#### **REVIEW ARTICLE**



# Metabolism and interactions of Ivermectin with human cytochrome P450 enzymes and drug transporters, possible adverse and toxic effects

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#### **Abstract**

The review presents metabolic properties of Ivermectin (IVM) as substrate and inhibitor of human P450 (P450, CYP) enzymes and drug transporters. IVM is metabolized, both in vivo and in vitro, by C-hydroxylation and O-demethylation reactions catalyzed by P450 3A4 as the major enzyme, with a contribution of P450 3A5 and 2C9. In samples from both in vitro and in vivo metabolism, a number of metabolites were detected and as major identified metabolites were 3"-O-demethylated, C4-methyl hydroxylated, C25 isobutyl-/isopropyl-hydroxylated, and products of oxidation reactions. Ivermectin inhibited P450 2C9, 2C19, 2D6, and CYP3A4 with IC<sub>50</sub> values ranging from 5.3 µM to no inhibition suggesting that it is no or weak inhibitor of the enzymes. It is suggested that P-gp (MDR1) transporter participate in IVM efflux at low drug concentration with a slow transport rate. At the higher, micromolar concentration range, which saturates MDR1 (P-gp), MRP1, and to a lesser extent, MRP2 and MRP3 participate in IVM transport across physiological barriers. IVM exerts a potent inhibition of P-gp (ABCB1), MRP1 (ABCC1), MRP2 (ABCC2), and BCRP1 (ABCG2), and medium to weak inhibition of OATP1B1 (SLC21A6) and OATP1B3 (SLCOB3) transport activity. The metabolic and transport properties of IVM indicate that when IVM is co-administered with other drugs/chemicals that are potent inhibitors/inducers P4503A4 enzyme and of MDR1 (P-gp), BCRP or MRP transporters, or when polymorphisms of the drug transporters and P450 3A4 exist, drug–drug or drug–toxic chemical interactions might result in suboptimal response to the therapy or to toxic effects.

**Keywords** Ivermectin · P450s · Transporters · Metabolism · Inhibition · Induction

## Introduction

Avermectin compounds are a class of 16-membered macrolide lactone compounds produced by the fungus *Streptomyces avermitilis*, which synthesizes a complex of eight individual avermectin compounds (Burg et al. 1979). Ivermectin (IVM), as one of the avermectin derivative, is a semisynthetic derivative of avermectin B1 and consists of an 80:20 mixture of the equipotent homologous 22,23-dehydro B<sub>1a</sub> and B<sub>1b</sub> (Fig. 1). IVM is a widely used FDA-approved broad-spectrum antiparasitic drug used also to treat pest insects and was found to be especially effective to decrease *P. vivax* transmission (Hotson 2020; Campbell et al. 1983; Pinilla et al. 2018). As an anthelminthic drug, IVM is used to treat various parasitic infestations in veterinary and human

medicine. Applications include river blindness (onchocerciasis) (African region Onchocerciasis Control Programme of West Africa, 1975–2002, (Mectizan) by Merck & Co. Inc., in 1988), head lice, scabies, lymphatic filariasis, ascariasis, enterobiasis, strongyloidiasis, and trichuriasis (Lindley 1987; Boatin and Richards 2006; Moncayo et al. 2008; Anselmi et al. 2015; Thomsen et al. 2016; King et al. 2018; Sharun et al. 2019).

In the treatment of onchocerciasis, the optimal dose of IVM is 150  $\mu$ g/kg once or three times yearly (González Canga et al. 2008). The drug is effective in most patients with scabies after a single oral dose of 0.2 mg/kg, which is the maximal FDA-approved dose, but often the regimen involves two or three repeated doses, separated by intervals of 1 or 2 weeks (Dourmishev et al. 2005). Pharmacokinetic data of IVM in humans following single oral dose (6 and 12 mg) showed that no unchanged or conjugated Ivermectin could be detected in urine, AUC and  $C_{\rm max}$  were dosing proportional and increased linearly with dose with  $t_{\rm max}$ 

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**Fig. 1** Ivermectin  $B_{1a}$  (IVM- $B_{1b}$ ) and  $B_{1b}$  (IVM- $B_{1b}$ )

approximately 4 h and  $t_{1/2}$  elimination, depending on the doses applied, was up to 91 h. Due to its high lipid solubility, IVM is widely distributed within the body and strongly binds to plasma proteins (93.2  $\pm$  4.4% (SD) (Klotz et al. 1990; Krishna and Klotz 1993; Guzzo et al. 2002; Muñoz et al. 2018).

In addition to its antiparasitic action, IVM was also reported to inhibit the growth of some human cancer cells. IVM has powerful antitumor effects, including the inhibition of proliferation, metastasis, and angiogenic activity in different cancer cells (Korystov et al. 2004; Melotti et al. 2014; Kwon et al. 2015; Dou et al. 2016; Zhu et al. 2017; Nambara et al. 2017; Juarez et al. 2018; Jiang et al. 2019; Liu et al. 2020; Tang et al. 2021).

IVM also possess broad-spectrum antiviral property in vitro against RNA and deoxyribonucleic acid (DNA) viruses, including human immunodeficiency virus-1 (HIV-1), dengue virus (DENV), influenza, Venezuelan equine encephalitis virus (VEEV), a flavivirus, and Zika virus (Wagstaff et al. 2012; Mastrangelo et al. 2012; Pandey et al. 2020; Yang et al. 2020). A single dose of IVM effectively removed SARS-CoV-2 viral RNA in cell culture by 48 h. It was suggested that IVM might be used for the treatment of COVID-19 patients in the early phase with mild to moderate symptoms, PCR positive for SARS-CoV-2 virus, and those without comorbidities such as chronic obstructive pulmonary disease, diabetes, hypertension, obesity, acute or chronic renal failure, and coronary diseases. IVM suggested anti-CoV-2 effects are based on in vitro studies, showing that this compound inhibits the importin alpha/beta-1 nuclear transporter, and thus, in cell cultures, reduces the replication of various viruses including SARS-CoV-2 (Bray et al. 2020; Caly et al. 2020; Rizzo 2020; Sharun et al. 2020; Heidary and Gharebaghi 2020; Dixit et al. 2020; Chaccour et al. 2020a, b; Formiga et al. 2020). In addition, in a silico-based analysis of Ivermectin's molecular interaction indicates positive interaction of Ivermectin with viral protein targets, which is leading for SARS-CoV 2 N-protein NTD (nucleocapsid protein N-terminal domain) (Kaur et al. 2021). Avermectins have also antifungal activity and are used for the treatment of several metabolic disorders. Ivermectin plasma concentrations required to reach an in vitro antiviral efficacy (about 2–10 µM) might be very toxic as very high oral doses would be needed for antiviral use and overdose or uncontrolled administration of IVM can lead to the development of severe toxicity (El-Saber Batiha et al. 2020). Ivermectin has not been approved for the treatment of viral diseases, including SARS-CoV-2, and in April 2020, FDA issued a warning not to use Ivermectin to treat COVID-19 in humans (Bray et al. 2020).

The pharmacokinetics and hepatic metabolism of IVM was investigated in a subpopulation of the CF-1 mouse stock naturally deficient in mdr1a P-glycoprotein (multidrug-resistance protein). Intestinal excretion of (<sup>3</sup>H)IVM enhances in P-gp (+/+) animals. The absence of P-gp resulted in higher blood concentrations of IVM after oral dosing, suggesting enhanced absorption of IVM in (-/-) mice. Concentrations of (<sup>3</sup>H)IVM were always greater in the brains of (-/-) mice compared with (+/+) mice after either i.v. or oral administration. In contrast, liver concentration was not different between (+/+) and (-/-) animals after an i.v. dose. These results show the P-gp (-/-) and (+/+) subpopulations of CF-1 mice are useful for studying the role of P-gp in systemic exposure and tissue



disposition of P-gp substrates (Umbenhauer et al. 1997; Kwei et al. 1999). A mechanism of antiparasitic action of IVM is inhibition of a glutamate-gated chloride channel and a GABA receptor, specific for some invertebrates. In mammals, IVM may also inhibit GABAergic neurotransmission by promoting the release of GABA and acting as a GABA receptor agonist exerting neurotoxic activity (Yang 2012; Ménez et al. 2012). Potential neurotoxic activity in humans depends on the absorption-extrusion activity of the drug from the gastrointestinal tract/blood-brain barrier which is regulated by P-gp, MRPs, ABCB1, and other ABC transporters. In the case of IVM, it was suggested that the severe central nervous system side effects seen in various vertebrates following IVM treatment may be due to an absence of, or functional deficiency in P-glycoprotein (Edwards 2003). Due to the wide range of its clinical applications and its possible co-administration with other drugs when used to treat various illnesses, there is a possibility that toxic and/or clinically significant drug-drug interactions might occur. Consequences of toxic drug-chemical and/or drug-drug interactions which might occur during co-treatment of IVM with other drugs/ chemicals will depend on metabolic and pharmacokinetic properties of IVM and co-administered drugs, i.e., their interactions with drug-metabolizing enzymes (cytochrome P450s) and drug transporters involved in IVM metabolic pathways. The interactions might thus result in changes of activity and/or expression of drug metabolism enzymes and/or transporters with consequences such as changed responses to drug treatment or significant adverse/toxic effects.

In the earlier paper discussed was metabolism and interactions of chloroquine and hydroxychloroquine with human cytochrome with P450 enzymes and drug transporters. These drugs are since years used in the treatment and prevention of malaria and chronic inflammatory diseases but also proposed to be used in treatment of COVID-19 (alone or in combination with other drugs). The data on their metabolism showed that they are metabolized by white range of P450 enzymes in vitro (1A2, 2C8, 2C19, 2D6, and 3A4/5) and by P450s 2C8 and 3A4/5 in vivo. In addition, chloroquine effectively inhibited P450 2D6 in vitro and in vivo in particular in persons with limited P450 2D6 activity. Concluded was that by co-administration of the drugs with other drugs, substrates of P450 2D6 enzyme, attention should be given to substratespecific metabolism by P450 2D6 alleles present in persons taking the drugs for possible drug-drug interactions (Rendic and Guengerich 2020).

The present paper summarizes data published on the metabolism of IVM by human cytochrome P450 enzymes, interactions with drug transporters, and the potential to cause drug-drug/drug-chemical interactions and toxicity which might occur in a wide range of its clinical applications.

## **Results and discussion**

# Metabolism of Ivermectin (IVM) by P450 enzymes

IVM is extensively metabolized by cytochrome P450 enzymes (P450s, CYP) both in vivo and in vitro. In vivo IVM metabolites are predominately excreted in feces with only 1% of the doses in the urine. Using human liver microsomes, human recombinant enzymes, hepatocytes and blood samples up to 13 metabolites were identified. The major metabolites of IVM-B<sub>1a</sub> in humans occurs by the following metabolic recitations: methyl-hydroxylation, O-demethylation, and oxidation reactions occurring at different positions of the molecule. The structure of the major five metabolites of IVM identified in vitro indicated that they occur by 3"-O-demethylation, C4-hydroxylation, C25 isobutyl-hydroxylation, and 3"-O-demethyl-4-hydroxymethyl metabolite formation (Fig. 2). The major contribution to the metabolism of IMV in vitro is attributed to P450 3A4. In addition, 3"-O-demethyl metabolite was also produced to a small extent by P450 3A5 and is considered as a minor metabolite. The other hydroxylated metabolite occurs also by P4502C8 (Table 1 and references therein). Of the major metabolites of IVM formed in vitro, three of them were identified also in human plasma and occur by 3"-O-demethylation, C4-hydroxylation, and the metabolite formed by both 3'-O-demethylation and C4-hydroxvlation reactions. These reactions are catalyzed in vitro by P450 3A4 and 3"-O-demethylation was also catalyzed in minor extent by P450 3A5 (Zeng et al. 1998; Tipthara et al. 2021). The data showed that IVM is predominately substrate of P450 3A4 and that co-administration of drugs that are potent inhibitors or inducers of the enzyme activity, or polymorphism of the enzyme might provoke pronounced drug-drug interactions. Potent inhibitors of P450 3A4 activity are, for example, ketoconazole, econazole, troleandomycin, azamulin, cyclosporine, while potent inducers of P4503A4 activity are, for example, carbamazepine, cortisol, hydrocortisone, rifampicin, amprenavir, and clotrimazole (Rendic 2002).

### Inhibition of P450 enzymes by Ivermectin (IVM)

IVM was demonstrated as a weak, no, or medium inhibitor of the reactions catalyzed by P450 enzymes (Table 2 and references therein). The potency of inhibition for  $IC_{50}$  and % of inhibition values were rated according to the values and ranges as presented in Table 3. Taking into account these data and the data on the inhibition of P450 3A4 IVM was assigned as a medium to week inhibitor

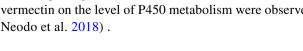


Fig. 2 Ivermectin (IVM) metabolites formed by cytochrome P450 enzymes

Table 1 Metabolism of Ivermectin (IVM) with human cytochrome P450 enzymes

| P450 | Method  | Reaction   | Reference PUBMED IDs | References   |
|------|---|--|----------------------|--|
| 3A4  | Recombinant enzyme, yeast expressing<br>human P450; microsomes, hepatocytes,<br>blood samples | Hydroxylation, C4-CH <sub>3</sub> -Hydroxylation, C26-CH <sub>3</sub> Demethylation, 3"-O- | •                    | Zeng et al. (1998), Tipthara et al. (2021)<br>Zeng et al. (1998), Tipthara et al. (2021)<br>Zeng et al. (1998), Tipthara et al. (2021) |
| 3A5  | Recombinant enzyme, yeast expressing human P450; microsomes, hepatocytes                      | Hydroxylation, C4-CH <sub>3</sub> -Demethylation, 3"-O-                                    | 33497030             | Tipthara et al. (2021)   |
| 2C8  | Recombinant enzyme, yeast expressing human P450; microsomes;                                  | Hydroxylation, CH <sub>3</sub>   | 33497030             | Tipthara et al. (2021)   |

of P450 3A4. The IC<sub>50</sub> values of inhibition of P450 2C9, 2D6, and 2C19 enzymes ranged from 5.3 to 13.8 µM, or no inhibition (Table 2 and references therein). Thus, the data reported show that they are at least somewhat dependent on the experimental model used, that the potential for drug-drug interactions due to inhibition of P450 enzymes is not likely, and that it will depend on the dose applied. As blood plasma concentrations of IVM in humans were reported to be rarely higher than 100 ng/ml (about 0.1 µM) at a standard oral dose, it might be concluded that IVM, when applied under clinically recommended dose, does not have the potential to cause drug-drug interaction potential on the level of inhibition of P450 enzymes. In addition, it was reported that co-administration of IVM and tribendimidine in rats did not significantly alter the pharmacokinetic behavior of each other and no clinically relevant drug-drug interactions between tribendimidine and Ivermectin on the level of P450 metabolism were observed (Neodo et al. 2018).





# Interaction of Ivermectin (IVM) with drug transporters

In the present chapter discussed is the role of transporters that belong to the ATP-binding cassette (ABC) transporter family and solute carrier family (SLC) transporters in the transport of IVM (Table 4 and references cited). ABC transporters are membrane-bound proteins that are present in all prokaryotes, as well as plants, fungi, yeast, and animals. In humans, the ABC transporters take part in the transport of a variety of compounds (drugs, environmental and physiological compounds) using the energy of ATP and have important role in drug disposition and effects/toxicities by affecting the absorption, distribution and excretion of drugs (Liu 2019).

The multiple drug-resistance transporter (MDR, P-glycoprotein, P-gp) is an export transporter discovered for its role in imparting resistance to drugs in cancer cells by pumping them out (Bradley et al. 1988). The transporter was in the course of research assigned in the literature under different names (Table 4). Its role as a drug export protein was

Table 2 Inhibition of cytochrome P450-catalyzed metabolic reactions by Ivermectin (IVM)

| P450 | Method  | Remarks  | Reference<br>PUBMED<br>IDs | References            |
|------|---|--|----------------------------|-----------------------|
| 1A2  | Recombinant enzyme, yeast expressing human P450, microsomes, fluorescence-based high-throughput screening assays                              | no inhibition, 8% of test reaction inhibition  | 11124226                   | Bapiro et al. (2001)  |
|      | Recombinant enzyme, yeast expressing human P450, microsomes, fluorescence-based high-throughput screening assays (Vivid CYP450 screening kit) | $IC_{50} > 10 \mu M$                           | 30323047                   | Neodo et al. (2018)   |
| 2C9  | Recombinant enzyme, yeast expressing human P450, microsomes, fluorescence-based high-throughput screening assays                              | no inhibition                                  | 11124226                   | Bapiro et al. (2001)  |
|      | Recombinant enzyme, yeast expressing human P450, microsomes, fluorescence-based high-throughput screening assays (Vivid CYP450 screening kit) | IC <sub>50</sub> 5.3 (range 4.5–6.2)           | 30323047                   | Neodo et al. (2018)   |
| 2C19 | Recombinant enzyme, yeast expressing human P450, microsomes, fluorescence-based high-throughput screening assays                              | no inhibition, 11% of test reaction inhibition | 11124226                   | Bapiro et al. (2001)  |
|      | Recombinant enzyme, yeast expressing human P450, microsomes, fluorescence-based high-throughput screening assays (Vivid CYP450 screening kit) | IC <sub>50</sub> 6.7 μM (range 5.3–8.9)        | 30323047                   | Neodo et al. (2018)   |
| 2D6  | Recombinant enzyme, yeast expressing human P450, microsomes, fluorescence-based high-throughput screening assays                              | no inhibition, 10% of test reaction inhibition | 11124226                   | Bapiro et al. (2001)  |
|      | Recombinant enzyme, yeast expressing human P450, microsomes, fluorescence-based high-throughput screening assays                              | IC <sub>50</sub> 13.8 μM (range 10.0–27.1)     | 30323047                   | Neodo et al. (2018)   |
| 3A4  | Recombinant enzyme, yeast expressing human P450, microsomes, fluorescence-based high-throughput screening assay                               | $K_i$ 27.2 $\mu M$                             | 11124226                   | Bapiro et al. (2001)  |
|      | Human liver microsomes using triazolam-4-hy-<br>droxylation, % of inhibition  | <10%   | 12948015                   | Perloff et al. (2003) |
|      | Fluorescence-based microtiter plate screening assay   | $IC_{50} > 50 \mu M$                           | 12699389                   | Schwab et al. (2003)  |
|      | Recombinant enzyme, yeast expressing human P450, microsomes, fluorescence-based high-throughput screening assays (Vivid CYP450 screening kit) | IC <sub>50</sub> 11.3 μM (range 9.1–19.8)      | 30323047                   | Neodo et al. (2018)   |

realized in the late 1990s and with time, the number of proteins involved in drug transport processes was grown. P-gp is present in intestinal epithelial cells, liver, kidney, and brain. It was also realized that drug's transport in or out of the cells, as well as passive diffusion, is dependent on the transporter's activity, having important role in many cases of drug transport, and can have major effects on tissue selectivity of drugs. Another point is that P-gp export transporter can also lower the level of a drug in a cell and so slow the rate of its metabolism and influence pharmacokinetics and therapeutic activity of drugs, including anticancer drugs (Lin and Yamazaki 2003; Jiang et al. 2019). Similarly, MRP1, 2, and 3 are efflux transporters that are present in hepatocytes with wide substrate transport specificity, including important

drugs. The main roles of the MRP transporter proteins are the efflux of xenobiotics (drugs and food components) and endogenous metabolites, transport of inflammatory mediators (e.g., LTC4), and defense against oxidative stress. In addition, P-gp and MRP1 play a role in the development of drug resistance of various types of cancer and contribute to inflammatory responses (Cole and Deeley 1998; Cole 2014; Nakanishi and Tamai 2015; Guengerich and Rendic 2010).

IVM is a potent inhibitor of the MDR1 (P-gp) (Pouliot et all 1997; Didier and Loor 2015) and MRP transporting proteins (Table 4 and referenced cited). For instance, RH123 accumulation into LS180V cells was used as an assay for P-gp inhibition in the screening of 17 clinically used drugs. The results showed that IVM was one of the most potent



**Table 3** Potency of inhibition of the drug/compound to inhibit P450 enzymes and transporters

| Inhibition                         | $IC_{50}(K_i) > 100 \mu M$   | Very weak inhibitor                 |
|------------------------------------|------------------------------|-------------------------------------|
|                                    | $IC_{50}(K_i) > 30 \mu M$    | Weak inhibitor                      |
|                                    | $IC_{50}(K_i)$ 1–30 $\mu M$  | Medium inhibitor                    |
|                                    | $IC_{50}(K_i) < 1 \mu M$     | Potent inhibitor                    |
|                                    | $IC_{50}(K_i) < 0.010 \mu M$ | Highly potent inhibitor             |
| % of inhibition of tested reaction | ≥99%                         | Highly potent inhibitor             |
|                                    | >90%                         | Potent inhibitor                    |
|                                    | 50—90%                       | Medium inhibitor                    |
|                                    | 20—50%                       | Weak inhibitor                      |
|                                    | 10—20%                       | Very weak                           |
|                                    | < 10%                        | No inhibitor or very weak inhibitor |
| % of the control reaction activity | <1%                          | Highly potent inhibitor             |
|                                    | < 10%                        | Potent inhibitor                    |
|                                    | 10—50%                       | Medium inhibitor                    |
|                                    | 50—80%                       | Weak inhibitor                      |
|                                    | 80—90%                       | Very weak inhibitor                 |
|                                    | >90%                         | No inhibitor or very weak inhibitor |
|                                    | <1%                          | Highly potent inhibitor             |

P-gp inhibitors, causing a ninefold increase in RH123 accumulation compared to control cells (Perloff et al. 2003). The data presented in Table 4 with IC $_{50}$  ranging from 0.1 to 2.5  $\mu$ M classify IVM to potent inhibitors of the P-gp mediated transport (Table 3). The data also show that MRP1, and to a lesser extent MRP2 and MRP3 take part also to IVM flux across physiological barriers (Schwab et al. 2003) and that IVM is also a substrate of P-gp. Thus, the MDR1 and MRP binding properties and its large molecular size (Fig. 1) make IVM potential competing inhibitor when coadministrated with other drugs/chemicals which are P-gp and MRPs substrates. IVM also inhibited the P-gp, MRP1, 2, and 3 ATPase activity after stimulation by their respective activators (Table 4 and references cited).

Organic-anion transporting polypeptides (OATP transporters, OATPs) are influx transporters having important role of drug transport in intestine and transport across blood—brain barrier (OATP1A2, OATP2B1) and hepatocytes (OATP1B1, OATP1B3). Thus, OATPs might have impact on drug disposition and efficacy by facilitating oral absorption and drug access to the liver and brain (Nakanishi and Tamai 2015; Varma et al. 2010; Shitara et al. 2013). IVM, when interacting with organic anion transporter polypeptide family, showed to be weak to a medium inhibitor of the activity (Table 3 and Table 4 and references cited). Thus, drug—drug interactions on the level of OATPs by IVM might be possible with application of high doses of IVM.

BCRP (ABCG2) is a member of the ATP-binding cassette (ABC) superfamily of transmembrane transporters and localized to the apical plasma membrane of cells. It actively transports xenobiotic and endogenous substrates out of the cell with energy derived from the hydrolysis of ATP. BCRP

was discovered in cancer cells but present also in normal cells preventing the accumulation of chemicals and their toxicity. It has a broad substrate specificity and important role by modulating efflux and having an impact on the pharmacokinetics of drugs, nutrients, and toxicokinetic of toxicants, and to have an important role in transporter-mediated drug-drug interactions (Bricsak and Aleksunes 2015; Safar et al. 2019). IVM inhibits BCRP (Table 4 and references cited) and according to the reported IC<sub>50</sub> values (1–3.1  $\mu$ M) IVM is classified as a potent to medium potency inhibitor (Table 3). Consequently, this interaction may also have implications in the metabolism and distribution of IVM and give rise to drug-drug interactions (Jani et al. 2011). The properties of IVM as a potent multidrug-resistance reversal agent, when combined with antitumor agents, led to its application as enhancer of antitumor activity. IVM drastically reversed the resistance of the tumor cells to the chemotherapeutic drugs both in vitro and in vivo. IVM reversed the resistance by inhibiting the proliferation of several tumor cells by regulating multiple signaling pathways and not by directly inhibiting P-gp activity. The inhibition of the transcriptional factor NF-κB led to the reduced P-gp transcription in tumor cells leading to reversal of the resistance of the tumor cells (Jiang et al. 2019; Tang et al. 2021).

On the level of P450 enzymes, drug—drug interaction might be provoked by co-administration with drugs which are potent inhibitors of P450 3A enzymes by lowering its extrusion from the body. Alternatively, potent inducers of P450 3A activity might affect IVM systemic exposure lowering its therapeutic effect. On the level of drug transport, by inhibition P-gp or MRPs, unexpected high plasma concentration and potential toxic effects of IVM might be elicited. On



Table 4 Interaction of Ivermectin (IVM) with human drug transporters

|   |           | 0                         |   |   |                            |   |
|---|-----------|---------------------------|---|---|----------------------------|---|
| Transporter   | Substrate | Substrate Inhibitor Model | Model   | Inhibition/remarks  | Reference<br>PUBMED<br>IDs | Reference no                                |
| BCRP1; BCRP; ABCG2; MRX; MXR;<br>ABCP; BMDP; MXR1; ABC15; CD338;<br>CDw338; ST157481; MGC102821 |           | X                         | Cellular and membrane assays  IC <sub>50</sub> values i Intact PLB-985 cells, PhenGreen (PG)-AM IC <sub>50</sub> 3.1 µM extrusion | $IC_{50}$ values in the 1–1.5 $\mu M$ range $IC_{50}$ 3.1 $\mu M$ | 20574995<br>33435273       | Jani et al. (2011)<br>Telbisz et al. (2021) |
| MDR1, P-glycoprotein, P-gp, ABCB1, ABC*20, CD243, CLCS, GP170, PGY                              | ×         | ×                         | LLCPK1-mdr1 cells fluorescent derivative BODIPY Ivermectin  | Substrate at low drug concentration                               | 12699389                   | Schwab et al. (2003)                        |
|   |           |                           | P-gp membranes (MDR1), calcein-AM assay   | $IC_{50}  0.1  \mu M$   |                            |   |
|   |           |                           | CaCo-2 assay, RHO-123 assay, inhibition relative to the positive control nicardipine  | 34%   |                            |   |
|   |           |                           | SH-SY5Y cells, H33342 dye efflux,   | Ki $0.24 \pm 0.08  \mu M$   | 25865432                   | Dalzell et al. (2015)                       |
|   |           |                           | CaCo-2 assay, RHO-123 assay   | $IC_{50} 0.10 \pm 0.04  \mu M$                                    | 15953199                   | Griffin et al. (2005)                       |
|   |           |                           | Sf9 cells, Membrane vesicles, activated ATPase activity   | $IC_{50} 2.5 \pm 2.0  \mu M$<br>Ki calc. 0.06 $\mu M$             | 16384,552                  | Lespine et al. (2006)                       |
|   |           |                           | Sf9 cells, Membrane vesicles, basal ATPase  | Half-inhibition at about 0.1–0.2 µM                               |                            |   |
|   |           |                           | LLCPK1-Mdr cells, Rho123  | $IC_{50}$ 0.4 $\mu M$   |                            |   |
|   |           |                           | PLB-985/ABCB1 Cells, Calcein-AM extrusion assay   | IC <sub>50</sub> 0.6 μΜ   | 33435273                   | Telbisz et al. (2021)                       |
|   |           |                           | HEK/-293 ABCB1 membrane vesicles, 3H-N-methyl-quinidine uptake  | IC <sub>50</sub> 0.3 µM   |                            |   |
|   |           |                           | Sf9 cells, activated ABCB1 ATPase activity  | inhibition  |                            |   |
|   |           |                           | Caco-2 cells, RH123 exclusion assay   | tenfold increase in RH123 cell accumulation                       | 12948015                   | Perloff et al. (2003)                       |
|   |           |                           | Modulation of basal P-gp ATPase activity<br>Modulation of verapamil-stimulated P-gp<br>ATPase activity                            | EC <sub>50</sub> 2.0 μМ;<br>Ki 0.05 μМ                            | 17134,887                  | Lespine et al. (2007)                       |
|   |           |                           | LLC-PK1-mdr1a cells, RH123 exclusion assay  | 0.44±0.07 μM  |                            |   |



|                       | Inhibition/remarks Reference no PUBMED IDs |
|-----------------------|--|
|                       | Substrate Inhibitor Model Inhi             |
| 7 Table 4 (continued) | Transporter                                |

| Transporter   | Substra | Substrate Inhibitor | itor Model  | Inhibition/remarks   | Reference<br>PUBMED  | Reference no                                    |
|---|---------|---------------------|---|--|----------------------|---|
|   |         |                     |   |  | IDs                  |   |
| MRP1, ABC29, ABCC1, GS-X, MRP                                 | ×       | ×                   | HL60-MRP1 cells, BODIPY Ivermectin fluorescent derivative                     | Substrate at higher concentrations   | 16384552             | Lespine et al. (2006)                           |
|   |         |                     | Sf9 cells membrane vesicles, activated ATPase                                 | $IC_{50} > 10 \mu M$   |                      |   |
|   |         |                     | Sf9 cells membrane vesicles, activated ATPase                                 | $IC_{50} 9 \pm 4 \mu M$<br>Ki calc. 0.5 $\mu M$                            |                      |   |
|   |         |                     | HL60-MRP1 cells, inhibition of Calcein-AM efflux                              | IC <sub>50</sub> 3.8 μΜ  |                      |   |
|   |         |                     | A549 tumor cells, calcein retention   | IС <sub>50</sub> 1 µМ  |                      |   |
|   |         |                     | A549 tumor cells, calcein retention   | $IC_{50}$ 2.5 $\mu M$  |                      |   |
|   |         |                     | Human SH-SY5Y, fluorescent dye H33342 transport                               | $IC_{50} 1.58 \pm 0.51  \mu M$   | 25865,432            | Dalzell et al. (2015)                           |
|   |         |                     | HL60/ABCC1 Cells, Calcein-AM extrusion assay                                  | $IC_{30}$ 3.3 $\mu M$  | 33435273             | Telbisz et al. (2021)                           |
|   |         |                     | Sf9/ABCC1 membrane vesicles, 3H-estradiol-17b-glucuronide vesicular transport | 80% inhibition at a 25 $\mu$ M concentration IC <sub>50</sub> 13.3 $\mu$ M |                      |   |
| MRP2, ABCC2, ABC30, CMOAT, DJS,                               | ×       | ×                   |   | Substrate at higher concentrations;  | 16384552             | Lespine et al. (2006)                           |
| cMRP  |         |                     | Membrane vesicles from Sf9 cells, activated ATPase activity                   | $IC_{50}$ 18±5 μM<br>Ki calc. 3.6 μM                                       |                      |   |
| MRP3, ABC31, EST90757, MLP2,                                  | ×       | ×                   |   | Substrate at higher concentrations   |                      |   |
| MOAT-D, cMOAT2  |         |                     | Membrane vesicles from Sf9 cells, activated ATPase activity                   | IC <sub>50</sub> 40±21 μM  |                      |   |
| OATP1A2, OATP-A, OATP1, OATP, SLC21A3, SLC01A2                |         | ×                   | A431-OATP1A2 cells, SR101 uptake  | IC <sub>50</sub> 5.2 μM  | 33435273             | Telbisz et al. (2021)                           |
| OATP1B1, SLC21A6, HBLRR, LST-1,<br>LST1, OATP-C, OATP2, OATPC |         | ×                   | Human embryonic kidney 293 (HEK293) cells, estradiol-17β-glucuronide uptake   | 64.7% inhibition   | 22541068             | Karlgren et al. (2012)                          |
| OATP2B1, SLCO2B1, OATP-B, OATPB, SLC21A9                      |         | ×                   | A431-OATP2B1 cells, pyranine uptake<br>Human embryonic kidney 293 (HEK293)    | IC <sub>50</sub> 8.6 μM<br>39.0% inhibition                                | 33435273<br>22541068 | Telbisz et al. (2021)<br>Karlgren et al. (2012) |
|   |         |                     | cells, estrone-3-sulfate uptake   |  |                      |   |
| OATPIB3, SLCOIB3, HBLRR, LST-2, LST-3TM13. LST3. OATP-8       |         | ×                   | Human embryonic kidney 293 (HEK293) cells, estradiol-17β-glucuronide uptake   | 55.2% inhibition   |                      |   |



another side, induction of transporter's activity in the host, as observed for induction of ABC transporters by rifampicin (Chaccour et al. 2017), or its synthesis by co-administered drugs, might result in subtherapeutic concentrations at the site of action, or cause the development of strains resistant to the drug (James and Davey 2009). Drug-drug interactions at the level of transporters may affect systemic exposure blocking influx transporters in the intestine, but increase it by modulating influx and efflux transporters in the liver and efflux transporters in the small intestine (Nakanishi and Tamai 2015). In addition to co-administered drugs, also polymorphism of drug transporters and P450 enzymes as well as foods might influence the metabolism and pharmacokinetics of IVM in humans. The plasma concentration of IVM in healthy volunteers was found as much higher in beer drinkers, but orange juice decreased AUC values of IVM. The latter effect was ascribed to the inhibition of drug transporters (González Canga et al. 2008). Thus, the results and discussion presented suggest that co-administration of IVM with other drugs, polymorphism of the drug-metabolizing enzymes and transporters or even kinds of food, might affect biotransformation and transport of IVM affecting its pharmacokinetic properties and provoke drug-drug interactions or toxic effects.

# **Concluding remarks**

In clinical practice, IVM is often co-administered with other drugs when different diseases are treated (parasitic, cancers, and COVID-19). Therefore, the effect of IVM and co-administered drugs on P450 enzymes and ABC-mediated transport have to be considered to provide the most effective treatments for patients. IVM is metabolized, both in vivo and in vitro, by C-hydroxylation and O-demethylation reactions catalyzed by P450 3A4, and with a contribution of P450 3A5 and 2C9 enzymes.

P-gp (MDR1) participates in IVM efflux at low drug concentration with slow transport rate. In the higher, micromolar concentration range, which saturates MDR1 (P-gp), MRP1, and to a lesser extent MRP2 and MRP3 take part in IVM flux across physiological barriers. IVM exert a potent inhibition of P-gp (ABCB1), MRP1 (ABCC1), MRP2 (ABCC2), and BCRP1 (ABCG2), and medium to weak inhibition of OATP1B1 (SLC21A6) and OATP1B3 (SLCOB3) transport activity. The data reported emphasize the potentially unsafe/ toxic effects when IVM is applied at higher doses, in the subjects with an impaired P-gp transporter function (Ménez et al. 2012). It might be well expected that in the clinic, following application of higher doses of IVM, the pharmacokinetics of co-administered drugs might be critically altered as result to drug-drug interactions and that toxic effect might be provoked as a result of the inhibition of P-gp and/or MRP

transporters. In addition, polymorphism of the enzymes and transporters which take part in IVM metabolism might additionally contribute to the effects.

The presented data show complex interactions of IVM being a substrate of P450 3A enzymes, and substrate and inhibitor of P-gp and MRP, and an inhibitor of BCRP transporter. Besides its FDA-approved antiparasitic activity, it exerted potent anticancer and antiviral effects. Therefore, co-administration of IVM with other drugs in complex treatments should take particular attention to develop the most effective treatment strategies for patients receiving multiple drug regimens to avoid unwonted and toxic effects.

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#### **Declarations**

**Conflict of interest** The author declare no conflict of interest, financial or otherwise.

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