



Full Length Article

Effects on Expression of Hepatic Phase-1 and Phase-2 Metabolism and Transporter Genes by *Swertia mussotii* Extract, A Hepatoprotective Herb from Tibetan Traditional Medicine

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Abstract

To study the effects of *Swertia mussotii* extract on the expression of drug processing genes, including Phase-1 (cytochrome P450 enzyme genes), Phase-2 (glucuronidation and sulfation genes) and Phase-3 (drug transporters) in mouse liver. Mice were orally administered *S. mussotii* extract at the dose of 0.1, 0.3, 1.0, 3.0 g/kg for consecutive eight days. Blood and livers were collected at the time of 24 h after dosing. RT-PCR was used to analysis of genes of interest. *S. mussotii* extract ingestion induced Cyp2b10 and inhibited Cyp3a11 at all doses, while increased the expressions of Cyp2e1 at the dose of 0.1 and 0.3 g/kg and decreased the expression gradually at the dose of 1.0 and 3.0 g/kg. For phase-2 enzyme genes, *S. mussotii* extract increased sulfotransferase (Sult1a1), heme oxygenase-1 (HO-1) and quinone oxidoreductase 1 (Nqo1), but had no effects on glucuronyltransferases (Ugt1a1 and Ugt1a6) and sulfotransferase (Sult1b1). It is very interesting that *S. mussotii* extract inhibited mRNA levels in the hepatic organic anion-transporting polypeptides (Oatp1a1 and Oatp2b1) and induced multidrug resistance-associated proteins, such as Mrp1 and Mrp4 were especially induced at 3.0 mg/kg. The metabolism of *S. mussotii* extract includes the induction of cytochrome P450 enzyme genes, hepatic transporters and phase-2 enzyme genes. The *S. mussotii* extracts influenced all types of metabolic drug processing genes and therefore could play a role in producing herb-drug interactions. © 2019 Friends Science Publishers

Keywords: Tibetan traditional medicine; *Swertia mussotii*; CytochromeP450; Hepatic transporters

Introduction

Herbal medications have been broadly utilized for centuries, and they are still used by 75–80% of the global population based on World Health Organization (WHO) statistics (Sieniawska *et al.*, 2013). And with the progress, there would up to 7000 chemical compounds used in phytotherapy are derived from herbal medicines (Sieniawska *et al.*, 2013). Because herbal medications have multiple pharmacologically active ingredients, more complex pathways are involved in absorption, distribution, metabolism, and excretion (*e.g.*, activating or suppressing metabolic enzymes).

The Gentianaceae herb *Swertia mussotii* Franch has hepatoprotective (Hase *et al.*, 1999; Karan *et al.*, 1999; Li *et al.*, 2005; Sadeghi *et al.*, 2006; Kotoky and Das, 2008), anticancer (Saha *et al.*, 2004; Balasundari *et al.*, 2005), antibacterial (Ramesh *et al.*, 2002) and antioxidant properties (Scartezzini and Speroni, 2000; Tripathi *et al.*, 2007) because of the existence of many therapeutically active phytoconstituents, including flavonoids, xanthenes,

terpenoids, secoiridoid, iridoids, swertiamarin, glycosides, and ursolic acid (Yamahara *et al.*, 1978; Kikuzaki *et al.*, 1996; Brahmachari *et al.*, 2004). It is used extensively in the Tibetan traditional system of medicine as a hepatoprotective agent since it is easily accessible to local communities. As rare wild plants of the Tibetan plateau, studies of the medicinal ingredients of *S. mussotii* are constantly deepening with the rapid development of national medicine. Active components, such as swertiamarin, sweroside, gentiopicroside and isoorientin, can be extracted from *S. mussotii* efficiently. The increase in the amount of pure components has allowed a better understanding of their pharmacological activity, which has shown distinct activities for some of the constituents of *S. mussotii*. For instance, sweroside inhibits the central nervous system (Zheng and Liu, 2009), while mangiferin stimulates the central nervous system (Cheng *et al.*, 1999), different doses of oleanolic acid have hepatoprotective effects or lead to cholestasis (hepatotoxicity), which are opposite effects (Liu, 2005; Lu *et al.*, 2013). The opposing activities of these compounds implies that different extraction methods may lead to the

production of different efficacies and demonstrate the diversity of medication efficacies that can be obtained when the whole plant is used. Our Laboratory has found that the median lethal dose (LD₅₀) of the 75% ethanol extract of *S. mussotii* is 27.24 g/kg, whereas the maximum tolerated dose (MTD) of the water extract of *S. mussotii* is 40 g/kg (Lv *et al.*, 2010). Because of these different mechanisms underlying the activity of medicinal ingredients, the pharmacological effect of *S. mussotii* extract remains a concern.

Drugs and other xenobiotics normally experience two metabolizing phases in the body, which are called phase 1 and 2 metabolism. During the phase 1 metabolism, drugs are catalyzed or detoxified mainly by Cytochrome P450 (CYP-450). Suppressing CYP-450 enzymes can cause elevated plasma levels due to a concomitantly dispensed pharmaceutical as the enzymes play a vital role in therapeutic pharmacology. The toxicology of xenobiotics could lower or extend the pharmacological power, broaden the manifestation of drug-induced toxicity, or reduce the therapeutic effect (Jarukamjorn *et al.*, 2006; Lee *et al.*, 2013). In the phase 2 metabolism, enzymes such as glucuronyltransferase (Ugts), sulfotransferase (Sults), heme oxygenase-1 (HO-1) and quinone oxidoreductase 1 (Nqo1), and some transporters all participating in metabolism and affect the disposition of chemicals in the body (Liu, 2007; Klaassen and Aleksunes, 2010). However, almost no reports about effects of *S. mussotii* extract on metabolism enzymes and liver drug transporters. Therefore, this study was designed to evaluate the transporter-mediated hepatobiliary disposition of *S. mussotii* extract to examine the effects of *S. mussotii* extract on major phase-1 CYP and phase 2 enzyme genes. To obtain a systemic understanding of the metabolism of the *S. mussotii* extract, the efflux transporters of multidrug resistance associated protein genes (primarily Mrp1, Mrp2 and Mrp4) and 4 major hepatic organic anion-transporting polypeptides (namely Oatp1a1, Oatp1b2, Oatp1a4 and Oatp2b1) were all assessed.

Materials and Methods

Reagent and Animals

Gentiopicroside (the purity of 99.9%), purchased from National Institute for Food and Drug Control (Beijing, China). All Kunming (KM) mice (*Mus musculus*) were eight-week-old, weighing 18–22 g, and were purchased from the Experimental Animal Center of Gansu University of Chinese Medicine (Gansu, China; Certificate No. SCXK 2015-0005), then kept in animal facilities at Northwest Institute of Plateau Biology, Chinese Academy of Sciences (NWIPB,CAS), with standard laboratory conditions of regular environment (12 h light /dark cycle, 24±1°C and 45 ± 2% humidity). All animals were acclimatized for a period of 15 days prior to the start of the experiment. All animal procedures were conducted in accordance with the WHO Guidance of Humane Care and Use of Laboratory Animals.

Plant Material and Preparation of Extract

Whole plants (include root) of *S. mussotii* were collected from Sichuan province, China (altitude, 2,729 m; latitude, 31°27.575'; longitude, 102°05.600'; slope, 10°; aspect, northeast), at the full-blooming stage in July 2015. Whole plants of *S. mussotii* were dried under shade. Air-dried *S. mussotii* plants were reduced to coarse powder, extracted with 75% ethanol four times (60 min each time) with a Soxhlet apparatus, and then concentrated in the midst of lowered temperature (45°C) and pressure in a rotary evaporator (Heidolph, Germany). The extracts were freeze-dried, then maintained in desiccators.

Standardization of Gentiopicroside through HPLC Method

The lyophilized extract of *S. mussotii* was standardized by using HPLC. Gentiopicroside was used as the standard. Agilent HPLC system equipped with G1379A degasser, G1316A column heater, G1312A pump, G1315B DAD Detector and manual injector. The column used for HPLC analysis was an Agilent Eclipse XDB-C18 column (Agilent Spherisorb, USA) with a length 250 mm × width 4.6 mm, and the column temperature set at 25°C.

Typical stock solutions of gentiopicroside and *S. mussotii* extract (1 mg/mL each) were set up in HPLC-grade methanol. The stock and extract solutions were sieved through a syringe filter. The loading volume was 10 µL. Elution was carried out with optimized mobile phase containing acetic acid (0.1%): methanol (0 min, 75:25 v/v; 17 min, 60:40 v/v; 18 min, 50:50 v/v; 30 min, 30:70 v/v; 35 min, 10:90 v/v; 40 min, 75:25 v/v) at flow rate of 1 mL/min and at wavelength of 254 nm. Prior to usage, the mobile phase was sonicated for 10 min and then degassed. The analysis was performed for a 40 min run time to take out the analyte.

The percent relative standard deviation (RSD), limit of detection (LOD) and quantification (LOQ) and the percent recovery of gentiopicroside from *S. mussotii* extract were all carried out in triplicate (n=3) and as per the recommended guideline of International Conference on Harmonization (ICH). The standard solutions were diluted to different concentration levels with methanol and were analyzed; the different concentration levels and peak area could be used to generate a calibration curve of the standard compound.

Experimental Design

All animals were randomly divided into six groups after adapted for five days, as the solvent control group (distilled water), first group (0.1 g/kg), second group (0.3 g/kg), third group (1.0 g/kg) and fourth group (3.0 g/kg) of *S. mussotii* extract, 20 mL/kg intragastric gavage (i.g.) daily, for eight days continuously. All doses were chosen according to our previous studies (Yang *et al.*, 2016). Mice were monitored

chiefly for animal activities and the body weight, during the 8 day dosing period. Blood and livers were collected at 24 h after dosing. Serum was separated and alanine aminotransferase (ALT) was analyzed with automatic blood analyzer in Northwest Institute of Plateau Biology, CAS. After recorded liver weights, part of the liver was stained with hematoxylin and eosin (H&E) after fixed in 10% neutral formalin for 48 h to examine morphology of the liver and approximately 100 mg liver was promptly put into 1.0 mL TRIzol (TakaRa Biotechnology, Dalian, China) solution for RNA isolation.

Real-time PCR Analysis

Isolated RNA was determined by the 260/280 ratios and gel-electrophoresis to analysis quality and quantity. Then the RNA was reversed transcribed with the High Capacity Reverse Transcriptase Kit (Applied Biosystems, Foster City, CA, USA). Primers were designed and listed in Table 1. Total RNA was extracted according to the manufacturer's instructions, followed by purification with Total RNA (Mini) Kit (Watson Biotechnology, Shanghai, China). RT-PCR analysis was performed with Power SYBR Green Master Mix (Applied Biosystems, Foster City, CA, USA). The procedure included 5 min of denaturing at 95°C, 40 cycles of annealing and extension for 45 s at 60°C. This was followed by denaturing at 95°C for 10 s. The dissociation curve was formed after completing 40 cycles to confirm the quality of amplification and the primers. The relative expression of the genes was determined with the $2^{-\Delta\Delta Ct}$ method. Normalization was done with the housekeeping gene β -actin using an identical sample.

Statistical Analysis

Statistical analysis was done by using SPSS 21.0 software. The data were expressed as mean \pm SEM. All data were analyzed by the one-way analysis of variance (ANOVA), followed by LSD multiple range test or Dunnett's Test. p -value < 0.05 was considered statistically significant.

Results

Quantification of Gentiopicroside by HPLC

Standard gentiopicroside presented a good linear relation in the range of 14 to 1000 $\mu\text{g/mL}$ in the calibration curve. The correlation between peak area and concentration was found to be good and had the higher correlation coefficient (r^2) value of 0.999. RSD was used to approximate the precision of instrument. The RSD values of instrumental precision and intra-assay precision of gentiopicroside were approximated as $< 1.8\%$. The LOD and LOQ of gentiopicroside were determined to be 0.064 and 0.58 $\mu\text{g/mL}$, respectively. This suggested that the HPLC procedure for gentiopicroside was exact and the sensitivity of the technique was good enough to identify the gentiopicroside in the extract. At various

concentrations, the retrieval of gentiopicroside was found to be 98% with RSD values that were $< 1.5\%$, suggesting that gentiopicroside can be fully retrieved from the *S. mussotii* extract. Gentiopicroside was determined and quantified in the *S. mussotii* extract through a comparison of the retention time ($R_t = 15.19$ min) of typical gentiopicroside. These outcomes were deemed satisfactory and appropriate for the quantitative analysis. Gentiopicroside present in *S. mussotii* extract was found to be 27.48% (w/w).

Animal General Health

Mice received oral administrations of distilled water and various amounts of *S. mussotii* extract (SME) (0.1, 0.3, 1.0, and 3.0 g/kg) for eight days. The body weight, movements, and overall health of the mice were noted. Body weight gain in various *S. mussotii* extract groups was obviously different compared to the controls, but all mice displayed typical movements and overall health. In comparison, with the doses increase, the animal body weight decreases significantly (23.35 ± 0.32 , 22.33 ± 0.42 , 21.77 ± 0.29 , 21.83 ± 0.06 and 21.35 ± 0.34 g for control, SME 0.1, 0.3, 1.0 and 3.0 g/kg group, respectively), especially the SME 0.3, 1.0 and 3.0 g/kg groups, were significantly different to control group ($P < 0.05$). At necropsy, the ratios of liver/body weight were not different from the control (5.67 ± 0.18 , 5.25 ± 0.24 , 5.06 ± 0.07 , 6.07 ± 0.19 and $5.50 \pm 0.16\%$ for control, SME 0.1, 0.3, 1.0, and 3.0 g/kg group, respectively). Blood biochemistry results indicated that ALT (45.01 ± 3.58 , 44.98 ± 4.14 , 44.61 ± 4.07 , 41.48 ± 3.72 and 40.39 ± 3.56 U/L for control, SME 0.1, 0.3, 1.0 and 3.0 g/kg group, respectively) were not observably different among groups.

Effect of *S. mussotii* Extract on Cytochrome P450 Enzymes

To verify the impact of *S. mussotii* extracts on P450 enzyme gene, we determined the mRNA levels of major P450 for metabolism (Table 2). The results showed the expression of Cyp2b10 was increased with the dose increase, while Cyp2e1 and Cyp3a11 were inhibition significantly at the highest doses (SME 3.0), which showed statistical difference when compare to the control group. The expression of Cyp2e1 increased at doses of 0.1 and 0.3 g/kg. It gradually decreased at doses of 1.0 and 3.0 g/kg. *S. mussotii* extract disclosed numerous induction activities toward CYP-450.

Effect of *S. mussotii* Extract on Phase 2 Enzyme Genes

Induction of phase-2 enzymes is broadly acknowledged as an effective method to lower the risk of various diseases associated with exposure to toxins, carcinogens and mutagens (Talalay *et al.*, 1995; Wilkinson and Clapper, 1997; Talalay, 2000; Zhang and Gordon, 2004). Therefore, the effect of *S. mussotii* extracts on Sults, Ugt, HO-1 and Nqo1 mRNA levels in this study were examined.

Table 1: Primer sequences for RT-PCR analysis

Gene	Accession	Forward	Reverse
β-actin	V01217	TGACCGAGCGTGGCTACAG	GGGCAACATAGCACAGCTTCT
Cyp1a2	NM_009993	GACATGGCCTAACGTGCAG	GGTCAGAAAAGCCGTGGTTG
Cyp2b10	AF128849	AAGGAGAAGTCCAACCAGCA	CTCTGCAACATGGGGTACT
Cyp2e1	NM_021282	TCCCTAAGTATCCTCCGTGA	GTAATCGAAGCGTTTGTGA
Cyp3a11	NM_007818	ACAAACAAGCAGGGATGGAC	CCCATATCGGTAGAGGAGCA
Cyp4a10	NM_010011	CACACCTGATCACCAACAG	TCCTTGATGCACATGTGGT
Sult1a1	NM_133670	GGATGTAGCTGAGGCAGAGG	CAGCTCCAGTGGCATTAT
Sult1b1	NM_019878	GGTGGGAAAAGAGGGAAGAG	AAGGCCTTTCATCCAAGGT
Ugt1a1	NM_201645	ACACCGAACTAGACCATCG	ATACCATGGGAGCCAGAGTG
Ugt1a6	NM_145079	ATACCATGGGAGCCAGAGTG	ACCAGAACTGTGAGGGTTGG
HO-1	M33203	CCTCACTGGCAGGAAATCATC	CCTCGTGAGACGCTTACATA
Nqo 1	BC004579	TATCCTTCCGAGTCATCTAGCA	TCTGCAGCTTCCAGCTTCTTG
Oatp1a1	NM_013797	ATCCAGTGTGTGGGACAAT	GCAGCTGCAATTTTGAAAACA
Oatp1a4	NM_030687	GGAAGATTGGACACGCATCT	GGCATTGTGACTGAAGCAGA
Oatp1b2	NM_020495	CAAACCTGCATCCAAGCAA	GGCTGCCAAAAAATATCCCTGA
Oatp2b1	NM_020495	CTAGGCCAAATGCCAGAAAG	TTGCTTGGATGCTGAGTTTG
Mrp1	NM_008576	GCCCTCTTTGCAATCATCTC	CAGTCTCTCCACTGCCACAA
Mrp2	NM_013806	TCCTAGACAGCGGCAAGATT	GCTAGAGCTCCGTGTGGTTC
Mrp4	BC150822	GCAAAGCCCATGTACCATCT	ACCACGGTAACAACCTACC

Table 2: Effect of *S. mussoitii* extract on mouse liver P450 gene expression

	control	SME 0.1	SME 0.3	SME 1.0	SME 3.0
Cyp1a2	47.728±1.774	49.306±4.452	46.714±0.862	48.970±4.191	45.806±7.416
Cyp2b10	4.394±0.851	4.516±0.738	5.890±1.319	7.094±1.915	8.537±0.841*
Cyp2e1	494.120±14.863	849.859±78.910*	760.111±24.777	545.213±69.469	351.232±9.561*
Cyp3a11	363.198±46.642	260.480±14.369	231.055±42.871	237.883±27.602	161.914±25.259*
Cyp4a10	0.583±0.196	0.297±0.066	0.942±0.071	1.083±0.034	1.244±0.339

**P* < 0.05 vs control

The expression of HO-1 and Nqo1 were increased 2-fold at highest doses of *S. mussoitii* extract and Sult1a1 was also increased, while it had no obvious effects on Sult1b1, Ugt1a1 and Ugt1a6 (Fig. 1).

Effect of *S. mussoitii* Extract on Hepatic Uptake Transporters

The initial step of xenobiotic metabolism elevates their uptake and every one of the similar treatments that can raise hepatic uptake could have a major effect on drug metabolism. As noted in the literature, in the liver, organic anion transporting peptides (Oatps) are the primary transporters (Cheng *et al.*, 2005). In this study, we choose Oatp1a1, Oatp1a4, Oatp1b2 and Oatp2b1 to observe. But Fig. 2 clearly demonstrates that *S. mussoitii* extract decreased the mRNA levels of Oatp1a1 and Oatp1b2. The change of expression of Oatp2b1 and Oatp1a4 was not evident.

Effect of *S. mussoitii* Extract on Hepatic Efflux Transporters

As the prime hepatic efflux transporters (Klaassen and Aleksunes, 2010), multidrug resistance proteins (Mrps) play a crucial role in the transfer of drug metabolites and drugs from the liver. *S. mussoitii* extract had remarkable effect on Mrp1, Mrp2 and Mrp4 mRNA levels, especially at the dose of 3.0 g/kg for Mrp1 and Mrp4 (Fig. 3).

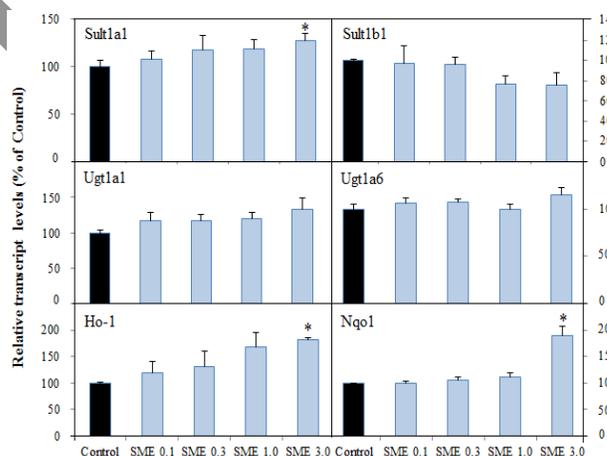


Fig. 1: Outcome of *S. mussoitii* extract on the expression of phase 2 enzyme genes in adult KM mouse livers. RT-PCR analysis the total RNA, which from the mice that were given *S. mussoitii* extract orally at doses of 0.1 g/kg, 0.3 g/kg, 1.0 g/kg and 3.0 g/kg for 8 days. The data represents the mean ± SEM. (*P* < 0.05 vs control)

Histopathological Alterations

Characteristics H & E microphotos are given in Fig. 4. The livers of the mice that were orally given *S. mussoitii* extract (SME 0.1, 0.3, 1.0 and 3.0 g/kg) for eight days were comparable to the controls. This indicates that no

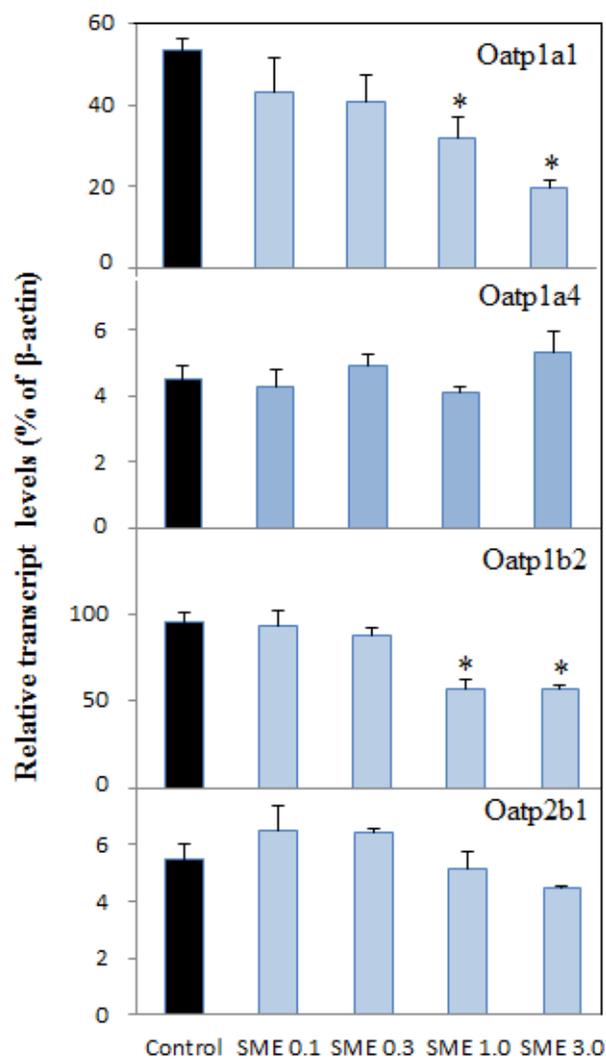


Fig. 2: Outcome of *S. mussotii* extract on the expression of uptake transporters in adult KM mouse livers. RT-PCR analysis the total RNA, which from the mice that were given *S. mussotii* extract orally at doses of 0.1 g/kg, 0.3 g/kg, 1.0 g/kg and 3.0 g/kg for 8 days. The data represents the mean \pm SEM. ($P < 0.05$ vs control)

obvious liver pathology lesions were visible with the dosage of the *S. mussotii* extract utilized in the current evaluation.

Discussion

Using herbs from natural sources as therapeutics has grown quickly (Malviya *et al.*, 2010). The current evaluation clearly shows that *S. mussotii* extract had a great impact on hepatic drug processing enzyme gene expression. It initiated CYP enzyme genes, had an impact on Sults and Ugts expression and elevated hepatic uptake and efflux transporter expression. Every one of these effects could have an integral part, and the outcomes from this systemic evaluation of the impact of *S. mussotii* extract on liver drug processing genes is pivotal

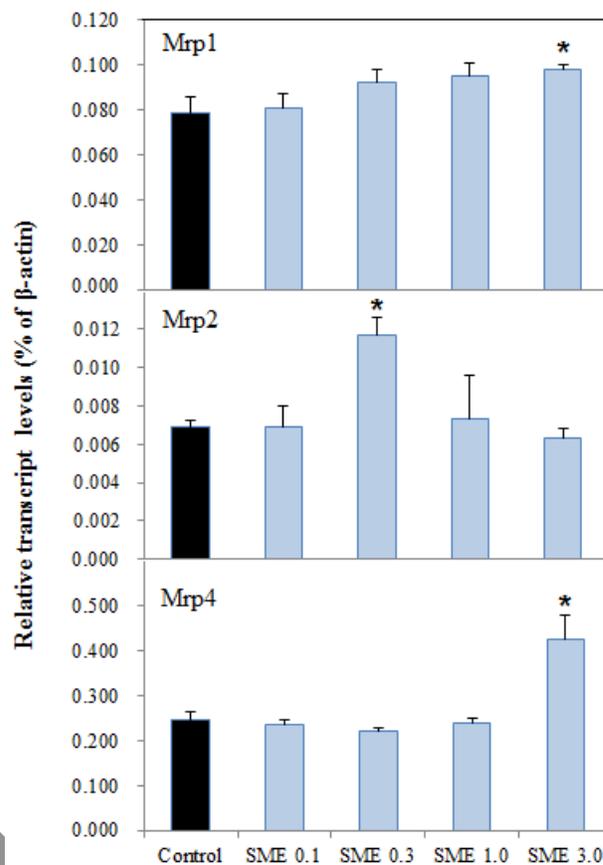


Fig. 3: Outcome of *S. mussotii* extract on the expression of efflux transporters in adult KM mouse livers. RT-PCR analysis the total RNA, which from the mice that were given *S. mussotii* extract orally at doses of 0.1 g/kg, 0.3 g/kg, 1.0 g/kg and 3.0 g/kg for 8 days. The data represents the mean \pm SEM. ($P < 0.05$ vs control)

when quantifying the specific bioactive molecule in the complex herbal extract.

S. mussotii extract has many substantial impacts on hepatic CYP450s. The hepatic CYP450 1 to 4 family gene expression levels we selected in the evaluation changed after the mice received *S. mussotii* extract for eight successive days. Cyp2b10 expression was elevated 2-fold at the largest dosage and Cyp3a11 expression was suppressed at all of the doses. This reinforced that the activation of Cyp genes is essential herb–drug interactions mechanization (Guengerich, 1997; Su *et al.*, 2007; Choi *et al.*, 2011). Besides the effects of *S. mussotii* extract on cytochrome P450s have been observed, the effects of extract on Phase 2 metabolism also have been studied for their importation for herb–drug interactions (Zamek-Gliszczynski *et al.*, 2006). The outcomes of this evaluation obviously showed that *S. mussotii* extract has a specific impact on the mRNA levels of the primary enzyme genes for sulfation conjugation-Sult1a1 at the largest dosage of 3 g/kg. In contrast, effect of *S. mussotii* extract on Ugt1a1 and Ugt1a6 was not obvious, and there was no relevant study to support our results.

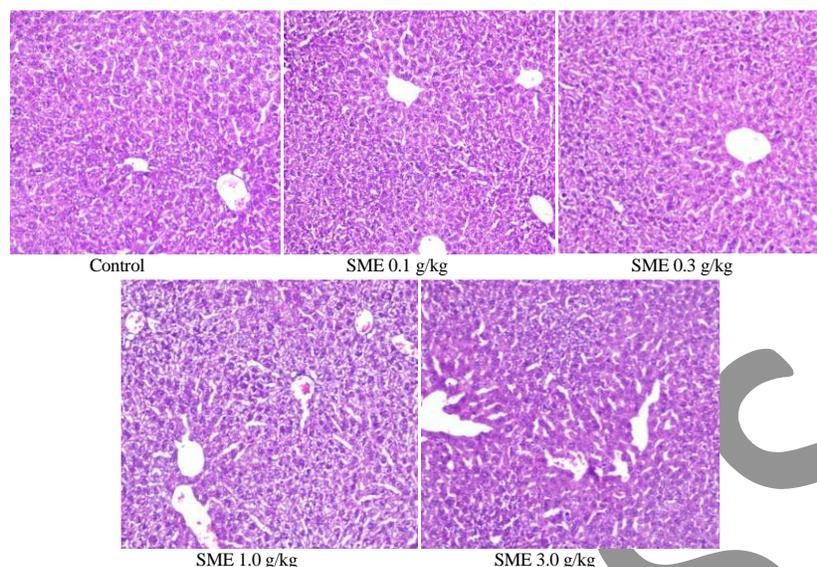


Fig. 4: Histopathological examination with representative photos from livers of mice orally administrated with distilled water (Control, 20 mL/kg) and *S. mussotii* extracts (SME 0.1, 0.3, 1.0 and 3.0 g/kg) for 8 days. Magnification: 200X

The Nrf2/ARE antioxidant pathways are primary defense mechanisms against oxidative stress (Liu *et al.*, 2013), which causes cells to restore toxic stimuli. Nrf2 separates from its partner, Keap1 and then it initiates a group of cell defense genes, including Nqo1, one of the primary Nrf2 target genes (Itoh *et al.*, 1997; Ishii *et al.*, 2002). In this study, highest dose of *S. mussotii* extract increased the expression of Nqo1 2-fold, which imply the potential liver injury sides effects, meanwhile, the expression of Ho-1 was changed in livers of mice that were given orally *S. mussotii* extract (0.3 g/kg) indicative of the oxidative stress to the liver. As a sensitive biomarker for oxidative liver injury (Lu *et al.*, 2011), the level change of Nqo1 and Ho-1 all indicated the further studies are necessary to expound the effects.

Organic anion-transporting polypeptides are related to drug-drug interactions (Koenen *et al.*, 2011; Badolo *et al.*, 2013), for they are responsible for cellular influx of some xenobiotics and endogenous substances (Obaidat *et al.*, 2012; Gong and Kim, 2013). As a kind of membrane uptake transporters, they are known to play a crucial part in transporting substrates from the blood in to the liver (Evers and Chu, 2008). Oatp1a1 is important in liver bile acid homeostasis and mainly expressed in liver and kidney (Cheng *et al.*, 2005; Fu *et al.*, 2012; Zhang *et al.*, 2012). The primary hepatic uptake transporter Oatp1b2 (Lu *et al.*, 2008) plays crucial roles in the hepatic uptake of therapeutics, environmental chemicals and toxicants (Meier-Abt *et al.*, 2004), as well as internal materials, such as bile acids (Csanaky *et al.*, 2011). In the present study, Oatp1a1 declined with an increase in dose, and similar decrease of Oatp1b2 was also evident, with over 40% decrease in SME 3 g/kg group compared to control group and the expression of Oatp1b2 was decreased at all doses of *S. mussotii* extract groups.

These results distinctly proved the effect of *S. mussotii* extract on Oatp1a1, Oatp1a4, Oatp1b2 and Oatp2b1 and the effect on the hepatic uptake or ganicanion transporting peptides could be regarded as a basis of *S. mussotii* extracts - drug interactions.

ATP-dependent plasma membrane transporters named Mrps, are in charge of the removal of xenobiotics (Klaassen and Aleksunes, 2010). In the present study, we have examined the hepatic efflux transporters, Mrp1, Mrp2 and Mrp4, they are multidrug resistance-associated protein genes. As showed in Fig. 3, *S. mussotii* extract had prominent effects on the Mrp1 and Mrp4 mRNA levels, especially at dose of 3.0 g/kg. MRP1 has been demonstrated to extrude natural product drugs, *S. mussotii* extract as a natural product drugs, maybe Mrp1 extrude it in a cotransport for the result of its high expression level (Borst *et al.*, 1999; Kool *et al.*, 1999). Mrp4 has been considered as a detoxification protein though the normal expression of Mrp4 in the liver is low (Chen and Klaassen, 2004; Russel *et al.*, 2008), it is also called a versatile efflux transporter for drugs and toxicants (Borst *et al.*, 2007) by transport of structurally diverse endogenous and xenobiotic compounds (Russel *et al.*, 2008). The results of our study provided a clue that high dose of *S. mussotii* extract may have a side effect. Mrp2 is in charge of biliary removal, the flow of compounds into bile, and detoxification of internal and xenobiotic organic anions (Anne and Dietrich, 2007; Klaassen and Aleksunes, 2010). Mrp2 is part of the multidrug resistance protein subfamily restricted solely to the apical membrane domain of hepatocytes, and it is a crucial participant in herb-drug interactions. In this study, effect of *S. mussotii* extract on Mrp2 had no dose-effect relationship and it is contrary to the results of Mrp1 and Mrp4.

Conclusion

This study systemically substantiated that multidrug resistance-associated protein gene expressions and hepatic organic anion-transporting polypeptides may be stimulated by *S. mussotii* extract. The metabolism of *S. mussotii* extract requires activation of hepatic transporters, cytochrome P450 enzyme genes, and phase-2 enzyme genes. The impact of *S. mussotii* extract on drug processing genes may have an integral part in generating herb-drug interactions. Besides these, the results implied that traditional herb medicine, especially those based on experience and custom, may have potential side effects, which require our more attention and in-depth study.

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