

Clinical Evidence of Herbal Drugs As Perpetrators of Pharmacokinetic Drug Interactions

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- herbal drugs
- botanical drugs
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- pharmacokinetics

Abstract



The use of herbal/botanical products, also referred to as complementary and alternative medicines (CAM), worldwide enjoys increasing popularity. It appears in particular highly prevalent in patient populations already exposed to complex treatment algorithms and polypharmacotherapy, frequently involving narrow therapeutic index drugs. Accordingly, the potential clinical dimension and relevance of herb-drug interactions has received considerable attention over the last years. However, review of pertinent literature indicates that the available clinical evidence in this regard is still limited and sometimes inconclusive. Also, communication of herb-drug interaction data in the biopharmaceutical/medical literature is often complex and confusing, not always unbiased, and in many cases appears not to strive for clear-cut and useful guidance in terms of the clinical relevance of such findings.

This systematic review summarizes and interprets the published evidence on clinical herb-drug interaction studies which examined the potential of six popular herbal drugs (*Echinacea*, garlic, ginkgo, ginseng, goldenseal, and milk thistle) as perpetrators of pharmacokinetic (PK) drug interactions. Reported effect sizes were systematically categorized according to FDA drug interaction guideline criteria. A total of 66 clinical PK interaction studies, meeting the scope of the present review, were identified. The clinical evidence was found to be most robust and informative for *Ginkgo biloba* (GB; 21 studies) and milk thistle/silymarin (MT; 13), and appears still limited for ginseng (9), goldenseal/berberine (GS; 8), garlic (8), and *Echinacea* (7). Collectively, the available evidence indicates that, at commonly recommended doses, none of these herbs act as potent or moderate inhibitors or inducers of cytochrome P450 (CYP) enzymes or P-glycoprotein (ABCB1). Weak effects in terms of either induction

or inhibition were found for GB (presystemic/hepatic CYP3A4 induction/inhibition, CYP2C19 induction at high doses), milk thistle/silymarin (CYP2C9 inhibition), GS/berberine (CYP3A4 and CYP2D6 inhibition), *Echinacea* (presystemic/hepatic CYP3A4 inhibition/induction, CYP1A2 and CYP2C9 inhibition at high doses). Information was found not always complete for the major drug metabolizing CYP enzymes in the less well-studied herbs and is largely limited to P-glycoprotein (ABCB1) when effects on drug transporters have been investigated.

Abbreviations



1-OH-MDZ:	1'hydroxy midazolam
6-OHCZX:	6-hydroxy chlorzoxazone
ABCB1:	ATP binding cassette transporter B1 (P-glycoprotein)
BBR:	berberine
b.i.d.:	twice daily
BP:	botanical products
CAM:	complementary and alternative medicine
CAR:	constitutive androstane receptor
CL/F:	apparent oral clearance
CYP:	cytochrome P450 enzyme
CZX:	chlorzoxazone
DDI:	drug-drug interaction
EH:	epoxide hydrolase
EM:	extensive metabolizer
GST:	glutathione S-transferase
IR:	immediate release
MDZ:	midazolam
NHP:	natural health product
Nrf2:	E2-related factor 2
OSC:	organo sulfur compounds
PD:	pharmacodynamics
P-gp:	P-glycoprotein (ABCB1)
PK:	pharmacokinetics
PM:	poor metabolizer

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p.o.: per oral
 PXR: pregnane X receptor
 q.d.: once daily

QR: quinine reductase
 t.i.d.: three times daily
 UGT: uridine diphosphate (UDP)-glucuronosyltransferase

Introduction

Historically, plants represent the origin of modern pharmacotherapy. Herbal or botanical drug products are complex mixtures of organic chemicals that are derived from raw or processed parts of plants, such as seeds, roots, bark, leaves, flowers or fruits, or mixtures thereof [1]. Many herbal products have been used for thousands of years, e.g., in traditional herbal medicine across Asia, Australia, South America. Likewise, herbal products enjoy increasing popularity as an essential part of the trend towards complementary and alternative medicine (CAM) practices in Europe and North America. The use of herbal drugs varies by regions and patient populations: In Western countries it was estimated to occur with a prevalence of 12.1–18.6% in the United States [2] and 9.0–23.2% in Canada [3] among the general adult population [4]. The concomitant intake of herbal drugs has been identified to occur in 16% of prescription drug users in the United States [5], whereas in Canada the combined use of herbal and prescription drugs was reported to be lower at 5.3% [4]. Many consumers in these countries perceive herbal drugs as natural and therefore as generally safe. However, the naturalness of herbal drugs does not guarantee that they are generally safe, free from adverse effects or toxicity, and devoid of any drug-drug interaction potential. Furthermore, herbal drugs represent a mixture of organic compounds that are cleared by the same system of xenobiotic metabolism and transport that eliminate synthetic drugs from the human body. Therefore, from a mechanistic perspective, pharmacokinetic (PK) herb-drug interactions appear inevitable and comparable with those known for drug-drug interactions with synthetic pharmaceutical products.

In today's drug development process of synthetic drugs, the assessment of drug-drug interactions comprises a systematic combination of *in vitro* and *in vivo* investigations aiming at identifying possible interactions with *in vitro* screens which are then followed up by dedicated *in vivo* studies. Thereby the role of *in vitro* investigations and perhaps nonclinical PK interaction studies is only to inform the clinical drug-drug interaction program, not to serve as substitutes for clinical drug-drug interaction studies. Nonclinical studies in animals are generally discouraged due to the marked species differences, which renders extrapolation of such results to humans difficult [6]. Extrapolation of *in vitro* data to the *in vivo* situation by physiologically-based pharmacokinetic modelling (PBPK), which is being used to guide *in vivo* drug-drug interaction trials and drug labelling, is difficult to apply to herbal drugs because it requires I) the identification of the herbal component(s) acting as a perpetrator of the herb-drug interaction, II) content uniformity of these components within and across marketed products, III) well designed *in vitro* studies characterizing the DDI potential, and IV) most importantly, knowledge about the systemic bioavailability and human PK of this (these) component(s). Information on PK characteristics of herbal components, however, is often lacking with only a few exceptions. Therefore, it is not surprising that the extrapolation from *in vitro* herb-drug interaction data, in most cases, did not correctly predict the outcome of the corresponding *in vivo* study, except for a few examples such as berberine (BBR), an active component of goldenseal that was shown to be a mechanism-based inhibitor of CYP3A4 *in*

vitro [7], which actually translated into a weak inhibition of CYP3A4 activity *in vivo* by marketed goldenseal and BBR products in various independent studies [8–10].

Review of pertinent literature indicates that the communication of herb-drug interaction issues in the biopharmaceutical/medical literature is often confusing, hardly providing useful guidance in terms of the clinical relevance of such interactions. Therefore, the issue still represents a considerable challenge for the health care professionals and a potential safety concern for patients and/or consumers.

The aim of this article is therefore to provide a systematic review of the publicly available clinical evidence of herbal drugs as perpetrators of PK drug interactions. The review focuses on detailed information gained from mechanistic herb-drug interaction studies and aims to carefully consider trial design aspects, employed products, dosages, and treatment durations in the analysis and interpretation of the data. Thereby, it was one of the key objectives of the review to guide the reader through the interpretation of the data and to provide clear assessments of reported interactions in terms of their generalizability and clinical significance. Case reports were not considered to provide robust and reliable evidence of herb-drug interactions and were therefore excluded from this review. Pharmacodynamic herb-drug interactions including overlapping toxicities, although not less important, are beyond the scope of this work but have been reviewed elsewhere [11, 12]. The present review is focused on *Echinacea*, garlic, ginkgo, ginseng, goldenseal, and milk thistle ranking among the most frequently used herbal drugs in the United States [13], Canada [4], and Europe [12]. St. John's Wort is not addressed because of its well-documented properties as a perpetrator of PK-based drug interactions.

Methods

The University of Washington Metabolism and Drug Transporter Database (DIDB) [14], SCOPUS, as well as PubMed were used to screen for 6 herb-drug interactions identifying a total of 66 clinical interaction studies. The following numbers of studies were found for the individual herbs and fruit juices: *Echinacea* (7), garlic (8), ginkgo (21), ginseng (9), goldenseal (8), and milk thistle (13).

Evaluation of Herb-Drug Interactions in Humans

Ginkgo

Ginkgo (*Ginkgo biloba*; family Ginkgoaceae), also known as maidenhair tree, is a 190 million years old, unique species of tree with no close living relatives [15]. Pharmaceutical *G. biloba* (GB) products represent leaf extracts and contain flavonoid glycosides (mainly quercetin, kaempferol, isorhamnetin) and terpene lactones (ginkgolides A, B, C, J, and bilobalides). GB products rank among the most widely used herbal products in the United States. Oral standardized dry extracts of GB usually contain between 22–27% flavones glycosides, 5–7% terpene lactones, and should not exceed more than 5 ppm of ginkgolic acids, constitu-

ents with known allergic potency [16]. Many studies have been conducted using EGB 761, a well-defined extract of GB.

GB is believed to have nootropic properties and is mainly used as a memory and concentration enhancer and for the treatment of cognitive impairment and dementia of various origins (e.g., Alzheimer's disease). GB products are also advocated for use in peripheral arterial disease (i.e., intermittent claudication) and to improve microcirculation in disease states such as tinnitus. GB supplements are usually taken in the range of 40 to 240 mg per day, with recommended doses of 120 to 240 mg daily for treatment of dementia and memory impairment.

GB preparations may confer appreciable antiplatelet effects, which are apparently mediated by various ginkgolides. Accordingly, the most frequently reported interactions associated with the use of GB are referring to enhancement of the PD action of platelet inhibitors or anticoagulants. Although clinically important, these PD interactions are beyond the scope of the present review on metabolism- and transporter-based PK interactions.

Various and overall inconsistent *in vitro* effects of GB extracts or specific constituents were demonstrated regarding their potential to alter activities of CYP enzymes. Findings suggesting either induction or inhibition of various CYP enzymes, sometimes concentration-dependently (i.e., inhibition at low concentrations, induction at high concentrations), sometimes in a substrate-dependent fashion have been described. Results have been obtained in part at very high concentrations, unlikely to be achieved at recommended doses *in vivo*. Collectively, the accumulated evidence from *in vitro* studies provides unfortunately little guidance in the reliable prediction of relevant metabolic or transporter-based GB-mediated drug interactions *in vivo*.

In the meantime, however, a considerable number of clinical studies have been conducted for the assessment of GB effects on various CYP isoforms, other drug metabolizing enzymes, and transporters, predominantly by using specific probe drugs, thereby allowing the identification and quantification of alterations of specific CYP enzyme and transporter activities. Overall, we identified a total of 21 human PK studies, reporting original drug interaction data on GB products.

Gurley and coworkers evaluated the effect of GB extract (containing 24% flavone glycosides and 6% terpene lactones), given at doses of 60 mg four times daily over 28 days to 12 healthy adult subjects (6 m/6 f) [17]. In this study, CYP 3A4, 1A2, 2E1, and 2D6 activities were estimated by the use of 1-OH-MDZ/MDZ serum ratios (1-hour sample), paraxanthine/caffeine serum ratios (6-hour sample), 6-OH-CZX/CZX serum ratios (2-hour sample), and debrisoquine urinary recovery ratios (8-hour collection), respectively. Comparisons of pretreatment and post-treatment ratios indicated that GB did not result in any appreciable alteration of CYP 3A4, 1A2, and 2D6 function but appeared to modestly enhance the activity of CYP2E1 by 23%, although this effect was not statistically significant [17].

In a similar study in 12 elderly subjects (6 m/6 f), using the same GB product, dose and treatment duration, as well as the same CYP probe drugs, no significant effects of GB in terms of modulation of CYP 3A4, 1A2, 2E1, and 2D6 activities were found, thereby not confirming the previously observed trend towards a modest CYP2E1 induction [18].

Consistent results were reported by Markowitz and colleagues, who assessed the influence of standardized GB extract on the activity of CYP2D6 and 3A4 in normal adult subjects, phenotyped as CYP2D6 extensive metabolizers. As probe substrates, dextromethorphan (CYP2D6 activity) and alprazolam (CYP3A4 activity)

were coadministered orally at baseline, and following treatment with GB (120 mg b.i.d. = 240 mg/day) for 14 days. No statistically significant differences were found between baseline and post-GB treatment on dextromethorphan metabolic ratios, indicating a lack of GB effect on CYP2D6 activity. For alprazolam there was a statistically significant, albeit modest 17% decrease in total exposure (AUC). Based on these findings, the authors concluded that standardized extracts of GB at recommended doses are unlikely to significantly alter the disposition of coadministered medications primarily relying in their clearance on CYP2D6 or CYP3A4 pathways [19].

The effects of GB supplementation (90 mg/day) administered for 30 days on the steady-state plasma concentrations of the CYP3A4 and CYP2D6 substrate donepezil were investigated in 14 Japanese patients (6 m/8 f; age range 65–80 years) with Alzheimer's disease [20]. The results of this study showed that relatively low doses of GB in elderly patients with Alzheimer's disease did not alter the steady-state plasma concentrations of donepezil, which implies that daily doses of 90 mg of ginkgo do not have appreciable inhibitory or inducing effects on the CYP3A4- and CYP2D6-mediated metabolism of donepezil in the target population.

Robertson and colleagues evaluated the effect of GB (120 mg b.i.d. = 240 mg/day) for two weeks on the steady-state exposure of lopinavir and ritonavir in 14 healthy subjects administered a lopinavir/ritonavir combination [21]. In addition, single oral doses of probe drugs MDZ and fexofenadine were administered prior to and after 4 weeks of GB (following washout of lopinavir/ritonavir) to assess the influence of ginkgo on CYP3A4 and P-gp activity, respectively. Lopinavir, ritonavir, and fexofenadine exposures were not significantly affected by GB administration, while total (AUC) and maximum (C_{max}) exposure of the CYP3A4 prototypic substrate MDZ were significantly lowered by 34% and 31%, respectively, relative to baseline. These results suggest that GB modestly induces – most probably presystemic (i.e., in the gut wall) – CYP3A metabolism, as indicated by the observed decrease in MDZ bioavailability. However, there was no change in the exposure of lopinavir, suggesting that the modest GB-mediated induction of CYP3A4 has most likely been superseded by ritonavir's potent inhibition of CYP3A4. Thus, GB appears unlikely to reduce the exposure of ritonavir-boosted protease inhibitors. As the H_1 -receptor antagonist fexofenadine is a substrate of P-gp (ABCB1), the intestinal uptake transporters organic anion transporting polypeptides 1A2 (OATP1A2) and 2B1 (OATP2B1) as well as the hepatic uptake transporter OATP1B1 [22], the observed unaltered disposition of fexofenadine in the study suggests that GB is unlikely to alter the function of these transporter proteins to a significant extent *in vivo*.

In a small single-dose study in 8 healthy male subjects, 240 mg of GB (Ginkgolon-24™, powder capsules) did not alter the mean exposure to the CYP3A4 substrate nifedipine or its major metabolite dehydronifedipine [23]. Although the observation of remarkably increased nifedipine C_{max} levels in two subjects upon GB coadministration was emphasized by the authors, and possible underlying mechanisms discussed, the known PK variability of nifedipine together with the overall study results indicating absence of any GB effects in terms of total nifedipine exposure (i.e., AUC) rather support the assumption of a lack of a systematic effect of *G. biloba* on the bioavailability of nifedipine. This interpretation would be consistent with reports from other GB interaction studies with CYP3A4 substrates (see above).

Uchida et al. assessed the influence of high repeat doses of GB (360 mg/d Egb 761, Ginkgold®, given as two 60 mg tablets

(120 mg) t.i.d. for 28 days = 360 mg/day) on the PK of orally administered CYP2C9 and CYP3A4 probe drugs tolbutamide (125 mg, single dose) and midazolam (8 mg, single dose) in 10 healthy male subjects, thereof 9 subjects with extensive metabolizer genotypes of CYP2C9 (CYP2C9*1/*1) and 1 subject with CYP2C9*1/*3 genotype [24]. This study showed that high-dose GB treatment slightly decreased the total exposure ($AUC_{0-\infty}$) of tolbutamide by 16% and the metabolic $AUC_{0-\infty}$ ratio of tolbutamide to 4-hydroxytolbutamide in plasma by 17% ($p < 0.05$ for both outcomes). In contrast, total exposure ($AUC_{0-\infty}$) of MDZ was significantly increased by 25% and oral clearance significantly decreased by 26% upon repeated high-dose GB treatment, whereby the metabolic $AUC_{0-\infty}$ ratio of MDZ to 1-OH-MDZ was not significantly altered. The study suggests evidence that a minor CYP2C9 induction can be achieved with supra-therapeutic doses (360 mg/day) of GB *in vivo*, and that some inhibition of intestinal CYP3A4 may go along with a modest increase in the bioavailability of sensitive CYP3A4 substrates. Overall the study results indicate a low potential of GB to confer CYP3A4- and CYP2C9-based metabolic drug interactions, and thus are consistent with reports from other studies that reported no effects of GB on the function of these enzymes at recommended doses.

By using diazepam as a model substrate of CYP2C19 and 3A4, when 10 mg single diazepam doses were given either alone or concomitantly with oral GB (120 mg b.i.d. = 240 mg/day) for 28 days to 12 healthy subjects, Zuo and coworkers demonstrated that total exposure (i.e., AUC) for diazepam and the main metabolite N-desmethyldiazepam were essentially unaltered, thereby indicating that the disposition of CYP2C19 and 3A4 substrates is unlikely to be substantially modified by recommended doses of GB products [25].

Jiang and colleagues investigated the effects of GB on the PK and PD of warfarin in 12 healthy male subjects (eight Caucasians, four Asians) [26]. A single 25-mg dose of warfarin (Coumadin™, 5 × 5-mg tablets) was administered to each subject with and without pretreatment with multiple doses of GB for 1 week (Tavonin™; 40 mg tablets, each tablet containing the standardized dry extract, EGb761, equivalent to 2 g of *G. biloba* leaf, 9.6 mg of ginkgo flavonglycosides, 2.4 mg of ginkgolides and bilobalide, 2 × 40 mg tablets, t.i.d. = 240 mg/day). Dosing of GB was continued for a further week after warfarin administration. The bioanalytics comprised enantiomer-selective quantification of S-warfarin, which is predominantly metabolized to S-7-hydroxywarfarin by CYP2C9, and R-warfarin, which is metabolized by CYP1A2 and CYP3A4, thereby allowing for a separate mechanistic assessment of any potential alteration of these metabolic pathways by concomitant GB treatment. The study demonstrated that GB, at the highest recommended dose, had no effect on the activity of CYP1A2, CYP3A4, or CYP2C9 in healthy subjects, as treatment did not affect the PK and clearance of both warfarin enantiomers in human subjects. GB also did not affect the apparent volumes of distribution or protein binding of warfarin enantiomers. PD endpoints of warfarin 25 mg single doses were also assessed in this study, and it was shown that GB did not significantly alter blood coagulation (i.e., INR) outcomes and platelet aggregation.

Greenblatt and colleagues used single 100 mg doses of the non-steroidal anti-inflammatory drug (NSAID) flurbiprofen as a probe substrate for CYP2C9, to examine the effects of short-term GB treatment (3 doses of 2 × 60 mg tablets = 120 mg, given in 12-hour intervals, as EGb761) in 12 healthy adult subjects (8 m/4 f), who were taking no other medication [27]. The study showed that pretreatment of healthy subjects with usual clinical doses

of GB has no detectable effect on the PK of a single dose of flurbiprofen or on the apparent extent of formation of the principal hydroxylated metabolite. The findings suggest that short-term exposure to GB does not inhibit CYP2C9 activity *in vivo*. As the study involved only short-term exposure to GB, it was not designed to capture possible CYP2C9 induction that could occur with long-term treatment.

Two PK studies in healthy subjects were performed using tolbutamide and diclofenac as probe substrates of CYP2C9. No interactions between GB extract and these CYP2C9 probe substrates were observed *in vivo* as evidenced by the lack of effect on the steady-state pharmacokinetics of diclofenac and the urinary metabolic ratio of tolbutamide [28].

Yin et al. examined the effects of GB on CYP2C19 and CYP3A4 metabolic pathways by using the proton pump inhibitor omeprazole (OPZ), as a well-established CYP2C19 substrate which undergoes CYP2C19-mediated hydroxylation to form its major metabolite 5-hydroxyomeprazole (5-OH-OPZ), and which also undergoes sulfoxidation via CYP3A4 to form omeprazole sulfone (OPZ-SUL) [29]. To decipher differential effects on both pathways, urinary 6 β -hydroxycortisol/cortisol ratios were determined as a marker of CYP3A4 activity before and after GB as well. The study was conducted in 18 healthy Chinese subjects genotyped for CYP2C19 (6 homozygous EMs, 5 heterozygous EMs, and 7 homozygous PMs), and all subjects received a single oral OPZ 40 mg dose at baseline and after a 12-day treatment period with GB (140 mg, two 70 mg tablets b.i.d. = 280 mg/day). Results displayed mean decreases in OPZ $AUC_{0-\infty}$ of 41.5%, 27.2%, and 40.4% ($p < 0.05$ or 0.01, each compared to control) in the homozygous EMs, heterozygous EMs and PMs, respectively. Similarly, decreases in OPZ-SUL $AUC_{0-\infty}$ of 41.2%, 36.0%, and 36.0% ($p < 0.05$ or 0.01, each compared to control), respectively, were noted. However, no significant differences in the $AUC_{OPZ}/AUC_{OPZ-SUL}$ metabolic ratio and the mean 6 β -hydroxycortisol/cortisol ratio were observed before and after GB in each genotype group. In contrast, GB increased the plasma concentrations of 5-OH-OPZ, with mean increases of 37.5%, 100.7%, and 232.4% in the homozygous EMs, heterozygous EMs and PMs, respectively. Accordingly, a significant decrease in the AUC ratio of OPZ to 5-OH-OPZ was observed in the homozygous EMs, heterozygous EMs, and PMs, respectively, whereby the decrease of the metabolic ratio was greater in PMs than EMs. The study results suggest that high-dose GB may induce OPZ hydroxylation in a CYP2C19 genotype-dependent manner but also appears to concurrently reduce the renal clearance of 5-OH-OPZ. Taken together, the observation that in PMs the seemingly inducing effect of GB was particularly pronounced is difficult to interpret, because of the obvious difficulty to induce CYP2C19 at all in PMs. On the other hand, the CYP3A4 activity was unaffected as shown by the lack of changes in the $AUC_{OPZ}/AUC_{OPZ-SUL}$ as well as 6 β -hydroxycortisol/cortisol ratios before and after GB. Overall, the study indicates that high-dose treatment with GB may significantly induce the clearance of CYP2C19 substrates, thereby possibly reducing their clinical effects.

Lei et al. examined the effects of GB (120 mg b.i.d. for 12 days) as an inducer of CYP2C19 in 14 healthy Chinese subjects genotyped as either CYP2C19 extensive (EMs; 7 carriers of 2C19*1/2C19*1) or poor metabolizers (PMs; 7 carriers of 2C19*2/2C19*2), by assessing the single dose PK of the orally administered CYP2C19 substrate voriconazole (200 mg) [30]. In EMs, the median voriconazole $AUC_{0-\infty}$ values were 5.17 $\mu\text{g} \times \text{h/mL}$ and 4.28 $\mu\text{g} \times \text{h/mL}$ after administration of voriconazole alone and after voriconazole with concomitant GB treatment, respectively, a 17% difference

suggesting a trend towards a modest induction of CYP2C19 by GB, although results did not achieve statistical significance. All other PK parameters of voriconazole such as AUC_{0-24} , time to reach maximum concentration (t_{max}), terminal disposition half-life ($t_{1/2}$), and apparent oral clearance (CL/F) also did not change significantly in the presence of GB in EMs of CYP2C19, and the results followed a similar pattern in PMs of CYP2C19. Based on the results of this study, it can be concluded that – even at the highest recommended doses – GB-mediated CYP2C19-based PK drug interactions appear to be weak and may have limited clinical significance.

Recently, the effects of GB (240 mg per day, administered as two 60-mg capsules b.i.d. for 14 days) on the activity of CYP2B6 was investigated in 14 healthy adult male Chinese subjects (age range 19–25 years), by assessment of the single dose PK of the antidepressant and smoking cessation drug bupropion (150 mg), which is almost exclusively metabolized by CYP2B6 to its active metabolite hydroxybupropion. The study showed that a 14-day oral administration of GB extract at the high end of the recommended dose range had no statistically significant effect on the total exposure of bupropion or its active metabolite hydroxybupropion, as measured by plasma AUC [31]. These results indicate that GB is unlikely to alter the disposition of sensitive CYP2B6 substrates.

Mauro and coworkers assessed the effect of recommended GB doses (240 mg/day, administered as two 40-mg tablets t.i.d. for 8 days) on the pharmacokinetics of the P-gp (ABCB1) substrate digoxin (0.5 mg) in eight healthy human subjects (7 m/1 f, mean age 23 ± 3 yrs) [32]. The total digoxin exposure ($AUC_{0-\infty}$) was observed with 21.0 ± 8.6 ng/mL \times h (digoxin alone) and 25.6 ± 13.2 ng/mL \times h (digoxin + GB), indicating a trend towards a modest (i.e., 22%) increase in digoxin exposure associated with a 1-week GB pretreatment. Although the authors emphasized that the digoxin AUC outcomes were not significantly different from each other, absence of GB treatment on P-gp (ABCB1) function should not be concluded from the study due to the lack of statistical power (small sample size).

The capacity of high-dose GB treatment (360 mg/day for 14 days) to alter the function of P-gp (ABCB1) was examined by Fan and coworkers in 10 healthy male subjects by using the P-gp (ABCB1) substrate talinolol (100 mg, single dose) [33]. GB increased the maximum observed plasma concentration of talinolol (C_{max}) by 36% (90% CI 10 to 68; $p = 0.025$), AUC_{0-24} by 26% (90% CI 11 to 43; $p = 0.008$), and $AUC_{0-\infty}$ by 22% (90% CI 8 to 37; $p = 0.014$), respectively, without significant changes in terminal disposition half-life ($t_{1/2}$) and the time to C_{max} (t_{max}). The authors concluded that the results indicate that repeated supra-therapeutic doses of GB may modestly inhibit intestinal P-gp, thereby increasing the bioavailability of sensitive P-gp transporter substrates. Comparable results were obtained by the same group of investigators in another study of similar design [34].

The long-term administration of GB (EGb 761 120 mg/day for 90 days) to 10 healthy adult subjects and 10 patients with type 2 diabetes mellitus (T2DM) did not significantly affect the PK of metformin single doses (500 mg in healthy subjects and 250 to 850 mg in T2DM patients) [35]. As metformin is a substrate of the plasma membrane monoamine transporters (PMAT), which are localized on the luminal side of enterocytes and were recently suggested to be the major transporter system responsible for the uptake of metformin from the gastrointestinal tract [36], the study suggests that long-term treatment with intermediate GB doses does not alter PMAT-mediated metformin absorption. As metformin, which is a hydrophilic organic cation (pK_a 12.4) that

is not significantly metabolized in humans is a good probe drug for organic cation transport (e.g., OCT1 and OCT3) in intestinal drug absorption, hepatic uptake (OCT1), and is also a substrate for several other polyspecific tubular OCT transporters in the kidney (e.g., OCTN1, OCTN2, MATE1, and MATE2-K may be involved in the renal excretion of metformin in humans) [37], it can be concluded that GB, at the dose level investigated, is unlikely to alter the disposition of OCT substrates to a significant extent.

Eight healthy subjects (4 m/4 f) were treated with single oral 250 mg doses of ticlopidine, a putative substrate of the organic anion transporting polypeptide OATP-B, either given alone or together with repeat dose GB (40 mg t.i.d. = 120/day; unspecified product) for 3 days [38]. In this study, short-term and low-dose coadministration of GB did not significantly change any of ticlopidine's PK parameters. These results are consistent with another study in which single coadministration of GB extract (80 mg) with ticlopidine (250 mg) in healthy male Korean subjects was not associated with any significant changes in the PK profile of ticlopidine compared with ticlopidine administered alone [39]. The information that can be derived from both studies in terms of clinical significance is limited, because of the low GB doses employed, and the short-term/single dose posology, respectively.

In the most recent study, a cocktail phenotyping design was employed to assess the interaction profile of GB extract EGb 761[®] on the activities of the major CYP enzymes CYP1A2, CYP2C9, CYP2C19, CYP2D6, and CYP3A [40]. EGb 761[®] extract or matched placebo were administered for 8 days to 18 healthy subjects (8 m/10 f) according to a 3-fold crossover design; total daily doses of 240 mg EGb 761[®] were given in one study period following a 120 mg b.i.d. posology (= 240 mg/day), and in the other study period 240 mg doses given q.d. in the morning. The phenotyping cocktail was orally administered before and after the EGb 761[®]/placebo treatment periods and employed the following probe substances and metrics: 150 mg caffeine (CYP1A2, paraxanthine/caffeine plasma ratio 6-h postdose), 125 mg tolbutamide (CYP2C9, plasma concentration 24-h postdose), 20 mg OPZ (CYP2C19, OPZ/5-OH-OPZ plasma ratio 3-h postdose), 30 mg dextromethorphan (CYP2D6, dextromethorphan/dextrorphan plasma ratio 3-h postdose), and 2 mg of MDZ (CYP3A, MDZ plasma concentration 6-h postdose).

Point estimates of EGb 761[®]/placebo ratios for phenotyping metrics were close to unity for all CYPs, except for CYP2C9 (0.834 for 120 mg b.i.d.; 0.848 for 240 mg q.d.) and CYP2C19 (0.874 for 120 mg b.i.d. and 0.896 240 mg q.d.), which may suggest a weak trend towards induction of these CYP2C family enzymes. Respective CIs for most CYP enzymes were within the prespecified acceptance margins (i.e., 0.70–1.43) for all ratios except CYP2C19 for EGb 761[®] 120 mg b.i.d. (90% CI 0.681–1.122) and CYP2D6 for EGb 761[®] 240 mg q.d. (90% CI 0.667–1.281). Taken together the data clearly indicate the absence of any CYP inhibitory potential of 240 mg daily doses of EGb 761[®] extract towards CYP1A2, CYP2C9, CYP2C19, CYP2D6, and CYP3A, regardless whether the product is given once or twice daily. The study also showed that EGb 761[®] extract does not induce the activity of CYP1A2, CYP2D6, and CYP3A enzymes, and may confer a weak induction of CYP2C9 and CYP2C19 enzymes, which, however, appears too small to be of clinical relevance.

Taken together, besides St. John's Wort, GB appears to be at the time being, the second best studied natural health product, in terms of the clinical investigation of its metabolic and transporter-based PK drug interaction potential. The available set of published mechanistic drug-drug interaction studies allows rea-

sonable inferences on the respective drug metabolizing enzyme and transporter effects of GB products *in vivo*. The information which can be derived from these studies is largely consistent, with some minor discrepancies that may be explained by differences in study design and populations, employed doses and treatment durations, and perhaps also differences in the composition of products investigated.

Collectively, the available evidence indicates that recommended doses of GB up to daily doses of 240 mg do not have significant or clinically meaningful effects on the activity of CYP enzymes such as CYP1A2, CYP2C9, CYP2D6, CYP2B6, CYP2E1, and CYP3A4. While data on PK alterations of CYP3A4 are not entirely consistent, sometimes suggesting the possibility of presystemic induction of CYP3A4 with a high-dose treatment of GB, sometimes – in turn – suggesting CYP3A4 inhibition, the overall CYP3A4-related effect sizes described are generally weak and unlikely to be of clinical relevance unless CYP3A4 substrates with a narrow therapeutic index are concerned.

CYP2C19 appears to be modestly inducible by GB products, with across study evidence that this effect appears to emerge at the highest recommended dose level of 240 mg/day [30] and may become somewhat more pronounced with increasing doses [29]. Differences in the effect sizes of CYP2C19 induction, however, may also depend on the individual CYP2C19 genotype of subjects [29] and perhaps on substrate characteristics. The available evidence in this respect, however, appears to be not entirely consistent and requires further investigation.

Although not as extensively studied as the metabolic interactions, the overall propensity of recommended doses of GB products to alter the functionality of transporter proteins appears low. Therapeutic and supra-therapeutic *G. biloba* doses (i.e., 240 and 360 mg/day) were shown to modestly increase the bioavailability of the P-glycoprotein (ABCB1) probe substrates digoxin and talinolol by about 22% to 26% [33, 34]. A study employing *G. biloba* treatment with 240 mg/day, however, did not show any alterations in the disposition of fexofenadine, which is a substrate of P-glycoprotein (ABCB1), the intestinal uptake transporters organic anion transporting polypeptide 1A2 (OATP1A2) and 2B1 (OATP2B1), and the hepatic uptake transporters OATP1B1 [41], thereby indicating that GB is unlikely to alter the function of these transporter proteins to a significant extent *in vivo* [21]. Similarly intermediate doses of GB (120 mg/day) do not appear to alter the function of membrane monoamine transporters (PMAT), organic cation transporters (e.g., OCT1 and OCT3), and other polyspecific tubular OCT-transporters in the kidney (e.g., OCTN1, OCTN2, MATE1, and MATE2-K), as shown by a recent metformin interaction study [37].

Milk thistle

Silybum marianum (family Asteraceae/Compositae), commonly known as milk thistle (MT), is one of the oldest and most extensively studied plants in the treatment of alcoholic, toxic, and viral liver diseases. Other treatment claims include prevention/amelioration of drug-induced liver toxicity (DILI), e.g., due to cancer chemotherapy, nonalcoholic fatty liver disease (NAFL), or steatohepatitis (NASH), lowering of cholesterol levels, and improvement of insulin resistance in patients with type 2 diabetes (T2DM) [42]. The active extract of MT is denoted as silymarin and represents a mixture of the flavonolignan constituents silybin (silibinin), isosilybin, silidianin, and silichristin. Silymarin is extracted from dried MT seeds, in which it is contained at higher concentrations than in other parts of the plant. Silymarin has

structural similarities to steroid hormones, which may be linked to its pharmacological actions. Silybin is the predominant and pharmacologically most active component, constituting approximately 60% to 70% of the isomers, followed by silychristin (20%), and silidianin (10%) [43]. The silymarin content in MT extracts may range from 40% to 80% [43]. Customary daily doses of MT extract range between 210 to 800 mg, depending on patient characteristics and therapeutic objectives.

In vitro studies implicate that silymarin and silybin can display inhibition of a variety of CYP enzymes such as CYP3A4, CYP2C8, CYP2C9, CYP2C19, CYP2D6, or CYP2E1, and also a couple of UGTs, such as UGT1A1, 1A6, 1A9, 2B7, and 2B15 [44–47]. In contrast, there is no conclusive *in vitro* evidence that MT would have significant effects on the activity of P-gp (ABCB1) [45, 48].

The available evidence from *in vitro* studies, has been acknowledged by clinical researches, and a number of clinical studies have been conducted for the *in vivo* assessment of the effects of MT products on various CYP isoforms, other drug metabolizing enzymes, and transporters, predominantly by using specific probe drugs, thereby allowing the identification and characterization of MT effects on specific metabolic and disposition pathways. Altogether, we identified a total of 13 pharmacokinetic drug interactions with orally administered MT extract in the public domain.

In the oldest published human MT interaction study, administration of silymarin at the lower end of usually recommended daily doses (Legalon[®], 70 mg t.i.d. = 210 mg/day) for 28 days to 16 healthy subjects had no effect on the PK of aminopyrine or phenylbutazone, two nonspecific CYP probes [49].

Gurley and colleagues studied 12 healthy subjects (6 m/6 f), who were randomly assigned to receive MT (175 mg b.i.d. = 350 mg/day, standardized to 80% silymarins) for 28 days [50]. Probe drug cocktails of midazolam (MDZ; CYP3A4) and caffeine (CYP1A2), followed 24 hours later by chlorzoxazone (CYP2E1) and debrisoquine (CYP2D6), were administered before (baseline) and at the end of supplementation. Presupplementation and postsupplementation assessments of CYP3A4, CYP1A2, CYP2E1, and CYP2D6 activities were made by using 1-hydroxymidazolam (1-OH-MDZ)/MDZ serum ratios (1-hour sample), paraxanthine/caffeine serum ratios (6-hour sample), 6-hydroxychlorzoxazone/chlorzoxazone serum ratios (2-hour sample), and debrisoquine urinary recovery ratios (8-hour collection), respectively. The results of this study indicate that daily doses of 350 mg of a standardized MT extract did not alter human CYP1A2, CYP2D6, CYP2E1, and CYP3A4 enzyme activities *in vivo* and thus, do not appear to pose a significant interaction potential for substrates of these enzymes [50].

In another study conducted by the same group of investigators, 19 healthy young subjects (10 m/9 f) received substantially higher daily doses of a standardized MT supplement for 14 days (900 mg/day). Subjects also received rifampin (300 mg b.i.d.) and clarithromycin (500 mg b.i.d.) for 7 days as positive controls for CYP3A induction and inhibition, respectively. Oral MDZ (8 mg) was administered as a CYP3A4 probe before and after supplementation and control periods [51].

In contrast to the positive controls rifampin and clarithromycin, no significant changes in MDZ PK were observed as a result of MT supplementation. Likewise, comparisons of 1-hour 1-OH-MDZ/MDZ ratios before and after supplementation/medication mirrored the effects observed for MDZ concentration-time profiles, i.e., no significant changes in mean 1-hour 1-OH-MDZ/MDZ serum ratios were noted after a 14-day course of MT

(900 g/day), thereby confirming that MT extract is not a clinically relevant modulator of CYP3A4 activity *in vivo*.

Likewise, coadministration of 560 mg silymarin (4 silymarin 140 mg capsules, Legalon®) administered 10 hours and 1.5 hours prior to the administration of the CYP3A4 substrate nifedipine (10 mg IR tablet) to 16 young healthy male subjects did not considerably change the extent of absorption or metabolism of nifedipine. Overall, the total extent of nifedipine exposure (i.e., AUC) was 13% higher (90% CI 0.97- to 1.32) in the silymarin period, while C_{max} values were on average 30% lower (90% CI 0.39- to 1.27), consistent with a trend towards a delayed nifedipine absorption in the silymarin period [52]. On the whole, also this study confirmed that silymarin does not act as a potent CYP3A4 inhibitor *in vivo*. The data suggest modest effects of silymarin on the absorption characteristics of nifedipine, resembling the known pattern of nifedipine food effects. The authors discussed that underlying mechanisms could be either a delay in the gastric emptying rate by silymarin or inhibition of OATP transporters expressed in the GI tract [52].

The effects of a 14-day treatment with a 900-mg daily dose of a standardized MT extract (300 mg t.i.d., standardized to contain 80% silymarin per capsule) on the activity of CYP2D6 were investigated in 16 healthy subjects (8 m/8 f), by administration of the CYP2D6 substrate debrisoquine (5 mg) before and at the end of MT supplementation, by using 8-h debrisoquine urinary recovery ratios [53]. Results did not indicate significant inhibition of CYP2D6 by MT supplementation (post/pre geom. mean ratio 0.97, 90% CI 0.93 to 1.10).

Two PK interaction studies, each of them conducted in 10 healthy male and female subjects, examined if MT extract may have the potential to alter the PK of the HIV protease inhibitor and CYP3A4 and P-gp substrate indinavir [54,55].

In the first study, subjects received four oral doses of indinavir 800 mg every 8 hours for baseline PK, followed by a 3-week MT treatment (175 mg tablets t.i.d. = 525 mg/day; each tablet containing 153 mg of silymarins, equivalent to 191 mg of MT extract standardized for 80% silymarins; Thisilyn). Then the course of four oral doses of indinavir 800 mg every 8 hours was repeated on top of continued MT treatment, and results were compared with baseline indinavir PK outcomes [55]. Indinavir plasma concentrations were followed for 8 h postdose. Results showed that regular MT treatment over 3 weeks did not significantly alter the overall exposure to indinavir, as evidenced by a nonsignificant average 9% reduction in indinavir + MT AUC_{0-8} compared to indinavir alone treatment at baseline. On examination of individual patients, the AUC_{0-8} of indinavir in the presence of MT was 70–107% of the AUC_{0-8} for indinavir alone. Average indinavir trough values taken at 8 hours postdose, displayed a statistical significant, albeit modest average decline of about 25% upon MT treatment.

In the second indinavir/MT interaction study, indinavir 800 mg t.i.d. was given for four doses, followed by MT 160 mg t.i.d. (= 480 mg/day standardized MT) for 13 days, with both drugs given at the same dosages over the following 2 days [54]. Indinavir plasma concentrations were followed up to 5 hours postdose. When given alone and combined with MT, respectively, the geometric mean (95% CIs) steady-state indinavir AUC was 20.7 hr·mg/L (15.3–28.2 hr·mg/L) and 19.4 hr·mg/L (15.8–23.6 hr·mg/L), and the trough plasma concentrations were 0.340 mg/L (0.232–0.497 mg/L) and 0.232 mg/L (0.129–0.419 mg/L). Average observed maximum exposure (C_{max}) was 8.85 (7.17–10.9) mg/L and 7.85 mg/L (6.55–9.30) for indinavir alone and indinavir + si-

lymarin, respectively. None of the PK outcomes for indinavir indicated a statistical significant alteration by MT treatment, and the authors concluded that a daily dosage of 480 mg of a standardized silymarin extract has no apparent effect on indinavir plasma concentrations. Taken together, both indinavir interaction studies yielded consistent results, thereby indicating that MT supplementation at commonly recommended doses does not alter the activity of CYP3A4 *in vivo* to a clinically relevant extent.

A study with another CYP3A4 and putative P-gp substrate, the H_2 receptor antagonist ranitidine, also yielded a consistent outcome in that regard [56]. In this trial, 12 healthy male subjects received oral ranitidine 150 mg either alone or after a 7-day pretreatment with silymarin 140 mg t.i.d. (= 420 mg/day). There was no influence of silymarin coadministration on the pharmacokinetics of ranitidine (i.e., C_{max} and AUC).

In a sample of 6 cancer patients, van Erp and colleagues investigated whether short-term MT supplementation affects the PK of the anticancer drug irinotecan, a substrate for CYP3A4 and UGT1-A1 enzymes [57]. Patients were treated with irinotecan (dose 125 mg/m²) given as a 90-minute infusion once weekly. Four days before the second irinotecan dose, patients received 200 mg MT capsules t.i.d. (= 600 mg/day), for 14 consecutive days (each capsule containing 200 mg MT seed extract, containing 80% silymarin). Short-term (4 days) or more prolonged (12 days) intake of MT in this study had no significant effect on irinotecan clearance (mean, 31.2 vs. 25.4 vs. 25.6 L/h; $p=0.16$). The AUC ratio of the active metabolite SN-38 and the irinotecan parent drug was slightly decreased by MT (2.58% vs. 2.23% vs. 2.17%; $p=0.047$), whereas the relative extent of glucuronidation of SN-38 was similar (10.8 vs. 13.5 vs. 13.1; $p=0.64$). Likewise, the AUC ratio of 7-ethyl-10-[4-N-(5-aminopentanoic acid)-1-piperidino]-carbonyloxycamptothecin and irinotecan was unaffected by MT (0.332 vs. 0.285 vs. 0.337; $p=0.53$). This study showed that the concurrent use of MT does not significantly influence the PK of irinotecan in cancer patients. The authors indicated that irinotecan is known to be highly susceptible to CYP3A4 and P-gp (ABCB1) inhibition and induction in humans and concluded that the results of the current study confirm that MT does not seem to modulate the activity and/or expression of these proteins. They also noted that this finding is consistent with the notion that the MT constituent silybin is not a ligand activator of the pregnane X receptor (PXR), a steroid- and xenobiotic-regulated transcription factor that is a key regulator of constitutive expression and induction of CYP3A4, UGT1A, and ABCB1. The results of the irinotecan study also provide additional information in support of an apparent lack of effect of MT on the function of human carboxylesterase 2-mediated hydrolysis and UGT1A-mediated phase II conjugation pathways in humans.

There are two studies regarding the assessment of potential MT effects on the PK of more or less sensitive CYP2C9 substrates available [58,59]. Han and colleagues investigated the effects of silymarin on the PK of the angiotensin II (ATII) receptor antagonist losartan (50 mg) and its active metabolite E-3174 and its possible relationship to CYP2C9 genotypes because current evidence suggests that losartan biotransformation to E-3174 is highly dependent on CYP2C9 activity [59]. Twelve healthy adult male subjects of known CYP2C9 genotype (six CYP2C9*1/*1 and six CYP2C9*1/*3) were enrolled, and the PK of losartan and E-3174 were assessed before and after a 14-day treatment with 140 mg of silymarin t.i.d. (= 420 mg/day; Legalon® 140 mg capsules). About ≥ 2 -fold differences in the AUC_{0-24} (409.9 \pm 76.9 vs. 854.3 \pm 236.9 ng h/mL, placebo vs. silymarin, $p < 0.05$), $AUC_{0-\infty}$

(425.0 ± 83.9 vs. 886.0 ± 221.1 ng h/mL, placebo vs. silymarin, $p < 0.05$) and C_{\max} (170.9 ± 54.6 vs. 323.3 ± 121.0 ng/mL, placebo vs. silymarin, $p < 0.05$) of losartan between the placebo and silymarin treatments were observed in carriers of CYP2C9*1/*1. Accordingly, the oral clearance (CL/F) of losartan was also significantly decreased by more than 2-fold after a 14-day silymarin treatment in these subjects (121.9 ± 27.2 vs. 59.7 ± 159.7 L/h, placebo vs. silymarin, $p < 0.05$). In contrast, in carriers of CYP2C9*1/*3, no significant differences in any of losartan PK parameters between the placebo and silymarin treatment groups were observed, e.g., AUC_{0-24} was 816.3 ± 282.9 vs. 805.8 ± 266.2 ng h/mL, and $AUC_{0-\infty}$ was 866.0 ± 292.8 vs. 871.9 ± 23 ng h/mL. Likewise, the metabolic ratio of losartan (ratio of $AUC_{0-\infty}$ of E-3174 to $AUC_{0-\infty}$ of losartan) was significantly decreased after silymarin treatment in carriers of CYP2C9*1/*1 ($p < 0.05$), but not in those with the CYP2C9*1/*3 genotype ($p = 0.065$). Notably, the total exposure parameters (i.e., AUC_{0-24} and $AUC_{0-\infty}$) in placebo-treated carriers of CYP2C9*1/*3 were well comparable to silymarin-treated CYP2C9*1/*1 subjects, thereby suggesting that the extensive CYP2C9 metabolizers (i.e., carriers of CYP2C9*1/*1 wild-type) were actually subject to a silymarin-mediated pharmacological conversion to heterozygous intermediate CYP2C9 metabolizers (i.e., CYP2C9*1/*3).

As CYP2C9 appears responsible for oxidative conversion of losartan to its pharmacologically active carboxylic acid metabolite E-3174, the data of the present study are consistent with a substantial CYP2C9 inhibitory potential of silymarin treatment at customary doses *in vivo*. As E-3174 has a higher ATII antagonistic potency and longer half-life than the parent compound losartan, the active metabolite may be possibly responsible for most of the antihypertensive effect, and the decrease in the metabolic ratio of losartan to E-3174 due to decreased CYP2C9 activity may diminish the efficacy of losartan in carriers of CYP2C9*1/*1 [59]. The study results of Han et al. were recently followed up by Brantley and colleagues, who identified *in vitro* that silybin A and silybin B were the most potent CYP2C9 inhibiting constituents of silymarin extract, with K_i values of 4.8 and 10 μ M, respectively [44]. In mechanistic terms, the available CYP2C9-related evidence suggests that MT treatment may have a potential to inhibit the clearance and to increase the exposure of other sensitive CYP2C9 substrates such as (S)-warfarin *in vivo*.

An earlier study examined the influence of a 9-day silymarin pretreatment at a low daily dose of 140 mg on the metabolism and disposition of metronidazole (3 × 400 mg/day), a substrate of CYP3A4 and CYP2C9, in 12 healthy adult subjects [58]. In this study, low-dose silymarin modestly increased the clearance of metronidazole and its major metabolite, hydroxy-metronidazole (HM), by 29.51% and 31.90%, respectively, with a concomitant decrease in half-life, C_{\max} , and AUC_{0-48} . However, urinary excretions of acid-metronidazole (AM), HM as well as metronidazole in 48 h were decreased. The authors concluded that the results indicate that silymarin might induce both intestinal P-gp and CYP3A4 upon multiple dose administration. However, this interpretation would not be consistent with all other published *in vivo* studies examining the effects of silymarin on CYP3A4 and P-gp function. Hence, the underlying mechanism(s) of the reported modest silymarin/metronidazole PK interaction remain(s) to be determined.

Two studies investigated the effect of MT administration on the PK of the prototypic P-gp (ABCB1) substrates digoxin [60] and the β -adrenoceptor antagonist talinolol [61], with the aim to examine potential *in vivo* effects of MT treatment on P-gp function.

Sixteen healthy young subjects (8 m/8 f) were randomly assigned to receive a standardized MT supplement for 14 days (300 mg t.i.d. = 900 mg/day, standardized to contain 80% silymarin), or to receive rifampin (300 mg b.i.d. = 600 mg/day, 7 days) and clarithromycin (500 mg b.i.d. = 1000 mg/day, 7 days) as positive controls for P-gp (ABCB1) induction and inhibition, respectively. Oral digoxin (Lanoxicaps™, 0.4 mg) was administered as a P-gp probe before and at the end of supplementation and control periods. Serial digoxin serum concentrations were obtained over 24 h [60].

As expected, rifampin produced significant reductions ($p < 0.01$) in measures of total (AUC_{0-3} , AUC_{0-24}) and maximum (C_{\max}) digoxin exposure, whereas clarithromycin increased these parameters significantly ($p < 0.01$). In turn, no remarkable effects on digoxin PK were observed following supplementation with MT, although digoxin AUC_{0-3} and AUC_{0-24} , and C_{\max} were altogether found modestly reduced following MT administration by about 9.5%, 12.5%, and 13.5%, respectively. These results do not support reported *in vitro* data suggesting a potential inhibition of P-gp function by MT constituents and also do not suggest a clinically relevant modulation of P-gp activity by MT supplementation at customary doses.

Han and coworkers examined the effect of concomitantly administered silymarin on the pharmacokinetics of talinolol, a drug mainly eliminated into the intestine by P-gp, in 18 healthy male Chinese subjects, and its association with a genetic single nucleotide polymorphism in exon 26 (C3435T) of the MDR-1 (ABCB1) gene [61], which was shown to be correlated with intestinal expression and *in vivo* activity of P-gp [62]. The PK of talinolol was assessed after coadministration of either placebo or 140 mg silymarin capsules t.i.d. (420 mg/day) for 14 days, and talinolol plasma concentrations were followed for 36 hours. Talinolol peak plasma concentrations (C_{\max}) were significantly increased after silymarin administration as compared with placebo ($p = 0.007$), and the total exposure estimates (AUC_{0-36} and $AUC_{0-\infty}$) of talinolol were increased on average by about 36% by silymarin coadministration. Accordingly, the oral clearance (CL/F) of talinolol was decreased by about 23% ($p < 0.001$). These effects were unrelated to the MDR-1 C3435T single nucleotide polymorphism, thereby suggesting a P-gp-unrelated mechanism of this finding, probably consistent with an upregulation of intestinal organic anion transporting polypeptide (OATP) mediated uptake of talinolol [63].

Deng and coworkers examined the interaction of MT extract and the cholesterol lowering HMG-CoA-reductase inhibitor rosuvastatin [64]. Eight healthy Korean male subjects were either treated with placebo or silymarin (Legalon® 140 mg capsules) t.i.d. (= 420 mg/day) for 5 days. Subjects received a single 10 mg dose of rosuvastatin on day 4 of silymarin supplementation, and serial blood samples were collected for 72 h. During plasma sampling, subjects continued to take silymarin or placebo capsules t.i.d. for 2 further days. The plasma concentration–time profile of rosuvastatin showed no significant effect of silymarin treatment. C_{\max} values (mean ± SD) for rosuvastatin were 13.7 ± 4.7 ng/mL and 14.8 ± 5.3 ng/mL in the presence and absence of silymarin, respectively. The AUC values (mean ± SD) for rosuvastatin with and without silymarin treatment were 134.6 ± 42.3 and 144.0 ± 38.7 ng h/mL, respectively. Thus, short-term silymarin treatment at customary doses did not significantly affect the PK of rosuvastatin, a multitransporter substrate, including the intestinal efflux/uptake transporters P-gp (ABCB1), BCRP (ABCG2), and OATP1A2, and the hepatic uptake transporters OATP1B1, 1B3, and

2B1 [65], indicating that silymarin is not a potent modulator of ABC and OATP transporters *in vivo*.

In conclusion, the available evidence from a total of 13 clinical PK interaction studies consistently indicates that marketed MT/silymarin products, given at customary or supra-therapeutic doses, do not alter the function of CYP3A4, CYP1A2, CYP2D6, and CYP2E1 enzymes, or intestinal ABC and OATP transporters to a relevant extent, although the findings regarding a potential modulation of the activity of certain OATP transporters is not sufficiently detailed and entirely conclusive as yet, and deserves further investigation. Due to the total number of studies available and the high degree of consistency in overall study outcomes, the evidence in terms of a lack of interaction potential is most robust for CYP3A4 and P-gp (ABC1). The results from a single study in irinotecan-treated cancer patients also suggest that the function of UGT1A1 and carboxylesterase 2 are unlikely to be substantially altered by MT/silymarin.

In contrast, the available evidence from one well-designed study in Chinese subjects regarding CYP2C9 [59], together with consistent *in vitro* data [44], strongly suggests the possibility of significant CYP2C9 inhibition by MT/silymarin. These data call for further investigation and confirmation of the clinical significance of this matter with narrow therapeutic index CYP2C9 substrates such as warfarin.

Ginseng

Several ginseng species (family Araliaceae) need to be distinguished including *Panax ginseng* (Asian ginseng), *Panax quinquefolius* (American ginseng), and Siberian ginseng (*Eleutherococcus senticosus* Maxim). For centuries ginseng has been used as a traditional herbal medicine in many Asian countries. Today ginseng products are consumed worldwide and rank among the most commonly used herbal products. Actually, ginseng is the 5th best-selling herb in the United States [66]. Ginseng products are advocated for many conditions, including maintenance of general health, physical and mental well-being, treatment of fatigue, weakness and mild depression, improving immune function, and for conferring antioxidant and antineoplastic pharmacological effects [67, 68].

Among the different principal ginseng constituents, which are derived from the dried roots of the plants, the major active components are ginsenosides, which represent a diverse group of steroidal saponins [68]. Some remarkable differences between ginseng species exist regarding the content and composition of ginsenosides, a factor that may be associated with a differential drug interaction potential. The recommended daily dose of *P. ginseng* is 200 mg of standardized (i.e., 4% total ginsenosides) extract.

The effects of ginseng extracts or individual ginsenosides on CYP enzymes have been examined in many published *in vitro* studies. Unfortunately the results have been largely inconsistent and remain controversial. Also the concentrations required to elicit CYP inhibition or induction *in vitro* were often very high and unlikely to be achieved *in vivo* with recommended doses of marketed products [69]. Hence, reliable predictions on the *in vivo* relevance of these findings are hardly possible.

Clinical studies on PK ginseng-drug interactions are scarce; we identified a total of 9 published PK interaction studies reporting original data.

In a phenotyping study measuring single timepoint, phenotypic metabolic ratios in elderly subjects with administration of the prototypic enzyme-specific probe drugs MDZ (CYP3A4), caffeine

(CYP1A2), chlorzoxazone (CYP2E1), and debrisoquine (CYP2D6), the potential of *P. ginseng*, at a dose of 500 mg t.i.d. (= 1.500 g/day, standardized to 5% ginsenosides) given for 28 days, to alter the activities of these CYP enzymes was investigated [18]. In this setting, *P. ginseng* did not alter the function of CYP3A4, CYP1A2, and CYP2E1, and displayed only a marginal reduction of CYP2D6 activity (~ 7%), an effect magnitude not being of clinical concern. In a similar study in healthy young subjects, using the same *P. ginseng* product, dose, and treatment duration, as well as the same CYP probe drugs, no significant effects of *P. ginseng* on any CYP enzyme activity were found [17].

The findings of Gurley et al. in terms of unaltered CYP3A4 activity were confirmed by another study, in which *P. ginseng* 100 mg doses (Ginsana™, standardized to 4% ginsenosides) were given b.i.d. (= 200 mg/day) for 14 days to 20 healthy male and female subjects (10 Caucasians, 10 Asians). In this study, the urinary 6- β -hydroxycortisol/cortisol ratio, a marker of CYP3A enzyme induction, remained unaltered, thereby indicating that *P. ginseng*, at recommended doses, does not confer significant CYP3A induction *in vivo* [70].

The effects of Siberian ginseng (SG; *Eleutherococcus senticosus* Maxim; 485-mg capsules of standardized extract twice daily) on the activity of CYP2D6 and 3A4 were assessed in a 2-week study in 12 healthy male and female subjects [71]. Dextromethorphan (CYP2D6) and alprazolam (CYP3A4) were used as probe drugs. There were no statistically significant differences between pre- and post-SG treatment urinary dextromethorphan/dextrorphan metabolic ratios, indicating a lack of an SG effect on CYP2D6 enzyme activity. For alprazolam there also were no significant differences in plasma concentrations and derived PK parameters (C_{max} , t_{max} , AUC, $t_{1/2}$) indicating that SG does not significantly induce or inhibit CYP3A4. These results are consistent with the outcomes reported for *P. ginseng* preparations as detailed above and indicate that standardized extracts of SG at generally recommended doses are unlikely to alter the disposition of coadministered medications primarily dependent on the CYP2D6 or CYP3A4 pathways for their elimination [71].

The effects of American ginseng (AG; *Panax quinquefolius*) on the disposition of the HIV protease inhibitor and sensitive CYP3A4 substrate indinavir were assessed in 14 healthy adult male and female subjects (12 African-American and two Caucasians) [72]. Indinavir at 800 mg doses was administered t.i.d., either alone or together with 1.0 g doses of AG three times daily over 2 weeks (dried whole root 500 mg capsules). The study demonstrated that AG did not alter the PK of the CYP3A4 substrate indinavir, thereby demonstrating the absence of appreciable CYP3A4 induction or inhibition.

Lee and colleagues assessed the effects of a 2-week treatment of 200 mg b.i.d. (= 400 mg/day) enriched AG extract (HT1001 100 mg capsules; REMEMBER-fx; total ginsenoside content of $8.5 \pm 0.5\%$) on the PK profile of the nucleoside reverse transcriptase inhibitor zidovudine (single 300-mg oral doses of Retrovir™) [73]. This study provided the first *in vivo* insights into the potential effects of AG on phase 2 drug metabolizing enzymes, as zidovudine is predominantly cleared through glucuronidation by UGT 2B7. The study results showed that AG does not significantly affect the formation clearance of zidovudine to its glucuronide (ratio post- to pre-AG = 1.17; 90% confidence interval: 0.95–1.45; $p = 0.21$), total clearance (ratio = 0.97; 0.82–1.14; $p = 0.70$), or total plasma zidovudine exposure (AUC_{0-8} ratio = 1.03; 0.87–1.21; $p = 0.77$).

Jiang and colleagues investigated the effects of *P. ginseng* on the PK and PD of warfarin in 12 healthy male subjects (eight Caucasians, four Asians) [74]. A single 25-mg dose of warfarin (Coumadin™, 5 × 5-mg tablets) was administered to each subject with and without pretreatment with multiple doses of ginseng for 1 week (Golden Glow = Korean ginseng, each capsule containing extract equivalent to 0.5 g *Panax ginseng* root and 8.93 mg ginsenosides as ginsenoside Rg1; 2 × 0.5-g capsules, t. i. d. = 3 g). Dosing of ginseng was continued for a further 1 week after warfarin administration. The bioanalytics comprised enantiomer-selective quantification of S-warfarin, which is predominantly metabolized to S-7-hydroxywarfarin by CYP2C9, and R-warfarin, which is metabolized by CYP3A4 and CYP1A2, thereby allowing for a separate mechanistic assessment of any potential alteration of these metabolic pathways by concomitant *P. ginseng* treatment. The study demonstrated that *P. ginseng* had no effect on the activity of CYP1A2, CYP3A4, or CYP2C9 in healthy subjects, as treatment did not affect the PK and clearance of both warfarin enantiomers in human subjects. *P. ginseng* also did not affect the apparent volumes of distribution or protein binding of warfarin enantiomers. PD endpoints of warfarin 25 mg single doses were also assessed in this study, and it was shown that *P. ginseng* did not significantly alter blood coagulation (i.e., INR) outcomes and platelet aggregation.

The results of this study were essentially confirmed by the same group of investigators in a warfarin interaction study of similar design, and in a comparable population (12 healthy male subjects, 6 Caucasian, 6 Asian), but using a different product (Blackmores Travel Calm Ginger, *Zingiber officinale*, each capsule containing extract equivalent to 0.4 g of ginger rhizome powder) at different daily doses (3 capsules, 3 × daily for 1 week) [26]. Similar as in the previous study, dosing of the herbal product was continued for a further week after warfarin administration. Also in this study, the PK and PD of warfarin were essentially unaltered by repeat-dose coadministration of recommended ginseng doses, thereby confirming the findings of the previous study [26]. Finally, the results of both studies [27, 75] were pooled and subjected to a population PK/PD modeling approach [74]. The pooled data analyses indicated that the ratio of S-warfarin's apparent clearance (CL/F) compared to control (= ratio warfarin + ginseng/warfarin alone) was 1.14 ± 0.04 after *P. ginseng* pretreatment, indicating that *P. ginseng* produced only a moderate increase in the CL/F of S-warfarin. None of the two ginseng products employed in the pooled studies had a direct effect on warfarin PD [74].

In another randomized, double-blind, placebo-controlled parallel group study in healthy adult subjects, the effects of AG (*P. quinquefolius*, 0.5 g capsules, total ginsenoside content of 5.19%) on the PK and PD of warfarin were studied [76]. Patients received oral warfarin (Coumadin, DuPont Pharmaceuticals), 5 mg daily, for the first 3 consecutive days during week 1. Beginning in week 2, patients were randomly assigned to receive either oral AG, 1.0 g, or placebo b. i. d. (= 2 g/day), for 3 consecutive weeks. During week 4, all patients again received oral warfarin, 5 mg daily, for the first 3 consecutive days. Ginseng or placebo assignment was randomized, and a total of 20 subjects were enrolled (9 m/11 f; 12 subjects in the ginseng group, 8 subjects in the placebo group). The authors compared changes in peak INR, INR AUC, peak plasma warfarin levels, and warfarin AUCs between the ginseng and placebo groups. The peak INR was shown to decrease statistically significantly after 2 weeks of ginseng administration compared with placebo (difference between ginseng and placebo, -0.19 [95% CI, -0.36 to -0.07]; $p = 0.0012$). The INR area

under the curve (AUC), peak plasma warfarin level, and warfarin AUC were also significantly reduced in the ginseng group as compared with the placebo group. Also, peak INR and peak plasma warfarin levels were positively correlated. The INR responses in the ginseng group were modest, albeit significantly different from the placebo group. The average group results, however, have to be interpreted with caution, as the INR data in the ginseng group were largely driven by one outlier patient displaying a high baseline INR (1.32) and a remarkably increased peak INR after warfarin administration on day 4. After ginseng administration, the peak INR of this patient declined remarkably from 5.16 to 2.75, and the corresponding AUC_{INR} decreased from 17.46 to 11.1. Although the authors emphasize that the results of their study remain statistically significant, when this patient is excluded from analysis, it becomes evident, that parallel group studies of such a small sample size are difficult to interpret.

Taken together, the available published evidence from clinical ginseng/warfarin interaction studies indicate that high repeated doses of *P. ginseng* and ginger over 1 week did not significantly alter the PK or PD of 25 mg warfarin single doses in healthy adult subjects [27, 74, 75]. In contrast, a small parallel group study in healthy subjects using a different warfarin posology (5 mg warfarin q. d. on 3 consecutive days) and a 2-week treatment with AG (*P. quinquefolius*) at the high end of the recommended dose range suggests that AG may reduce warfarin exposure (i.e., increases warfarin clearance), thereby reducing the anticoagulant effect of warfarin to a modest extent. The differences in the outcome of these studies might be either due to differential properties of *P. ginseng* and AG (*P. quinquefolius*) products, differences in the study designs (1-week vs. 2-week treatments), or simply due to the limitations of the small parallel group study which was apparently confounded by an outlier subject [76].

While all ginseng/warfarin interaction studies available so far suffer the principle design limitation of short-term warfarin treatments in healthy adult subject populations, which may not be predictive for long-term warfarin treatment in typical patient populations, the totality and consistency of data derived from mechanistic PK studies as detailed above, strongly suggest a lack of relevant CYP enzyme inhibitory or induction properties of ginseng products at recommended therapeutic doses.

As yet, no clinical studies on the investigation of ginseng products regarding their capacity to confer transporter-based interactions have been published. Therefore, the available published evidence from clinical trials suggests an overall low potential of virtually all ginseng species for CYP-based ginseng-drug interactions, while clinical studies on transporter-based interactions are still lacking.

Goldenseal

Goldenseal (*Hydrastis canadensis*, orangeroot) is a perennial herb in the buttercup family Ranunculaceae, indigenous to southeastern Canada and the northeastern United States. Botanical supplement goldenseal (GS) preparations are taken for the prevention and treatment of the common cold, upper respiratory and gastrointestinal infections, as well as menstrual disorders. In addition, GS is being used as a topical antimicrobial remedy to disinfect cuts and scrapes. GS contains a number of isoquinoline alkaloids such as hydrastine, berberine, berberastine, hydrastinine, tetrahydroberberastine, canadine, and canalidine. Hydrastine and berberine are the main active constituents [77].

As a supplement, standardized extracts of GS are available in liquid, tablet, and capsule forms. Recommended doses of the pow-

dered root vary widely and range from 750 mg to 6 grams in tablet or capsule form per day, to be given 2 to 3 times daily. GS preparations enjoy widespread popularity and are among the top 20 bestselling botanical products in the US [78].

A total of 8 published pharmacokinetic *in vivo* drug interactions with orally administered GS root extract (5 studies) or berberine products (3 studies) were identified.

In the first published study, the influence of GS root extract (GS root herbal single 570-mg capsules, two capsules b.i.d. = 2.280 mg/day, given for 14 days) on the disposition of the HIV protease inhibitor as well as CYP3A4 and P-gp substrate indinavir was examined in 11 healthy subjects (6 m/5 f) [79]. Indinavir was given as a single 800-mg oral dose; blood samples were collected for 8 hours postdose, and the PK of indinavir was characterized before and after 14 days of GS treatment. No statistically significant alterations in the PK of indinavir were observed (AUC values not reported). The oral clearance (CL/F) of indinavir was lowered on average by 4.6% following GS treatment, with individual changes ranging from -46.1% (subject 5) to a 46% increase (subject 4), indicating a remarkable intersubject variability in this study. Accordingly, the 95% CI for the difference in oral clearance between treatments was large (-10.6 to 3.4 L/h). Similar changes were observed in C_{max} , with the 95% CI for the difference in peak concentration with goldenseal root ranging from -0.6 to 2.5 mg/L. Overall, the study results suggest that GS supplementation at customary doses appears unlikely to affect the activity of CYP3A4 and/or P-gp (ABCB1) to a relevant extent *in vivo*, although the remarkable intersubject variability observed may pose questions regarding the robustness and reliability of the study. The authors also discussed the possibility of a limited study sensitivity regarding inhibition of presystemic CYP3A4, as indinavir was previously shown in a grapefruit juice interaction study to be a poor probe substrate for assessing CYP3A4-mediated gut wall extraction [80].

In a subsequent study, single timepoint, metabolic ratios of CYP probe drugs were used in 12 healthy subjects (6 m/6 f, extensive metabolizers of CYP2D6) to determine whether a 28 day supplementation of GS root extract (900 mg t.i.d. = 2.700 mg/day; no standardization claim) affects CYP1A2, CYP2D6, CYP2E1, or CYP3A4/5 activity [8]. Probe drug cocktails of MDZ (CYP3A4) and caffeine (CYP1A2), followed 24 hours later by chlorzoxazone (CYP2E1) and debrisoquine (CYP2D6) were administered before (baseline) and at the end of supplementation. Presupplementation and postsupplementation assessments of CYP3A4, CYP1A2, CYP2E1, and CYP2D6 activities were made by use of 1-OH-MDZ/MDZ serum ratios (1-hour sample), paraxanthine/caffeine serum ratios (6-hour sample), 6-hydroxychlorzoxazone/chlorzoxazone serum ratios (2-hour sample), and debrisoquine urinary recovery ratios (8-hour collection), respectively.

Comparisons of pre- and postsupplementation metabolic ratios indicated significant inhibition of about 40% of CYP2D6 [geom. mean post-/pre-ratio (90% CI) of 0.60 (0.51 to 0.70)] and CYP3A4/5 [geom. mean post-/pre ratios (90% CI) of 0.60 (0.53 to 0.67)] activities due to GS supplementation, while there was no appreciable effect on CYP1A2 and CYP2E1 activities. Hence, this was the first clinical study strongly suggesting that GS root extract, given at recommended doses, appeared to inhibit CYP2D6 and CYP3A4/5 to a notable extent *in vivo*.

For CYP3A4/5, the same group of authors confirmed this first clinical indication by a more comprehensive PK interaction study that comprised the assessment of complete plasma concentration-time profiles of the respective probe drugs [9]. In this study,

the effects of GS supplementation (GS root extract 1.323 mg t.i.d. = 3.969 mg/day, standardized to contain 24.1 mg isoquinoline alkaloids per capsule) on human CYP3A activity were evaluated using MDZ as a phenotypic probe. Sixteen healthy young subjects (8 m/8 f) were assigned to receive goldenseal for 14 days. Oral MDZ (8 mg) was administered before and after goldenseal supplementation, and serum concentrations of MDZ and the main metabolite 1-OH-MDZ were followed up to 6 hours postdose. In order to demonstrate the sensitivity of the study to adequately capture any relevant effects on CYP3A4 activity, the well-established CYP3A4 inducing/inhibiting probe drugs rifampin (300 mg b.i.d. for 7 days) and clarithromycin (500 mg b.i.d. daily for 7 days) were employed in the study as well. Comparisons of pre- and post-GS supplementation MDZ PK parameters indicated significant inhibition of CYP3A by GS ($AUC_{0-\infty}$ 107.9 ± 43.3 vs. 175.3 ± 74.8 ng·h/mL; CL/F/kg, 1.26 ± 0.59 vs. 0.81 ± 0.45 l/h/kg; $T_{1/2}$ 2.01 ± 0.42 vs. 3.15 ± 1.12 h; C_{max} 50.6 ± 26.9 vs. 71.2 ± 50.5 ng/mL). Comparisons of these data show that mean MDZ $AUC_{0-\infty}$ and C_{max} data were increased by about 63% and 40%, respectively, upon GS supplementation. In contrast, the potent CYP3A4 inhibitor clarithromycin produced about 5.5-fold (i.e., 548%) and about 2-fold (217%) increases in MDZ $AUC_{0-\infty}$ and C_{max} outcomes, respectively. Taken together, these data categorize GS root extract, at the given dose, as a weak inhibitor of CYP3A4 (i.e., <2-fold increase in AUC of the sensitive CYP3A4 probe MDZ), according to current FDA drug interaction guideline recommendations [81].

In a follow-up study addressing GS potential to inhibit CYP2D6, the effects of a 14-day treatment with standardized GS root extract (1.070 mg, standardized to contain 24.1 mg isoquinoline alkaloids per capsule, t.i.d. = 3.210 mg/day) on the activity of CYP2D6 was investigated in 16 healthy subjects (8 m/8 f), after administration of the CYP2D6 substrate debrisoquine (5 mg) before and at the end of supplementation, by using 8-h debrisoquine urinary recovery ratios [53]. Comparisons of pre- and postsupplementation ratios indicated a significant inhibition of CYP2D6 activity by GS supplementation by about 47% (post/pre geom. mean ratio 0.53, 90% CI 0.44 to 0.64), thereby confirming previously reported single-point phenotyping data [18]. Quantitatively, these data categorize GS root extract, at the given dose, as a weak inhibitor of CYP2D6 (i.e., <50% decrease in substrate clearance) according to current FDA drug interaction guideline recommendations.

To examine potential effects of GS root extract supplementation on P-gp (ABCB1) function *in vivo*, 20 healthy young subjects (10 m/10 f) were randomly assigned to receive a standardized GS supplement (1.070 mg t.i.d. = 3.210 mg/day) for 14 days or to receive the established positive control probe drugs rifampin (300 mg b.i.d. for 7 days) and clarithromycin (500 mg b.i.d. for 7 days). Supplementation/medication phases were separated by 30-day washout periods. Digoxin (Lanoxin™, 0.5 mg) was administered p.o. before and at the end of each supplementation/medication period, and serial digoxin serum concentrations were determined over 24 h postdose [82].

Statistically significant increases ($p < 0.001$) in digoxin total and maximum exposure parameters [AUC_{0-24} (57%), AUC_{0-3} (83%), C_{max} (95%)], and terminal disposition half-life (79%) were observed after 7 days of clarithromycin treatment. Accordingly, clarithromycin produced a 53% decrease in CL/F ($p < 0.001$). In turn, statistically significant reductions ($p < 0.05$) in digoxin AUC_{0-24} (-16%), AUC_{0-3} (-27%), and C_{max} (-28%) were noted following rifampin administration. Taken together, the results of

the P-gp probe inhibitor/inducer indicate adequate sensitivity of the study to accurately capture relevant effects on P-gp function. As a result of GS supplementation, there were no significant changes in digoxin PK observed, apart from a 14% increase in digoxin C_{max} ($p < 0.05$). Based on the results of this study, it can be concluded that GS root extract supplementation at daily doses of about 3 g does not modify P-gp function to a clinically relevant extent.

Two studies examined the effect of berberine, a major alkaloid constituent in GS extract, on the PK of cyclosporine A (CsA) in Chinese renal transplant patients [83] and in healthy adult Chinese subjects [84].

In a randomized, controlled clinical trial in 104 renal transplant recipients, Wu and colleagues examined the effects of oral berberine supplementation [BBR hydrochloride tablets 200 mg (unspecified source), t.i.d. for 3 months] on the PK of CsA [83]. Patients were randomly assigned in a 1:1 fashion to a BBR-free group or to the BBR treatment group. Blood trough concentration of CsA and markers for hepatic and renal functions were determined. For a more comprehensive PK substudy, six renal transplant recipients were treated with a 3-mg/kg dose of CsA b.i.d. before and after oral coadministration of 200 mg BBR t.i.d. for 12 days.

In the clinical study, the final CsA trough concentrations and the ratios of concentration/dose of CsA in BBR-treated patients were modestly increased, i.e., 29.3% and 27.8% higher than those in BBR-free patients, respectively ($p < 0.05$). In the dedicated PK study in 6 patients, the average increase in the total CsA exposure (AUC) upon BBR treatment was 34.5% ($p < 0.05$), while average increases in C_{ss} and C_{min} values of CsA were 34.5% and 88.3%, respectively, without significant alterations in C_{max} . The average decrease in CL/F was 40.4%. Overall, the study results suggest a weak inhibition of CYP3A4 by BBR supplementation at a 600 mg/day dose level.

In the second BBR/CsA interaction study, the effects of 300 mg BBR, b.i.d. (= 600 mg/day), given for 10 days on the pharmacokinetics of CsA were examined in 12 healthy male Chinese subjects. CsA PK assessments at a dose of 6 mg/kg were done in 6 subjects both before and at the end of the BBR treatment period, whereby the second CsA dose was given ≥ 12 h after the last BBR dose. Another 6 subjects received 3 mg CsA/kg, the second CsA dose being given concomitantly with a single oral dose of 300 mg BBR. In the study with 6 mg CsA/kg treatment and time-separated intake of the last BBR dose, BBR did not significantly alter the PK of CsA. However, in the trial with 3 mg CsA/kg and the concomitant administration of CsA and BBR, a modest average increase in the AUC of CsA of 19.2% ($p < 0.05$) was observed, without any accompanying changes in the terminal disposition half-life or apparent oral clearance (CL/F) of CsA. These results indicate that BBR can modestly increase exposure of CsA, when both drugs are administered together [84].

Taken together, although not entirely consistently, both BBR/CsA studies are suggestive for a modest BBR-mediated inhibition of CYP3A4, which may need to be considered in case of concomitant treatment with narrow therapeutic index drugs that are sensitive substrates of CYP3A4, such as CsA.

In the most recent study, 18 Chinese healthy male adult subjects received orally either placebo or berberine (BBR) capsules at a dose of 300 mg t.i.d. (= 900 mg/day) for 14 days, and phenotypic effects of BBR supplementation on the activities of CYP1A2, CYP2C9, CYP2C19, CYP2D6, and CYP3A4 were examined by the administration of the probe substances caffeine (93 mg), losartan

(30 mg), omeprazole (20 mg), dextromethorphan (30 mg), and MDZ (2 mg), respectively [10]. The PK profiles of omeprazole, MDZ, and caffeine were assessed over 12 h postdose, and the losartan/E-3174 as well as dextromethorphan/dextrorphan urinary ratios were assessed by urine sampling over 8 h postdose.

A remarkable decrease in CYP2D6 activity was observed in this study as the 0–8 h urinary dextromethorphan/dextrorphan ratio increased about 9-fold ($p < 0.01$). In addition, the losartan/E-3174 ratio doubled ($p < 0.01$) after BBR administration, indicating a notable decrease in CYP2C9 activity. In contrast, CYP3A4 activity was only modestly inhibited, as the C_{max} , $AUC_{0-\infty}$, and AUC_{0-12} of MDZ were increased 38% ($p < 0.05$), 40% ($p < 0.01$), and 37% ($p < 0.05$), respectively, after BBR treatment. Accordingly, the oral MDZ clearance was also modestly decreased by 27% ($p < 0.05$). There were no significant alterations of the probe substances caffeine (CYP1A2) and omeprazole (CYP2C19). Taken together, repeated BBR administration (900 mg/day) decreases CYP2D6 activity significantly, appears to modestly modify 2C9 and CYP3A4 activities, and does not alter CYP1A2 and CYP2C19 function.

In conclusion, 5 pharmacokinetic interaction studies with different GS root extract products at daily doses up to 4 g are overall largely consistent and characterize GS root extract as a weak inhibitor of CYP3A4 and CYP2D6 enzymes, when FDA drug interaction guideline criteria are considered. The available evidence for this assessment is most robust for CYP3A4, and based on phenotypic serum ratios or metabolite urinary recovery data only, for CYP2D6. Results from single studies indicate that GS root extract does not alter the activities of CYP1A2 and CYP2E1 enzymes, and the efflux transporter P-gp (ABCB1) to an appreciable extent. No information is available regarding potential effects of GS on CYP2C9 and CYP2C19 enzymes.

The available *in vivo* evidence on the PK drug interaction potential of BBR products is derived from 3 studies, altogether conducted in Chinese subjects. These studies consistently point to a weak potential of BBR products to inhibit CYP3A4. In addition, a single phenotyping study suggests that BBR decreases CYP2D6 activity and modestly modifies CYP2C9 function, but is unlikely to alter CYP1A2 and CYP2C19 functions. These findings and their clinical relevance, however, need to be confirmed by future studies. So far, caution should be applied when GS root extract or BBR products are coadministered with narrow therapeutic index drugs, which are sensitive substrates of CYP3A4, CYP2D6 or CYP2C9 enzymes.

Garlic

Garlic (*Allium sativum*; family Amaryllidaceae) has been cultivated and used for the alleviation of a variety of medical conditions for thousands of years. Today, garlic is worldwide still one of the most popular and most commonly used herbal supplements with purported antioxidative, antibacterial, antiparasitic, antilipidemic, antihypertensive, antiatherosclerotic, antithrombotic, antidiabetic, immune-stimulating and anticancer properties [85–87].

Pharmaceutical products derived from garlic bulbs/cloves are provided in the forms of powders, oily preparations, or aqueous-alcoholic extracts of fresh or aged garlic. Due to the complex chemistry, differing manufacturing processes result in preparations of quite different chemical composition and pharmaceutical properties, which in turn may result in different pharmacological, safety, and bioavailability characteristics [12, 88].

The so far identified active constituents of garlic include alliin, allinase, diallyl disulfide, ajoens, and others. Alliin is enzymatically

converted to allicin, the major garlic component, which, however, displays considerable chemical instability and is rapidly degraded to a variety of organosulfur compounds (OSCs such as diallyl sulfide, diallyl disulfide, diallyl trisulfide) that are believed to ultimately confer the main pharmacological effects [89].

Recommended doses range around 4 g of fresh garlic daily, which is equivalent to approximately 8 mg garlic oil or 600 to 900 mg garlic powder preparations standardized to 1.3% alliin content [89].

Despite the widespread use of garlic preparations along with reported preclinical evidence and anecdotal clinical case reports on the potential role of garlic in metabolism-based PK interactions and also some susceptibility for PD interactions, well-designed clinical studies evaluating drug interactions with garlic preparations in a systematic fashion are notably very limited.

Nonclinical studies suggest that garlic components may exert enzyme inducing effects on CYP1A1/2 and CYP2B1/2 family members via CAR and Nrf2 activation [90], and also to strongly induce GST activity [91], whereby upregulation of GST has been implicated in the detoxification of carcinogens. In turn, inhibition of CYP2E1, CYP2C9, and 2C19 by garlic and QSCs was suggested by preclinical data [69].

Unfortunately, the clinical relevance of these findings for commonly employed human doses has not been completely followed up as yet by well-designed mechanistic *in vivo* studies for many of the described metabolic pathways. Hence, based on currently published evidence, the question of clinical relevance for the described effects on CYP2B1/2 family members and UGTs, GST, QR, and EH enzymes remains virtually still unaddressed to the best of our knowledge.

A total of 8 studies in healthy adult young and elderly subjects using different garlic products (600 mg garlic tablets, 3 tablets b.i.d. [92]; garlic oil 500 mg, t.i.d. [18, 19], garlic oil extract, diallyl sulphide (DAS) [93]) and study designs have so far addressed various aspects of the PK drug interaction potential of garlic treatment. Apparently, the most recent human *in vivo* study addressing garlic-related PK drug interaction issues was published in 2005, thereby indicating some stagnation rather than ramp-up of clinical activities in this area of research.

A total of 3 interaction studies with administration of prototypic enzyme-specific probe drugs assessed the potential of various garlic preparations to alter activities of CYP2E1 [93], CYP2D6 and 3A4 [92], CYP1A2, CYP2D6, CYP2E1, or CYP3A4 [17] in healthy subjects, and one study used a similar approach to probe for alterations of CYP1A2, CYP2D6, CYP2E1, and CYP3A4 activities in elderly individuals [18].

These studies indicate consistently that neither single nor repeat dose administrations of various garlic formulations up to 28 days may have potential to exert notable alterations (i.e., inhibition or induction) of CYP1A2, CYP2D6, or CYP3A4 activities in humans, which are important metabolic pathways for many drugs likely to be coadministered with garlic supplements. In clinical terms, these results indicate that garlic preparations are unlikely to alter the disposition of coadministered drugs that are sensitive substrates of these oxidative metabolic pathways.

In contrast, repeat administration of garlic oil over 28 days was shown to modestly reduce CYP2E1 activity [estimated by 6-hydroxychlorzoxazone (6-OH-CZX)/chlorzoxazone (CZX) metabolic ratio] by 39% in healthy adult subjects [17] and by approximately 22% in elderly individuals (mean age 67 years) [18]. These results are consistent with an earlier study on single dose effects of 0.2 mg/kg BW garlic oil extract, diallyl sulphide (DAS), on CYP2E1

activity, which also employed the 6-OH-CZX/CZX metabolic ratio for the estimation of CYP2E1 function [93]. However, as the overall extent of CYP2E1 inhibition by garlic is modest, i.e., conceptually metabolic ratios always display quantitatively greater changes than respective clearance alterations of the prototypic substrate, and hardly any currently approved drugs in widespread use are relying on CYP2E1 as a sole or predominant clearance pathway (except chlorzoxazone), the observed CYP2E1 effects of garlic do not appear to have important clinical implications. This perspective is further supported by a study in healthy adult subjects, in which a 3-month administration of aged garlic extract (10 mL q.d.) did not show any notable effect on oxidative metabolism of acetaminophen, to which CYP2E1 is known to be a significant contributor, besides CYP3A4 and CYP1A2 [94]. Interestingly, in this study also no effects of garlic on GST activity could be noted, as no changes in the excretion of three acetaminophen metabolites arising from GST-mediated glutathione conjugation with the reactive acetaminophen metabolite were observed [94]. This finding casts doubts on the human *in vivo* relevance of the consistently reported inducing effects of garlic components on GST activity from animal and *in vitro* studies.

The lack of a 3-week garlic treatment (GarliPure™ 1.200 mg/day) on the functional activity of intestinal and hepatic CYP3A4 in healthy adult subjects was further confirmed by demonstration of an unaltered PK of the HIV-protease inhibitor saquinavir, a sensitive CYP3A4 substrate which undergoes extensive CYP3A4-mediated presystemic gut wall metabolism, and by means of an erythromycin breath test, a noninvasive marker of hepatic CYP3A4 function [95].

These findings are also in line with another healthy subject study demonstrating that a 4-day short-term treatment with garlic capsules (10 mg garlic extract q.d., provided as 2 × 5-mg liquid-filled soft gelatin capsules, equivalent to 1 g of fresh garlic) did not significantly alter the single dose PK of ritonavir, another HIV-protease inhibitor that is mainly metabolized by CYP3A4. However, the authors cautioned that in view of the complex time-dependent effects of ritonavir on drug metabolizing enzymes and transporters (ritonavir is also a potent inhibitor of CYP3A4 and the ABC efflux transporter P-gp), the results of the study may be difficult to interpret and should not be extrapolated to steady-state conditions [96].

The only study reporting a PK alteration of the HIV-protease inhibitor saquinavir upon a 3-week garlic treatment (GarliPure™ caplets b.i.d.) of ten healthy subjects suggested about 51%, 49%, and 54% reduced mean saquinavir steady-state exposure (AUC_τ), trough levels (C_{trough}), and maximum concentrations (C_{max}), respectively [55]. However, most notably, after a 10-day washout period of garlic treatment, the saquinavir AUC, trough, and C_{max} values did not return to baseline, but still remained at 60–70% of their baseline values. As the study employed a fixed sequence rather than a randomized crossover design, and in light of published evidence on time-dependent saquinavir PK characteristics in healthy subjects [97] and HIV patients [98], it is most conceivable that the study results merely mirror saquinavir-mediated induction processes of its own metabolism and disposition rather than indicating a relevant garlic-mediated drug interaction.

Taken together, there is well-consistent evidence from a variety of mechanistic PK studies that common doses of various kinds of garlic preparations do not exert any notable alteration of CYP3A4 function *in vivo*.

Whether or not garlic may be involved in transporter-based drug interactions has apparently not yet been addressed in mechanis-

tic human PK studies, to the best of our knowledge. Also the pre-clinical evidence in this respect is less well documented [69].

Taken together, from the limited available evidence of human *in vivo* drug interaction studies, there is currently no indication that garlic may confer any notable or clinically significant metabolism-based PK interactions with drugs cleared by CYP1A2, CYP2D6, CYP2E1, or CYP3A4 enzymes. In contrast, unfortunately, no published evidence from human *in vivo* studies is currently available for the characterization of the potential effects of garlic on CYP2B family members and UGT, GST, QR, and EH enzymes or drug transporter proteins. Hence, translation of published *in vitro* findings into clinical interaction studies is still largely incomplete.

Echinacea

Echinacea is a genus of herbaceous flowering plants (purple coneflowers, family Asteraceae), overall comprising nine species, which are native and endemic to eastern and central North America. Marketed and studied medicinal products are made of three different species (*E. purpurea*, *E. angustifolia*, *E. pallida*), whereby the majority (i.e., about 80%) of the products are based on *E. purpurea*. Different parts of the plants (aerial parts and roots) are used, and different pharmaceutical preparations exist (extracts and expressed juice). The German Monograph issued by the E Commission recommends alcoholic extracts from the root of *E. pallida* and juices pressed from the aerial parts of the plants [99]. However, the diversity in the overall composition and contents of pharmaceutical *Echinacea* products is considerable and may contribute to some across study inconsistencies.

Echinacea is considered to be an unspecific immunostimulant and is primarily used for the prevention and treatment of acute virus infections of the upper respiratory tract, such as the common cold and influenza. Dose recommendations for *Echinacea* products vary widely and depend on the formulation and concentration of the product. Recommended daily doses for adult subjects range between 900 to 1000 mg, to be given in 3 to 4 divided doses.

Despite the popularity and widespread use of *Echinacea* preparations and their pharmaceutical diversity, well-designed clinical studies evaluating PK drug interactions with different *Echinacea* preparations are notably very limited. Overall, we identified only 7 clinical PK interaction studies of *Echinacea* products in the public domain, which primarily addressed CYP- and P-gp-based interaction issues.

The effect of *Echinacea* (*E. purpurea* root) on CYP activities *in vivo* was assessed by use of single dose administrations of the CYP probe drugs caffeine (CYP1A2), tolbutamide (CYP2C9), dextromethorphan (CYP2D6), and midazolam (CYP3A) in 12 healthy subjects (6 m/6 f) [100]. The probe substances caffeine (200 mg), tolbutamide (500 mg), dextromethorphan (30 mg), and oral and intravenous midazolam (MDZ; 0.05 mg/kg BW i.v., and 5 mg per oral) were administered before and after a short course of *Echinacea* [*E. purpurea* root extract 400 mg 4 times daily (= 1600 mg/day) for 8 days], and respective plasma concentration-time profiles of the probe substances were assessed.

The disposition (i.e., $AUC_{0-\infty}$, oral clearance, $t_{1/2}$) of the CYP2D6 probe dextromethorphan in CYP2D6 genotyped subjects (11 CYP2D6 EMs, 1 CYP2D6 PM) was not significantly altered by *Echinacea*. Likewise, the ratio of the dextromethorphan/dextrorphan $AUC_{0-\infty}$ values and the dextromethorphan-to-dextrorphan urinary metabolic ratio were not significantly affected by *Echinacea* coadministration.

The oral clearance of the CYP2C9 probe drug tolbutamide was significantly, albeit modestly, reduced on average by 11% after *Echinacea* dosing, with two individuals (1 m/1 f) displaying a 25% or greater reduction in the oral clearance of tolbutamide. In line with this, the geometric mean total tolbutamide exposure ($AUC_{0-\infty}$) before and after *Echinacea* dosing was modestly increased by 14% (90% CI 9–20%). The maximum tolbutamide concentration was not significantly altered by *Echinacea*.

Echinacea dosing significantly reduced the mean oral clearance of the CYP1A2 probe caffeine by 27%, with considerable intersubject variability (> 50% in 2 individuals). Maximum serum concentrations of caffeine after a 200-mg oral dose were on average increased by 30%, and the geometric mean total caffeine exposure ($AUC_{0-\infty}$) was increased by 29% after *Echinacea* dosing.

The systemic clearance of intravenously administered CYP3A4 probe MDZ was significantly increased by 42%, and there was a corresponding significant decrease in mean $AUC_{0-\infty}$ of 23%, together suggesting that hepatic CYP3A activity was increased. In contrast, the oral clearance and the $AUC_{0-\infty}$ after oral MDZ dosing were not significantly altered by *Echinacea* administration. However, the mean absolute oral bioavailability (F) of MDZ was increased from 24% to 36% (i.e., by 50%), consistent with the inhibition of intestinal CYP3A by *Echinacea*.

The authors discussed that the differential effects of *Echinacea* on intestinal and hepatic CYP3A activity may be due to a variety of possible mechanisms such as locally acting CYP3A inhibiting constituents of *Echinacea* that do not become systemically available, or rapid absorption of CYP3A4 inducing components of *Echinacea*, thereby limiting intestinal exposure and intestinal CYP3A induction, or a systemically formed metabolite of a constituent of *Echinacea* capable of inducing hepatic CYP3A but not intestinal CYP3A. However, these mechanistic considerations remain speculative and deserve further investigation.

Taken together, the study showed that an 8-day treatment with relatively high doses of *E. purpurea* root extract (i.e., 400 mg 4 times daily = 1600 mg/day) displayed differential effects on the activity of various CYP enzymes. There were no alterations of CYP2D6 noted, negligible to modest inhibition of CYP2C9, modest inhibition of CYP1A2, and differential (i.e., inductive/inhibitory) effects on hepatic and intestinal CYP3A4.

Although the observed mean effects on CYP1A2 and CYP2C9 activities were generally modest, they displayed some more pronounced effect sizes in individual subjects. Considering CYP-based *Echinacea*-drug interactions of potential clinical significance, the study suggests that the bioavailability of orally administered CYP3A4 substrates with low oral bioavailability may be significantly increased by *Echinacea* coadministration, and that the exposure to CYP1A2 and 2C9 substrates (e.g., theophylline, clozapin) may be modestly increased, at least in individual subjects. The observed effects of high doses of *E. purpurea* root extract on CYP1A2 and CYP2C9, however, are overall small and therefore unlikely to be of clinical significance in clinical practice, unless substrates with a narrow therapeutic index are concerned [e.g., (S)-warfarin in case of CYP2C9 and theophylline in case of CYP1A2].

In another study, 12 healthy subjects (6 m/6 f) were randomly assigned to receive *E. purpurea* (800 mg b.i.d., no standardization claim) for 28 days [50]. Probe drug cocktails of MDZ (CYP3A4) and caffeine (CYP1A2), followed 24 hours later by chlorzoxazone (CYP2E1) and debrisoquine (CYP2D6) were administered before (baseline) and at the end of supplementation. Presupplementation and postsupplementation assessments of CYP3A4, CYP1A2,

CYP2E1, and CYP2D6 activities were made by use of 1-OH-MDZ/MDZ serum ratios (1-hour sample), paraxanthine/caffeine serum ratios (6-hour sample), 6-hydroxychlorzoxazone/chlorzoxazone serum ratios (2-hour sample), and debrisoquine urinary recovery ratios (8-hour collection), respectively.

In this study, high-dose *E. purpurea* given over 28 days did not significantly change the activities of CYP3A4, CYP2E1, and CYP2D6 as estimated by single timepoint metabolic ratios. The only perceptible change in mean phenotypic ratios occurred with 6-hour paraxanthine/caffeine values, where an approximate 13% decrease (from 0.514 ± 0.181 to 0.447 ± 0.190) pointed to a possible inhibitory effect of *E. purpurea* on CYP1A2. This minor difference, however, was not statistically significant ($p = 0.07$), nor was it considered clinically relevant.

The effects of supplementation of a standardized *E. purpurea* extract (267 mg t.i.d. = 801 mg/day, each capsule standardized to contain 2.2 mg isobutylamides; Gaia Herbs, Inc.) on the activity of CYP2D6 was investigated in 16 healthy subjects (8 m/8 f) after administration of the CYP2D6 substrate debrisoquine (5 mg) before and at the end of supplementation, by using 8-h debrisoquine urinary recovery ratios [53]. Respective results did not indicate significant inhibition of CYP2D6 by *E. purpurea* supplementation.

Two studies examined the effects of *Echinacea* treatment of the PK of two different protease inhibitor/ritonavir combination products. The most recent study was conducted in 15 HIV-infected patients (14 m/1 f, median age 49 yrs, range 43 to 67 yrs) receiving antiretroviral therapy including darunavir-ritonavir (600/100 mg b.i.d.) for at least 4 weeks. The patients were concomitantly treated with *E. purpurea* root extract capsules (500 mg every 6 h) for 14 days on top of their antiretroviral treatment [101]. The HIV protease inhibitor darunavir undergoes extensive metabolism by CYP3A4, while ritonavir is a potent inhibitor of CYP3A4 and P-gp, and is combined with darunavir to boost (i.e., increase) its systemic exposure. Darunavir and ritonavir plasma concentrations were determined up to 12 h after a morning dose of darunavir-ritonavir on days 0 (darunavir-ritonavir alone) and 14 (darunavir-ritonavir plus *Echinacea*). Mean ritonavir PK parameters were essentially unchanged by coadministration of *E. purpurea*, while small changes in average darunavir PK parameters were noted, with trough concentrations (C_T) and measures of total exposure over a complete dosing interval (AUC_T) decreasing on average by 16% and 10%, respectively. However, individual patients displayed declines in darunavir C_T or AUC_T by as much as 40% and 30%, respectively, while maximum observed darunavir concentrations (C_{max}) were essentially unchanged by *E. purpurea* coadministration. Overall, the study indicates that high-dose *E. purpurea* coadministration does not alter the PK of low dose ritonavir and has on average little effect on the exposure characteristics of darunavir. It is to be noted, however, that studies with ritonavir containing anti-HIV combination products provide little assay sensitivity to capture any *Echinacea*-mediated effects on CYP3A4 activity because ritonavir, as a potent CYP3A4 inhibitor, is likely to mask any less potent influence of *Echinacea* on CYP3A4 activity. However, the finding that darunavir exposure was reduced in individual patients by 30 to 40% upon *E. purpurea* coadministration, would be consistent with the assumption of a modest induction of hepatic CYP3A4 by *Echinacea* administration as previously reported by Gorski and colleagues [50]. This finding would be further compatible with the hypothesis, that low-dose ritonavir (100 mg b.i.d.) may

not effectively inhibit hepatic CYP3A4 activity in individual patients.

In a comparable study in thirteen healthy subjects (8 m/5 f) receiving a combination of lopinavir/ritonavir 400/100 mg b.i.d. for 29.5 days, subjects were treated concomitantly with *E. purpurea* (500 mg t.i.d.) for 28 days: 14 days in combination with lopinavir/ritonavir and 14 days of *E. purpurea* alone [102]. Lopinavir and ritonavir plasma concentrations were determined at the end of each treatment period up to 12 h postdose. In order to assess CYP3A and P-gp activity, subjects also received single oral doses of the probe drugs MDZ 8 mg (CYP3A4) and fexofenadine 120 mg (P-gp/OATP2B1), respectively, before and after the 28 days of *E. purpurea*. In this study, neither lopinavir nor ritonavir PK were significantly altered by 14 days of *E. purpurea* coadministration. The post-*Echinacea*:pre-*Echinacea* geometric mean ratios (GMRs) for the lopinavir AUC_{0-12h} and for maximum concentrations (C_{max}) were 0.96 (90% CI 0.83–1.10, $p = 0.82$) and 1.00 (90% CI 0.88–1.12, $p = 0.72$), respectively. Conversely, GMRs for the total exposure of MDZ ($AUC_{0-\infty}$) and oral MDZ clearance were 0.73 (90% CI 0.61–0.85, $p = 0.008$) and 1.37 (90% CI 1.10–1.63, $p = 0.02$), respectively, thereby suggesting the induction of CYP3A4-mediated MDZ clearance. Fexofenadine PK, in contrast, did not significantly differ before and after *E. purpurea* administration ($p > 0.05$).

Overall, the study outcome regarding the observed changes in CYP3A4 activity is consistent with the results reported by Moltó for the effects of *E. purpurea* supplementation on a darunavir-ritonavir combination [101]. Both studies consistently indicate that *E. purpurea* coadministration is unlikely to significantly alter the PK of ritonavir-boosted protease inhibitors, most likely due to the presence of the potent CYP3A inhibitor ritonavir, but still may be capable to modestly induce hepatic CYP3A activity. The lack of any effects of *E. purpurea* treatment on the PK of fexofenadine – a substrate of several drug transporters including P-gp, OATP1A2, OATP1B1, and OATP2B1 – in the presence of low-dose ritonavir administration is difficult to interpret, because ritonavir is also a potent inhibitor of P-gp and hence, may mask any modest *Echinacea*-mediated effects on P-gp (ABCB1).

Abdul et al. investigated the PK and PD interactions of *Echinacea* with warfarin in 12 healthy male subjects who received a single oral 25 mg dose of warfarin alone and after 2 weeks of pretreatment with high-dose *Echinacea* (1275 mg four times daily = 5100 mg/day, containing a mixture of 600 mg of *E. angustifolia* roots and 675 mg of *E. purpurea* root; standardized to contain 5.75 mg of total alkylamides per tablet) [103]. In this study, the apparent clearance of (S)-warfarin was found slightly increased by 9% (geom. mean ratio 1.09; 90% CI of ratio 1.01, 1.18) during concomitant *Echinacea* treatment, a negligible change that achieved, however, statistical significance. Accordingly, the total exposure (i.e., $AUC_{0-\infty}$) to (S)-warfarin was increased by 9% [geom. mean ratio 1.09 (1.01, 1.18)]. The exposure characteristics of (R)-warfarin were not changed significantly by *Echinacea*. Similarly, the PD endpoints were not altered significantly by *Echinacea* with geometric mean ratios of 1.04 (0.95, 1.13) and 1.09 (0.91, 1.31) for INR_{max} and AUC_{INR} , respectively. As (S)-warfarin is metabolized by CYP2C9 and CYP3A4, and (R)-warfarin is metabolized by CYP3A4 and CYP1A2 [104], it can be concluded that a 2-week treatment with high doses of *Echinacea* (i.e., 5100 mg/day) does not substantially alter the activities of these CYP enzymes. Overall, the small effect sizes observed are unlikely to be of clinical relevance.

The effects of *Echinacea* supplementation on P-gp function was studied in 18 healthy subjects, randomly assigned to receive a standardized *Echinacea* supplement (267 mg t.i.d. = 801 mg/day) for 14 days [105]. Subjects were also randomized to receive rifampin (300 mg b.i.d., 7 days) and clarithromycin (500 mg b.i.d., 7 days) as positive controls for P-gp induction and inhibition, respectively. Digoxin (Lanoxin™ 0.25 mg), a well-acknowledged P-gp probe, was administered orally before and after each supplementation and control period. Serial digoxin plasma concentrations were obtained over 24 h. Comparisons of area under the curve values (AUC_{0-3} , AUC_{0-24}), terminal disposition half-life ($t_{1/2}$), and maximum serum concentrations (C_{max}) were used to assess the effects of *Echinacea*, rifampin, and clarithromycin on digoxin disposition. As expected, rifampin produced significant reductions ($p < 0.05$) in AUC_{0-3} , AUC_{0-24} , and C_{max} , while clarithromycin, a known P-gp (ABCB1) inhibitor, increased these parameters significantly ($p < 0.05$). *Echinacea* supplementation, in contrast, did not affect digoxin PK, indicating that clinically significant P-gp-mediated herb-drug interactions with *Echinacea* are unlikely to occur.

In conclusion, the available clinical evidence indicates that *Echinacea* products at recommended doses do have little potential to cause clinically relevant or worrisome metabolism- and transporter-based PK interactions that involve CYP1A2, CYP2C9, CYP2D6, CYP2E1, and P-gp. In turn, there is consistent clinical evidence suggesting that *Echinacea* products actually do have potential to modestly induce hepatic CYP3A4 activity, but at the same time also may inhibit the presystemic metabolism of CYP3A4 substrates in the intestine. As both mechanisms in principle counteract each other in terms of the net effects on the systemic exposure of CYP3A4 substrates, may require different times for their onset (i.e., onset of inhibition occurs faster than onset of induction), and also will depend in their effect size on specific substrate characteristics (e.g., oral bioavailability of CYP3A4 substrates), general predictions on the clinical relevance of *Echinacea*/CYP3A4 substrate interactions are difficult to make. However, caution is advised when CYP3A4 substrates with low oral bioavailability due to pronounced intestinal CYP3A4-mediated metabolism such as verapamil, cyclosporine, or tacrolimus or narrow therapeutic index drugs predominantly cleared by CYP3A4 are coadministered with *Echinacea* supplementation.

Evidence from two phenotyping studies employing daily *Echinacea* doses of 1600 mg [50, 100] suggests the potential of a weak inhibitory effect of high-dose *Echinacea* supplementation on CYP1A2 and 2C9. However, reported effect sizes are overall small and therefore unlikely to be of clinical significance in clinical practice, unless substrates with a narrow therapeutic index are concerned, such as warfarin (CYP2C9) or theophylline (CYP1A2). No clinical studies are available addressing the potential impact of *Echinacea* products on other important metabolic pathways such as CYP2C19 or phase II metabolism (e.g., glucuronidation), and no clinical studies on potential transporter interactions other than P-gp (ABCB1) are available as yet.

Summary and Conclusions of Herb-Drug Interactions in Humans

Our review shows that the total number of clinical herb-drug interaction trials with primary PK objectives has increased considerably over the last years. A systematic review of clinical herb-drug interaction studies published in 2005 [106] identified a total

of 18 studies only, reporting original drug interaction data on ginkgo (5 studies), garlic (5), ginseng (3), milk thistle (3), golden-seal and *Echinacea* (each 1). This comparison indicates that the total number of PK herb-drug interaction studies referring to these six particular herbs has increased by more than 3.5-fold (i.e., from 18 to 66) over the last 7 years. Major progress since 2005 has been made for ginkgo (5 vs. 21 studies), milk thistle (3 vs. 13 studies), goldenseal (1 vs. 8 studies), ginseng (3 vs. 9 studies), and *Echinacea* (1 vs. 7 studies), while only 3 additional studies on garlic were published (5 vs. 8). Emerging contributions of Chinese working groups to this recent development need to be acknowledged in this context. It is also important to note that not just the mere number of studies has increased, but that the overall quality in study design (e.g., employed doses, measures to assess and confirm study sensitivity) and analysis points to a trend towards conceptual improvements.

It is worthy to note that this and virtually all previous reviews on PK-based herb-drug interactions only refer to herbal drugs as perpetrators of PK drug interactions. However, herbal drugs can also be objects of drug interactions (i.e., their exposure can be altered by coadministered perpetrator pharmaceuticals), a mechanism that may be of considerable importance for triggering or aggravating adverse events or secondary PD interactions. It is likely that this particular aspect of herb-drug interactions has not become part of the respective reviews as yet, because of the general paucity of published data. Such studies would require identification of the major active constituents in herbal products along with the establishment of sensitive bioanalytical assays to quantify alterations of the exposure characteristics of herbal drugs due to concomitantly administered perpetrator (i.e., inducer/inhibitors) drugs. As so far significantly underrepresented, the issue of herbal drug products as objects of PK-based drug interactions clearly represents an important field of future work.

Despite the progress that has been made over the last years, the overall number of clinical herb-drug interaction studies appears still limited, at least for some of the herbal products reviewed. Interactions studies with warfarin, a narrow therapeutic index drug and a probe substrate for CYP2C9, are numerous. However, studies with other narrow therapeutic index or critical dose drugs are sparse, except for CsA and antiretroviral drugs. With regards to mechanistic interaction studies, few major CYP enzymes such as CYP1A2, CYP2D6, and CYP3A4 have been well investigated, however the interaction potential of herbal drugs with members of the CYP2C family appears less well established, and we identified only one study addressing a CYP2B family enzyme (CYP2B6) [31]. In this context, however, it is probably also worthy to note that inducibility and susceptibility to inhibition are not of equal clinical relevance for the major CYP enzymes in drug metabolism. CYP2D6 for instance, is hardly inducible at all (i.e., no inducers of CYP2D6 are known), while in turn no potent or moderate *in vivo* inhibitors of CYP2B6 have been identified as yet [81]. Hence, *in vivo* studies with herbal products on the inducibility of CYP2D6 and the inhibition of CYP2B6 enzymes are in principle dispensable or at least of secondary importance. In turn, inducibility of CYP1A2, CYP2B6, and CYP3A should always be investigated, when *in vitro* data indicate induction potential of an herbal product. Systems-pharmacological knowledge on co-regulation of CYP enzymes (e.g., coinduction of CYP3A and CYP2C enzymes by the pregnane X receptor) or CYP enzymes and transport proteins (e.g., CYP3A and ABCB1) should also be considered to guide and prioritize smart and efficient *in vivo* drug interaction studies or programs.

Studies on the effects of herbal drugs on other important drug metabolizing enzymes, e.g., involved in phase II metabolic reactions (e.g., UGTs, N-acetyltransferases, etc.) are virtually lacking, further highlighting the need for more well-designed clinical trials that will increase our understanding about the underlying mechanisms of herb-drug interactions.

With the exception of ABCB1 and OATP2B1, there is very little information on herb-drug interaction trials investigating the effect on drugs that are mainly absorbed, distributed, or cleared by drug transporters. Given the evidence of clinically relevant drug-drug interactions involving these transporters, there is a still existing need to investigate whether herbal drugs could act as potential perpetrators of these important players in drug absorption, disposition, and elimination.

In light of the limited predictability of herb-drug interactions on the basis of *in vitro* studies, simultaneous administration of a mixture of substrates of multiple CYP enzymes and transporters in one study (i.e., a “cocktail approach”) in human subjects is an attractive approach to evaluate an herbal drug’s inhibition or induction potential [107]. There are phenotyping drug cocktails available that are scientifically sound and represent a valuable, safe, and cost-effective tool specifically suited to screen for metabolic and P-gp (ABCB1)-mediated perpetrator interactions of herbal drugs. In case of cocktail or other herb-drug interaction studies relying on the assessment of metabolic ratios, two aspects need to be considered in the assessment of study outcomes: first, urinary metabolic ratios often have a much weaker correlation to enzyme activity and are also not closely correlated to plasma-based metrics. Second, changes in metabolic ratios (regardless of the matrix/compartiment concerned, i.e., serum/plasma or urine), i.e., the ratio of its metabolite over its parent compound, are generally greater in magnitude and should not be compared in quantitative terms to changes in standard PK parameters derived from complete plasma concentration–time profiles, such as AUC or clearance. Therefore, it needs to be considered that positive interaction results from studies reporting metabolic ratios always necessitate further *in vivo* studies to provide quantitative data on exposure changes (AUC, C_{max}) and clearance alterations.

There is an apparent tendency in the published studies to over-interpret and over-communicate the statistical significance of identified herb-drug interactions, without adequate consideration of the overall magnitude of findings and their potential (lack of) clinical significance. One of the possible reasons in that regard might be the frequent reference to metabolic ratios, without adequate consideration how these actually may translate in clearance alterations. The recent FDA Guidance on Drug Interaction Studies lists *Echinacea* as a weak inhibitor of CYP1A2 and CYP2D6 and ginkgo as well as goldenseal as weak inhibitors of CYP3A [81]. According to FDA definitions, weak inhibitors are defined as perpetrators increasing the AUC of a sensitive substrate by less than 2-fold but equal or more than 1.25-fold or that produce a 20–50% decrease in clearance. Therefore, the clinical relevance of herb-drug interactions summarized in this review, at first sight, appears to be restricted to interactions with narrow therapeutic index or critical dose drugs. However, it needs also to be acknowledged that categorization of inhibitor/inducer characteristics (i.e., effect sizes) inherently refers to average study outcomes, and that individual responses might differ substantially from population means, based on a variety of extrinsic factors (e.g., diet, nature and number of concomitant medications) and the patients’ genetic makeup and metabolizing capacity. Hence, it would be misleading to conclude that herbal drugs conferring

only a weak inhibitory/inducing effect on certain transporters/enzymes would be generally safe in each patient and clinical setting. In this respect, it is interesting to note that the use of herbal drugs appears to be particularly high in patient populations already exposed to complex treatment algorithms and polypharmacotherapy, often involving narrow therapeutic index drugs, e.g., in oncology [108], HIV-infected patients [109], recipients of organ transplants [110], and in the general elderly population, often affected by multimorbidity. Surveys indicate an overall prevalence for herbal preparation use of 13% to 63% among cancer patients [111], about two-thirds (67%) in HIV-infected patients receiving antiretroviral (ARV) therapy [112], and about 75% and 49% in dialysis and renal transplant patients, respectively [110]. As in these clinical settings and vulnerable patients even modest herb-drug interaction may contribute to treatment failure (e.g., organ rejection) or increased incidences or severity of adverse events, general caution is advised, and health care professionals in these fields should pay attention to the herbal supplement use of their patients and offer proactive advice in this respect. We do hope that the present review provides adequate and useful scientific support in this regard.

In conclusion, substantial progress has been made over the last years in the clinical assessment of pharmacokinetic herb-drug interactions, as exemplified by the identified clinical study data on six popular herbal drugs. However, there is still a need for additional well-designed clinical trials that will increase our understanding about the underlying mechanisms of herb-drug interactions, and the proper communication of the clinical relevance and implications of respective findings, in order to improve the overall sensitivity of health care professionals and patients and to focus on herb-drug interactions that are likely to be of clinical concern, thereby ultimately enabling and continuously improving informed clinical decision making.

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Conflict of Interest



The authors have no conflict of interest to declare.

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