate was incubated with shaking for 10 minutes, and the absorbance at 490 nm was read using a microplate reader (BioTek Elx800, USA). Wells with no cells were used as blank control wells, and wells which contained cells without the test drug were used as negative control wells.

2.5. Transport Experiment. Cells (1 × 10⁵ cells/cm²) were seeded onto polycarbonate filter membranes in a 12-well plate. The cell monolayers could be used for transport studies after the cells were allowed to grow for five to seven days. The transepithelial electrical resistance (TEER) of each MDCK-MDRI cell monolayer was measured and exceeded 500 Ω/cm². The cell monolayers were washed twice with HBSS and then incubated for 30 min. The HBSS was removed, and test drug solutions were added to either the apical (AP, 0.5 mL) or basolateral side (BL, 1.5 mL). Samples were taken from the receiving side after 2h incubation. The samples were diluted with the same volume of methanol, mixed, and centrifuged at 16000 r/min at 4°C for 20 min to precipitate proteins. Brucine and strychnine were determined by LC-MS/MS.

2.6. LC-MS/MS Analysis. Brucine and strychnine in the samples were analysed by LC-MS/MS using an AB Triple Quad 4500 system (AB SCIEX). Brucine, strychnine, and the internal standard matrine were separated on a Gemini C₁₈ column (150 mm × 4.6 mm ID, Phenomenex, USA) using acetonitrile-water containing 10 mmol/L ammonium acetate (30:70) as the mobile phase. An electrospray ionization (ESI) source was used and operated in the positive-ion mode. Brucine, strychnine, and matrine were detected at m/z 395.3 → 324.4, m/z 335.2 → 184.2, and m/z 249.3 → 148.3, respectively. Linear calibration curves were obtained for brucine and strychnine in the concentration range of 1.02–102.20 ng/mL. The lower limits of quantitation for brucine and strychnine, defined as a signal-noise ratio of 10, were 1.02 ng/mL and 1.20 ng/mL, respectively. The extraction recoveries were 95.75%–102.54% and 95.82%–104.64%, respectively. The precision, accuracy, and stability of the analytes met the requirements. The results showed that the method was effective and convenient for the detection of strychnine and brucine in the transport samples.

2.7. Flow Cytometric Analysis of the Intracellular Accumulation of Rh123. For the flow cytometric analysis, 1 mL aliquots of the cell suspension (1 × 10⁶ cells/mL) were added in Eppendorf (EP) tubes, and the test drug solutions were put into the tubes. The mixtures were incubated for 1h at 37°C. 1 mL of Rh123 (10 μM) was added and the mixtures were incubated for 1h at 37°C and then resuspended in 0.5 mL of HBSS. The samples were detected by flow cytometry [21]. Verapamil is a well-known P-gp inhibitor as a positive control medicine.
2.8. **qRT-PCR Analysis of MDRI mRNA Expression.** The cells (5 × 10^4 cells/well) were seeded into a 12-well culture plate and then cultured for five days. The cells were incubated with test drugs for 2 h and washed twice with HBSS. Total RNA was extracted using TRIzol on the basis of the manufacturer’s specifications. First-strand cDNA synthesis and amplification were performed on the basis of the Prime Script RT reagent kit protocol. qRT-PCR was performed as described in the Power SYBR Green PCR Mix protocol with a real-time fluorescent quantitative PCR instrument (Prist 7500, ABI, USA). The following primers (Sangon, China) were used: MDRI (F) 5′-TGG GGC TGG ACT TTC CAT GAT GC-3′, MDRI (R) 5′-GCA GCA ACC AGC ACC CCA GCA CCA AT-3′, RpPip1v (F) 5′-CCC TCA TTC TGC ACG ACG AT-3′, and RpPip1v (R) 5′-GGC TCA ACA TTT ACA CCG GC-3′. The RPP1 mRNA levels were used for normalization. The fold-change in gene expression was calculated using the 2^{−ΔΔCT} method.

2.9. **Western Blotting Analysis of the Effect of SSE Combined with RPAE on P-gp Expression.** Cells (5 × 10^4 cells/plate) were seeded into 10 cm culture plates and cultured for five days. The cells were treated with the test drug solutions for 24 h and then washed twice with HBSS. The cells were lysed with RIPA Lysis Buffer containing protease inhibitors and phosphatase inhibitors on ice on the basis of the manufacturer’s specifications. The protein concentration of the samples was determined according to the BCA assay protocol, and equal amounts of protein were added to the Power SYBR Green PCR Mix protocol with a real-time fluorescent quantitative PCR instrument (Prist 7500, ABI, USA). The following primers (Sangon, China) were used: MDRI (F) 5′-TGG GGC TGG ACT TTC CAT GAT GC-3′, MDRI (R) 5′-GCA GCA ACC AGC ACC CCA GCA CCA AT-3′, RpPip1v (F) 5′-CCC TCA TTC TGC ACG ACG AT-3′, and RpPip1v (R) 5′-GGC TCA ACA TTT ACA CCG GC-3′. The RPP1 mRNA levels were used for normalization. The fold-change in gene expression was calculated using the 2^{−ΔΔCT} method.

2.10. **P-gp ATPase Activity Assay.** P-gp ATPase activity was measured using Pgp-Glo Assay System which provided the necessary reagents for performing luminescent P-gp ATPase assays [21–23]. P-gp is an energy-dependent protein, which relies on the energy released by ATP hydrolysis to transport the substrate outside the cell. However, ATP enzymes need to be activated during the hydrolysis of ATP. So compounds that interact with P-gp can be identified as stimulators or inhibitors of its ATPase activity.

The Pgp-Glo Assay detects the effects of compounds on recombinant human P-gp in a cell membrane fraction. The assay relies on the ATP dependence of the light-generating reaction of firefly luciferase. ATP is first incubated with P-gp; then the P-gp ATPase reaction is stopped, and the remaining unmetabolized ATP is detected as a luciferase-generated luminescent signal. Pgp-dependent decreases in luminescence reflect ATP consumption by P-gp; thus, the greater the decrease in signal, the higher the P-gp activity. Accordingly, samples containing compounds that stimulate the P-gp ATPase will have significantly lower signals than untreated samples.

Pgp-Glo Assay System included Pgp-Glo assay buffer, ATP detection substrate (lyophilized), recombinant human P-gp membranes, MgATP, ATP detection buffer, Na₂VO₄, and verapamil. The Pgp-Glo assays were performed in two steps. The first step was the P-gp reaction. In this step, a recombinant human P-gp membrane fraction was incubated in Pgp-Glo assay buffer with MgATP (5 mM) for 40 min at 37°C. Untreated (NT) and Na₂VO₄ (100 μM) treated control samples were included except for verapamil (200 μM) treated samples (positive control). When an ATP standard curve was used, the ATP standards were added to the plate at the end of this step. The second step was the ATP detection reaction. ATP detection reagent was added to the P-gp reaction described above and incubated for 20 min at 37°C. The samples were measured by a Varioskan Flash reader (Thermo Scientific). The statistics of the luminescent signals were on the basis of the manufacturer’s specifications, and the influence of the test drug on P-gp ATPase activity was then estimated.

2.11. **Statistical Analysis.** The data were analysed by SPSS 19.0. The statistical difference between two groups was measured by t-test.

### 3. Results

3.1. **MTT Assay.** The MTT assay results shown in Figure 1 indicated that SSE was nontoxic below the concentration of 1.06 μg/mL and that RPAE was nontoxic in the range of 0–203.40 μg/mL. When SSE and RPAE were combined, the nontoxic concentration range of SSE increased significantly. In particular, when the ratio of SSE to RPAE was 1:3, the nontoxic concentration range of SSE was 0–8.00 μg/mL.

3.2. **Effect of RPAE on Brucine and Strychnine Transport in MDCK-MDRI Cells.** As shown in Tables 1 and 2, the \( \text{Papp}_{\text{AP/BL}} \) values of brucine and strychnine in the SSE + RPAE groups were lower than those in the SSE group significantly, whereas the \( \text{Papp}_{\text{BL/AM}} \) values were higher. Moreover, the ER values in the SSE + RPAE groups were higher than those in the SSE group. As the RPAE concentration increased, the ER value increased. These results showed that the combination of SSE and RPAE inhibited the absorption of brucine and strychnine. In addition, the ER values were significantly lower in the groups treated with verapamil than in the groups without verapamil, indicating that P-gp was participated in the transport of brucine and strychnine.
3.3. Effect of RPAE and/or SSE on the Accumulation of Rh123.

The accumulation of Rh123 was determined by flow cytometry. As shown in Table 3 and Figure 2, RPAE and SSE both decreased the intracellular accumulation of Rh123 compared with the control group. The intracellular accumulation of Rh123 in the SSE + RPAE groups was significantly lower than that in the SSE group, indicating that RPAE could activate the efflux function of P-gp.

3.4. Effect of RPAE and/or SSE on the Relative Expression Level of MDRI mRNA. As shown in Figure 3, in contrast to the control group, the relative expression level of MDRI mRNA was decreased significantly in the SSE group but increased in RPAE groups. The relative expression levels of MDRI mRNA in the SSE and RPAE combination groups were higher than that in the SSE group. The relative expression levels of MDRI mRNA in the SSE (1 μg/mL) + RPAE (3 μg/mL) and SSE
Table 1: Effect of RPAE on brucine transport in MDCK-MDR1 cells ($\bar{x} \pm SD, n = 3$).

<table>
<thead>
<tr>
<th>Groups</th>
<th>$\text{Papp}_{\text{AP-BL}} (\times 10^{-6} \text{ cm/s})$</th>
<th>$\text{Papp}_{\text{BL-AP}} (\times 10^{-6} \text{ cm/s})$</th>
<th>ER</th>
</tr>
</thead>
<tbody>
<tr>
<td>1 $\mu$g/mL SSE</td>
<td>6.81 $\pm$ 1.76</td>
<td>36.72 $\pm$ 0.23</td>
<td>5.39 $\pm$ 0.13</td>
</tr>
<tr>
<td>1 $\mu$g/mL SSE + 1 $\mu$g/mL RPAE</td>
<td>5.97 $\pm$ 0.58</td>
<td>33.82 $\pm$ 0.67</td>
<td>5.66 $\pm$ 1.16</td>
</tr>
<tr>
<td>1 $\mu$g/mL SSE + 3 $\mu$g/mL RPAE</td>
<td>4.63 $\pm$ 1.59*</td>
<td>34.65 $\pm$ 1.30</td>
<td>7.48 $\pm$ 0.82</td>
</tr>
<tr>
<td>1 $\mu$g/mL SSE + 6 $\mu$g/mL RPAE</td>
<td>4.46 $\pm$ 0.97*</td>
<td>48.23 $\pm$ 1.66*</td>
<td>10.81 $\pm$ 1.71</td>
</tr>
<tr>
<td>1 $\mu$g/mL SSE + 100 $\mu$M verapamil</td>
<td>24.98 $\pm$ 2.17*</td>
<td>28.78 $\pm$ 2.16*</td>
<td>1.15 $\pm$ 0.99</td>
</tr>
</tbody>
</table>

* $P < 0.05$ with contrast to the SSE $1 \mu$g/mL group; $^*$ $P < 0.05$ with contrast to groups without verapamil.

Table 2: Effect of RPAE on strychnine transport in MDCK-MDR1 cells ($\bar{x} \pm SD, n = 3$).

<table>
<thead>
<tr>
<th>Groups</th>
<th>$\text{Papp}_{\text{AP-BL}} (\times 10^{-6} \text{ cm/s})$</th>
<th>$\text{Papp}_{\text{BL-AP}} (\times 10^{-6} \text{ cm/s})$</th>
<th>ER</th>
</tr>
</thead>
<tbody>
<tr>
<td>1 $\mu$g/mL SSE</td>
<td>28.33 $\pm$ 2.67</td>
<td>39.39 $\pm$ 11.75</td>
<td>1.39 $\pm$ 4.40</td>
</tr>
<tr>
<td>1 $\mu$g/mL SSE + 1 $\mu$g/mL RPAE</td>
<td>24.49 $\pm$ 1.63</td>
<td>30.73 $\pm$ 0.64</td>
<td>1.25 $\pm$ 0.39</td>
</tr>
<tr>
<td>1 $\mu$g/mL SSE + 3 $\mu$g/mL RPAE</td>
<td>22.29 $\pm$ 8.19</td>
<td>37.39 $\pm$ 0.95</td>
<td>1.68 $\pm$ 0.12</td>
</tr>
<tr>
<td>1 $\mu$g/mL SSE + 6 $\mu$g/mL RPAE</td>
<td>19.64 $\pm$ 2.33*</td>
<td>41.92 $\pm$ 1.53*</td>
<td>2.13 $\pm$ 0.67</td>
</tr>
<tr>
<td>1 $\mu$g/mL SSE + 100 $\mu$M verapamil</td>
<td>44.19 $\pm$ 4.04*</td>
<td>39.57 $\pm$ 1.53*</td>
<td>0.90 $\pm$ 0.38</td>
</tr>
</tbody>
</table>

* $P < 0.05$ with contrast to the SSE $1 \mu$g/mL group; $^*$ $P < 0.05$ with contrast to groups without verapamil.

Table 3: Accumulation of Rh123 in MDCK-MDR1 cells ($\bar{x} \pm SD, n = 3$).

<table>
<thead>
<tr>
<th>Group</th>
<th>Fluorescence value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>111.95 $\pm$ 1.04</td>
</tr>
<tr>
<td>100 $\mu$M verapamil</td>
<td>1113.13 $\pm$ 56.80*</td>
</tr>
<tr>
<td>1 $\mu$g/mL SSE</td>
<td>46.14 $\pm$ 1.05*</td>
</tr>
<tr>
<td>1 $\mu$g/mL SSE + 1 $\mu$g/mL RPAE</td>
<td>30.37 $\pm$ 1.62**</td>
</tr>
<tr>
<td>1 $\mu$g/mL SSE + 3 $\mu$g/mL RPAE</td>
<td>26.07 $\pm$ 1.05**</td>
</tr>
<tr>
<td>1 $\mu$g/mL SSE + 6 $\mu$g/mL RPAE</td>
<td>24.77 $\pm$ 4.82**</td>
</tr>
<tr>
<td>1 $\mu$g/mL RPAE</td>
<td>26.94 $\pm$ 0.84**</td>
</tr>
<tr>
<td>3 $\mu$g/mL RPAE</td>
<td>25.70 $\pm$ 0.38**</td>
</tr>
<tr>
<td>6 $\mu$g/mL RPAE</td>
<td>19.66 $\pm$ 0.30**</td>
</tr>
</tbody>
</table>

* $P < 0.05$ with contrast to the control group; $^* P < 0.05$ with contrast to the SSE $1 \mu$g/mL group.

3.5. Effect of RPAE and/or SSE on P-gp Expression. As shown in Figure 4 and Table 4, in contrast to the control group, P-gp expression significantly increased in the SSE and RPAE combination groups and significantly decreased in the SSE group.

3.6. Effect of RPAE and/or SSE on P-gp ATPase Activity. As shown in Figure 5, the $\Delta$RLU of all groups was greater than $\Delta$RLU$_{basal}$, demonstrating that all of the test compounds are stimulators of P-gp ATPase activity. Because the $\Delta$RLU in the RPAE groups was greater than that in the SSE group, the affinity of RPAE for P-gp was greater than that of SSE.

4. Discussion

Semen Strychni can effectively be used to treat inflammation and pain, but its use is limited by its toxic effects. Some studies have reported that Radix Paeoniae Alba can reduce the toxic effects and increase the therapeutic effects of Semen Strychni [4–6]. In the research, the mechanism of P-gp modulation by Semen Strychni combined with Radix Paeoniae Alba was investigated. First, the cytotoxic effects of SSE, RPAE, and the combination of SSE and RPAE on MDCK-MDR1 cells were estimated by an MTT assay with verapamil as the positive control. We found that the nontoxic concentration range of SSE increased significantly when combined with RPAE.

To investigate the underlying mechanism, the effect of RPAE on brucine and strychnine transport in MDCK-MDR1 cells was investigated, and the results showed that RPAE could inhibit brucine and strychnine transport in MDCK-MDR1 cells which might explain how RPAE reduces the toxicity of SSE to the central nervous system. Our results are consistent
Table 4: The effect of RPAE and SSE on the relative expression level of P-gp was measured by Western blot.

<table>
<thead>
<tr>
<th>Relative protein quantification</th>
<th>Control</th>
<th>A</th>
<th>B</th>
<th>C</th>
<th>D</th>
</tr>
</thead>
<tbody>
<tr>
<td>P-gp</td>
<td>1.00</td>
<td>0.88</td>
<td>1.53</td>
<td>2.17</td>
<td>2.79</td>
</tr>
<tr>
<td>GAPDH</td>
<td>1.00</td>
<td>1.00</td>
<td>1.00</td>
<td>1.00</td>
<td>1.00</td>
</tr>
</tbody>
</table>

In Table 4, A is the SSE (1 μg/mL) group, B is the SSE (1 μg/mL) + RPAE (1 μg/mL) group, C is the SSE (1 μg/mL) + RPAE (3 μg/mL) group, and D is the SSE (1 μg/mL) + RPAE (6 μg/mL) group.

Figure 2: Effect of the combination of Semen Strychni and Radix Paeoniae Alba on P-gp function.
interact with P-gp. The effect of RPAE on the activity of ATPase activity, demonstrating that SSE and RPAE directly upregulate the expression of P-gp. Therefore, we speculated that SSE combined with RPAE could activate P-gp to increase the expression of P-gp.

Results indicated that the combination of SSE and RPAE could significantly increase the efflux of Rh123 in a proportion-dependent manner. Webb et al. [10] reported that P-gp was regulated by the combination of SSE with RPAE and found that Semen Strychni combined with Radix Paeoniae Alba might not be related to membrane lipid fluidity. This result indicated that the mechanism through which P-gp is regulated by Semen Strychni combined with Radix Paeoniae Alba might not be related to membrane lipid fluidity.

Finally, although P-gp is the most well-known active efflux transporter, there are many other multidrug transporters, such as members of the multidrug resistance protein family. Our study focused on P-gp, but future studies should be performed to investigate the other transporters.

5. Conclusions

In the current study, we focused on the mechanism through which P-gp was regulated by the combination of SSE with RPAE and found that Semen Strychni combined with Radix Paeoniae Alba could induce P-gp-mediated drug transport, likely by inhibiting brucine and strychnine transport in MDCK-MDR1 cells, enhancing the P-gp efflux function and stimulating P-gp ATPase activity. This investigation can not only improve the clinical therapeutic use of Semen Strychni but also might facilitate the exploration of possible preventative methods to decrease the toxic effects of Semen Strychni through its combination with Radix Paeoniae Alba.

Conflicts of Interest

The authors have stated that there are no conflicts of interest.
Authors’ Contributions
Li-Li Liu and Yong-Mei Guan contributed equally to this study and share first authorship.

Acknowledgments
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