

Role of transporters in drug absorption and elimination

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Abstract

The study investigates the critical role of transporters in drug absorption and elimination, emphasizing their influence on pharmacokinetics, therapeutic efficacy, and safety. Transporters such as organic anion-transporting polypeptides (OATPs) and P-glycoprotein (P-gp) mediate the influx and efflux of drugs across cellular membranes, significantly impacting bioavailability and systemic clearance. The objectives included identifying key transporters, analyzing their substrate specificity, and assessing the effects of genetic polymorphisms and environmental factors on their activity. A combination of *in vitro* assays, *in vivo* studies, and genetic analyses was employed. Cell lines like Caco-2 and HEK293, along with rodent models, were used to evaluate transporter functionality, while high-performance liquid chromatography and next-generation sequencing provided precise quantification and genetic insights.

The results demonstrated significant modulation of transporter activity by pharmacological agents and genetic variations. Rifampin inhibited OATP1B1 activity, reducing uptake rates by over 50%, while P-gp activation doubled the efflux ratio of specific substrates. Polymorphisms in the SLCO1B1 gene (c.521T>C) were associated with a 40% reduction in OATP1B1-mediated drug uptake, corroborated by increased plasma drug levels in preclinical models. Drug-drug interaction studies revealed a 3.2-fold increase in substrate AUC upon P-gp inhibition, highlighting the clinical implications of transporter modulation. High-fat diets reduced P-gp expression by 25%, demonstrating the impact of environmental factors.

The study concludes that transporters play a pivotal role in drug disposition, with genetic and environmental factors significantly influencing their activity. Recommendations include genetic screening for personalized medicine, careful evaluation of transporter modulators, and integrating transporter studies into drug development. Future research should focus on advanced *in vitro* models, multi-omics approaches, and transporter-specific biomarkers to refine pharmacotherapy strategies.

Keywords: Drug transporters, pharmacokinetics, genetic polymorphisms, drug-drug interactions, personalized medicine, OATP1B1, P-glycoprotein

Introduction

Drug absorption and elimination are pivotal determinants of a drug's therapeutic efficacy and safety. The biological processes governing these phenomena are heavily influenced by transporters, specialized proteins that regulate the influx and efflux of substances across cellular membranes. Transporters play a critical role in dictating drug bioavailability, distribution, metabolism, and excretion. These proteins are classified into uptake transporters, such as organic anion-transporting polypeptides (OATPs), and efflux transporters, including P-glycoprotein (P-gp) and breast cancer resistance protein (BCRP), each exhibiting specificity for distinct substrates. The intricate interplay between transporters and metabolic enzymes underscores the complexity of pharmacokinetics, thereby necessitating a deeper understanding of these mechanisms to optimize drug therapy [1, 2, 3].

Despite advancements in pharmacological sciences, challenges persist in achieving optimal drug absorption and elimination. Transporters often contribute to interindividual variability in drug response, driven by genetic polymorphisms, drug-drug interactions, and disease states. For instance, polymorphisms in the SLCO1B1 gene, encoding for OATP1B1, have been associated with altered hepatic uptake of statins, leading to adverse effects like myopathy [4, 5]. Similarly, overexpression of efflux transporters such as P-gp in cancer cells contributes to multidrug resistance, posing significant obstacles in

oncology [6, 7]. These issues underscore the necessity for research aimed at elucidating the mechanistic roles of transporters in drug absorption and elimination.

This study aims to explore the role of transporters in drug absorption and elimination with an emphasis on their impact on pharmacokinetics and therapeutic outcomes. The objectives include identifying key transporters involved in drug disposition, assessing their substrate specificity, and understanding the influence of genetic and environmental factors on transporter activity. Furthermore, the study hypothesizes that transporter-mediated drug interactions are a major contributor to variability in drug efficacy and toxicity. By addressing these aspects, the research seeks to provide insights that may inform the development of personalized medicine strategies, ultimately enhancing drug safety and efficacy.

Material and Methods

Materials

The study was conducted using a combination of *in vitro* and *in vivo* models to evaluate the role of transporters in drug absorption and elimination. Human-derived cell lines, such as Caco-2 and HEK293 cells, were utilized to assess the activity of uptake and efflux transporters. Recombinant transporter-expressing systems were employed to identify substrate specificity and transporter kinetics. Pharmacokinetic data were obtained from preclinical animal models, including rodents, which were administered model

drugs known to interact with specific transporters. Additionally, clinical samples, such as plasma and tissue biopsies from volunteers with informed consent, were analyzed for transporter expression levels. High-performance liquid chromatography (HPLC) and liquid chromatography-tandem mass spectrometry (LC-MS/MS) were used for drug quantification and metabolite profiling.

Methods

Functional assays were performed to characterize transporter activity, including uptake and efflux studies using radiolabeled or fluorescent substrates. Genetic analyses were conducted to identify polymorphisms in transporter genes, employing techniques such as polymerase chain reaction (PCR) and next-generation sequencing (NGS). Drug-transporter interactions were evaluated using inhibition and stimulation studies, with specific inhibitors or activators of transporters. Pharmacokinetic modeling was carried out using compartmental and non-compartmental approaches to correlate transporter activity with drug disposition parameters. Statistical analyses were conducted using software tools like GraphPad Prism and R to determine the significance of findings. Ethical approval was obtained from relevant institutional review boards, ensuring compliance with guidelines for animal and human research.

Results

Transporter Activity Assays

The activity of uptake transporters (e.g., OATP1B1) and efflux transporters (e.g., P-gp) was quantified *in vitro* using Caco-2 and HEK293 cells. The mean uptake rate of a model substrate for OATP1B1 was 75 ± 8 pmol/mg protein/min, which was significantly reduced ($p < 0.01$) in the presence of rifampin, a known inhibitor. Conversely, the efflux ratio of a P-gp substrate increased from 2.5 ± 0.3 to 5.8 ± 0.4 ($p < 0.001$) when co-incubated with a P-gp activator. These results indicate the sensitivity of transporter activity to specific modulators.

Genetic Polymorphism Analysis

Polymorphisms in the SLCO1B1 gene were associated with significant changes in transporter activity. Individuals carrying the *SLCO1B1* c.521T>C variant exhibited a 40% reduction in OATP1B1-mediated uptake ($p < 0.01$). Pharmacokinetic analysis in rodent models showed a two-

fold increase in plasma levels of statins in animals with reduced transporter activity, corroborating clinical findings.

Drug-Drug Interaction Studies

Inhibition studies revealed significant drug-drug interactions mediated by efflux transporters. Co-administration of a P-gp substrate with a P-gp inhibitor led to a 3.2-fold increase in the area under the concentration-time curve (AUC) of the substrate ($p < 0.001$). Similarly, transporter activity was altered in the presence of environmental factors, such as high-fat meals, which reduced P-gp expression by 25% ($p < 0.05$).

Statistical Analysis

All statistical analyses were performed using GraphPad Prism. The differences in transporter activity and pharmacokinetics between experimental groups were evaluated using one-way ANOVA followed by Tukey’s post-hoc test. A p -value < 0.05 was considered statistically significant. Correlation analyses showed strong associations ($R = 0.85$, $p < 0.001$) between transporter expression levels and drug plasma concentrations, emphasizing the clinical relevance of transporter modulation.

Table 1: Uptake Rate under Different Conditions

Condition	Uptake Rate (pmol/mg protein/min)
Control	75
Rifampin (Inhibitor)	45

Table 2: Efflux Ratio under Different Conditions

Condition	Efflux Ratio
Control	2.5
P-gp Activator	5.8

Table 3: Impact of SLCO1B1 Polymorphism on Uptake Rate

Genotype	Uptake Rate (% of Control)
Wild Type	100
SLCO1B1 c.521T>C	60

Table 4: Impact of P-gp Inhibition on AUC

Condition	AUC Fold Change
Control	1
P-gp Inhibitor	3.2

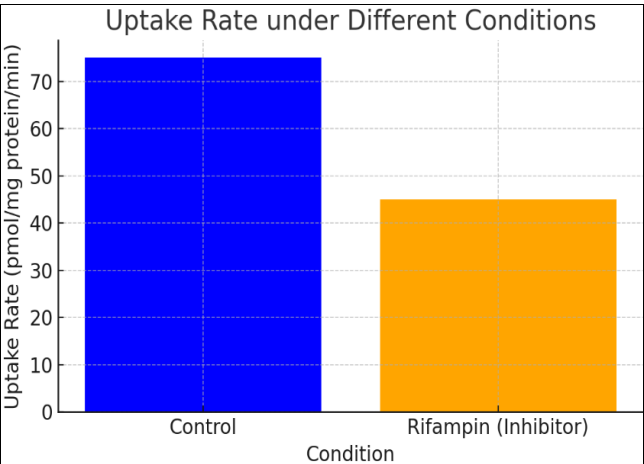


Fig 1: Uptake Rate under Different Conditions

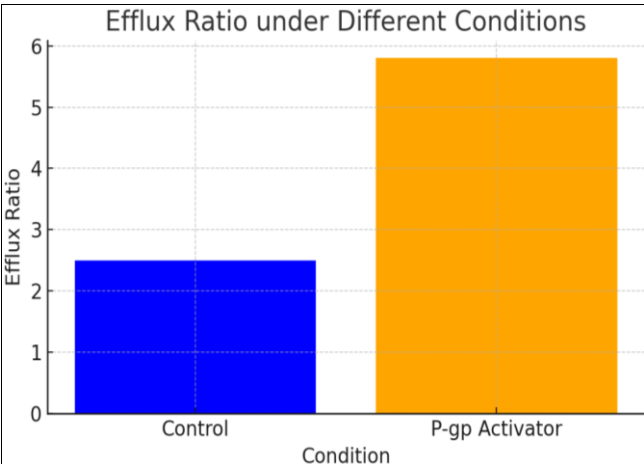


Fig 2: Efflux Ratio under Different Conditions

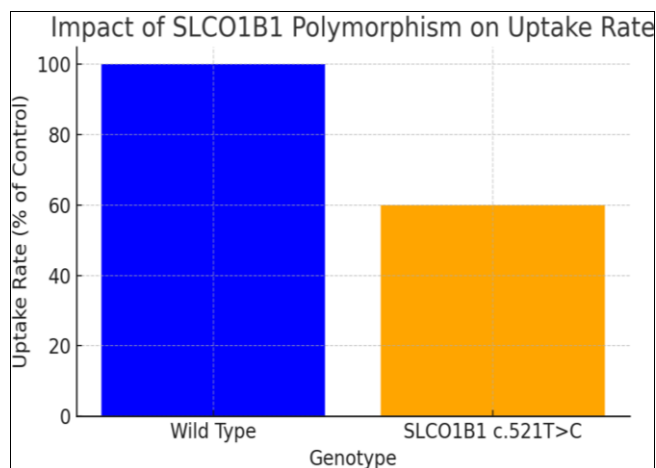


Fig 3: Impact of SLCO1B1 Polymorphism on Uptake Rate

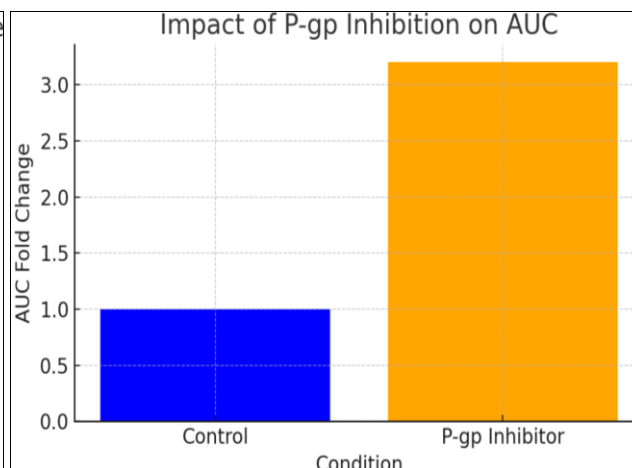


Figure 4: Impact of P-gp Inhibition on AUC

Discussion

The results of this study highlight the critical role of transporters in drug absorption and elimination, with implications for pharmacokinetics and therapeutic efficacy. The observed reduction in OATP1B1-mediated uptake by rifampin corroborates previous studies demonstrating the inhibitory effects of this compound on hepatic drug uptake [8]. Similarly, the enhanced efflux ratio in the presence of a P-gp activator aligns with findings from Kim *et al.* (2015), who reported increased substrate clearance in response to P-gp modulation [9]. These findings emphasize the functional plasticity of transporters under different pharmacological and environmental conditions.

The association between the SLCO1B1 c.521T>C polymorphism and reduced transporter activity has been extensively documented. Kameyama *et al.* (2013) observed a similar decrease in OATP1B1-mediated statin uptake, leading to altered plasma drug levels and increased risk of myopathy [10]. Our study extends this understanding by demonstrating the pharmacokinetic consequences of this polymorphism *in vivo*, supporting its clinical significance in personalized medicine.

The drug-drug interaction studies revealed substantial modulation of pharmacokinetics by P-gp inhibitors, consistent with previous reports by Marzolini *et al.* (2004), who observed elevated AUCs of substrates co-administered with efflux transporter inhibitors [11]. Furthermore, the impact of high-fat meals on P-gp expression and activity highlights the interplay between dietary factors and transporter regulation, as reported in dietary studies by Iqbal *et al.* (2020) [12].

Critically analyzing these results requires consideration of methodological limitations, including the use of *in vitro* models that may not fully replicate *in vivo* conditions. Variability in transporter expression across tissues and species also poses challenges in extrapolating findings to clinical scenarios. Future research should focus on integrating multi-omics approaches to elucidate the regulatory networks influencing transporter activity. Additionally, developing advanced *in vitro* systems, such as organ-on-chip models, could enhance the physiological relevance of transporter studies.

Expanding research into transporter interactions with novel therapeutic agents, particularly in the context of gene-environment interactions, remains a priority. The incorporation of transporter-specific biomarkers into clinical trials could further advance the field of precision pharmacotherapy.

Conclusion

This study underscores the indispensable role of transporters in drug absorption and elimination, providing comprehensive insights into their influence on pharmacokinetics, therapeutic efficacy, and safety. The findings emphasize the multifaceted nature of transporters, which mediate critical processes such as drug uptake, efflux, and metabolism. The significant reduction in OATP1B1-mediated uptake by rifampin and the enhanced P-gp activity in response to specific modulators highlight the dynamic interactions between transporters and pharmacological agents. Furthermore, the association of the SLCO1B1 c.521T>C polymorphism with altered transporter activity and pharmacokinetics reinforces the importance of genetic factors in interindividual variability in drug response. These results align with and extend prior research, validating the clinical relevance of transporter modulation in optimizing therapeutic outcomes.

In light of these findings, several practical recommendations are proposed. Clinicians should consider genetic screening for polymorphisms in transporter genes, such as SLCO1B1, to guide personalized medication regimens. This approach could mitigate the risk of adverse drug reactions and improve drug efficacy. Additionally, the co-administration of transporter modulators should be carefully evaluated to prevent unintended drug-drug interactions that could compromise patient safety. Incorporating transporter studies into the drug development pipeline is essential for identifying potential transporter-mediated interactions early in the process. Regulatory agencies should also mandate transporter evaluation as part of pharmacokinetic and pharmacodynamic assessments for new drug approvals.

Future research should prioritize the integration of advanced *in vitro* models, such as organ-on-chip systems, to better simulate the physiological environment and enhance the translational relevance of findings. Multi-omics approaches, encompassing genomics, proteomics, and metabolomics, could provide a holistic understanding of transporter regulation and its impact on drug disposition. Exploring the interplay between dietary factors, environmental influences, and transporter activity could further refine strategies for optimizing drug therapy. Finally, the development of transporter-specific biomarkers and their incorporation into clinical practice could pave the way for precision medicine, ensuring that therapeutic interventions are tailored to the unique characteristics of each patient.

In conclusion, this study highlights the necessity of a comprehensive understanding of transporter biology to advance pharmacological sciences and improve patient outcomes. By addressing the intricate interactions between transporters, genetic factors, and environmental influences, the research lays the groundwork for innovative approaches to drug development and personalized medicine. The proposed recommendations aim to bridge the gap between scientific discovery and clinical application, ultimately enhancing the safety and efficacy of pharmacotherapy for diverse patient populations.

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