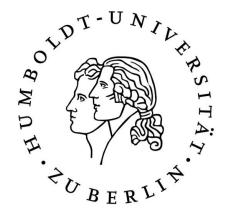
# Captopril in Focus: Open Pharmacokinetic Dataset and PBPK Modeling



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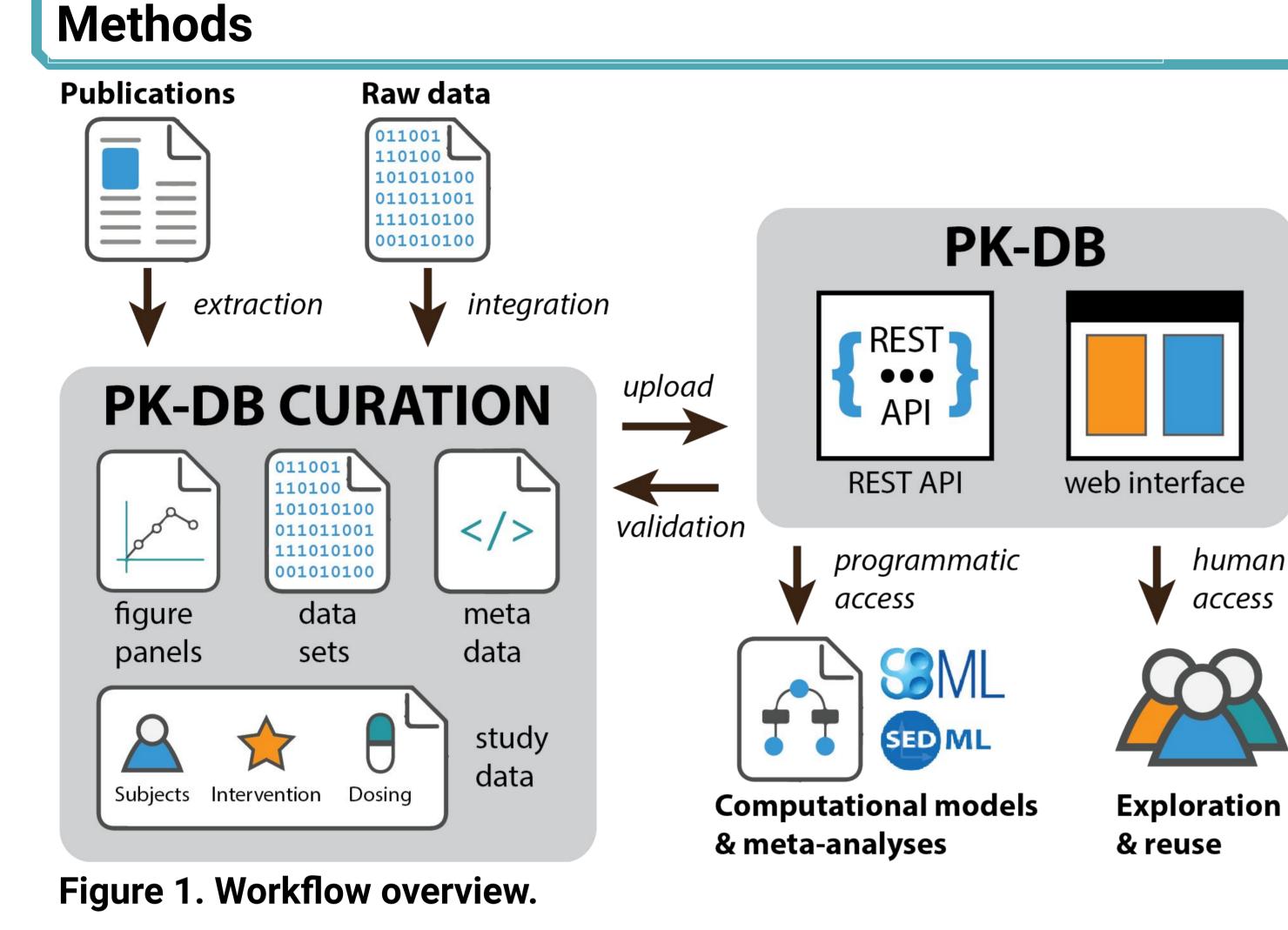
## Background

Our group has developed PK-DB [1], an open pharmacokinetics database from clinical and preclinical research. The objective of this study was to expand PK-DB with data on the angiotensin converting enzyme (ACE) inhibitor captopril, develop an open and FAIR [2] PBPK/PD model, and improve our understanding of the intraindividual variability in captopril pharmacokinetics.

## Results

Identifier		Data		Metabolites				Health					PK			PD RAAS			PD cardiovascular					
Study	PK-DB	pharmaco- kinetics	pharmaco- dynamics	all captopril	total captopril	captopril disulfide	S-methyl- captopril	healthy	hyper- tension	renal impairment	hepatic impairment	cardiac impairment	plasma	urine	feces	ACE activity	renin	aldosterone	arterial pressure	blood pressure	heart rate	cardiac output	PCWP	SVR
AlFuraih1991	PKDB00810	$\checkmark$	$\checkmark$					$\checkmark$					$\checkmark$											
Anuta2021	PKDB00811	$\checkmark$						$\checkmark$					$\checkmark$											
Arroyo1997	PKDB00748	$\checkmark$						$\checkmark$					$\checkmark$											
Cody1982	PKDB00812	$\checkmark$	$\checkmark$		$\checkmark$							$\checkmark$	$\checkmark$										$\checkmark$	$\checkmark$
Cohen1984	PKDB00813	$\checkmark$			$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$					$\checkmark$											
Creasey1986	PKDB00814	$\checkmark$			$\checkmark$		$\checkmark$	$\checkmark$					$\checkmark$	$\checkmark$										
Creasey1988	PKDB00815	$\checkmark$		$\checkmark$				$\checkmark$					$\checkmark$											
Drummer1987	PKDB00816	$\checkmark$	$\checkmark$		$\checkmark$				$\checkmark$	$\checkmark$			$\checkmark$							$\checkmark$				
Duchin1982	PKDB00817	$\checkmark$						$\checkmark$					$\checkmark$	$\checkmark$	$\checkmark$									
Duchin1984	PKDB00818	$\checkmark$								$\checkmark$			$\checkmark$	$\checkmark$	$\checkmark$									
Foster2009	PKDB00819	$\checkmark$			$\checkmark$			$\checkmark$					$\checkmark$											
Giudicelli1984	PKDB00820	$\checkmark$	$\checkmark$						$\checkmark$				$\checkmark$				$\checkmark$	$\checkmark$						
Giudicelli1987	PKDB00821	$\checkmark$	$\checkmark$		$\checkmark$				$\checkmark$				$\checkmark$				$\checkmark$			$\checkmark$	$\checkmark$			
Jankowski1995	PKDB00822	$\checkmark$						$\checkmark$					$\checkmark$											
Jehanli1996	PKDB00823	$\checkmark$						$\checkmark$					$\checkmark$											
Kok1997	PKDB00824	$\checkmark$			$\checkmark$			$\checkmark$					$\checkmark$											
Kripalani1980	PKDB00825	$\checkmark$		$\checkmark$		$\checkmark$		$\checkmark$					$\checkmark$	$\checkmark$	$\checkmark$									
Li1996	PKDB00826	$\checkmark$											$\checkmark$											
McElnay1995	PKDB00827	$\checkmark$	$\checkmark$					$\checkmark$					$\checkmark$											
McElnay1996	PKDB00828		$\checkmark$														$\checkmark$				$\checkmark$			
Ohman1985	PKDB00829	$\checkmark$	$\checkmark$		$\checkmark$				$\checkmark$				$\checkmark$	$\checkmark$			$\checkmark$							
Onoyama1981	PKDB00830							$\checkmark$					$\checkmark$											
Rezende2007	PKDB00831	$\checkmark$											$\checkmark$											
Richer1984	PKDB00832		$\checkmark$						$\checkmark$				$\checkmark$				$\checkmark$				$\checkmark$			
Salem2005	PKDB00833							$\checkmark$					$\checkmark$											
Schaefer1998	PKDB00834		$\checkmark$																					
Shaw1985	PKDB00835																				$\checkmark$			
Singhvi1982	PKDB00836							$\checkmark$						$\checkmark$	$\checkmark$									
Vancea2009	PKDB00837																							





**Over 600 publications** on captopril pharmacokinetics were reviewed through a systematic literature search, with 29 studies curated in PK-DB. The PBPK model was developed using a compartmental approach and encoded in Systems Biology Markup Language (SBML) [3] for accessibility and reproducibility based on established workflows [4, 5].

Figure 3. Studies used in calibration of model parameters.

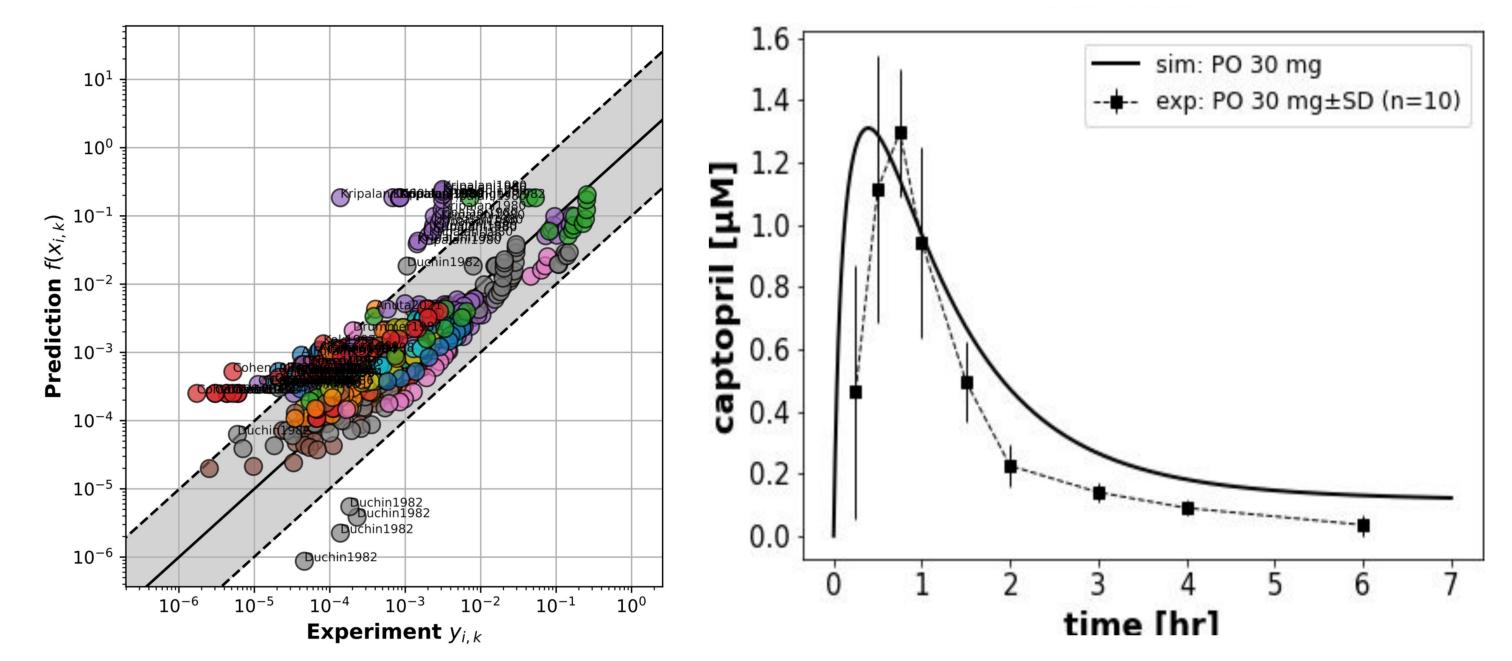
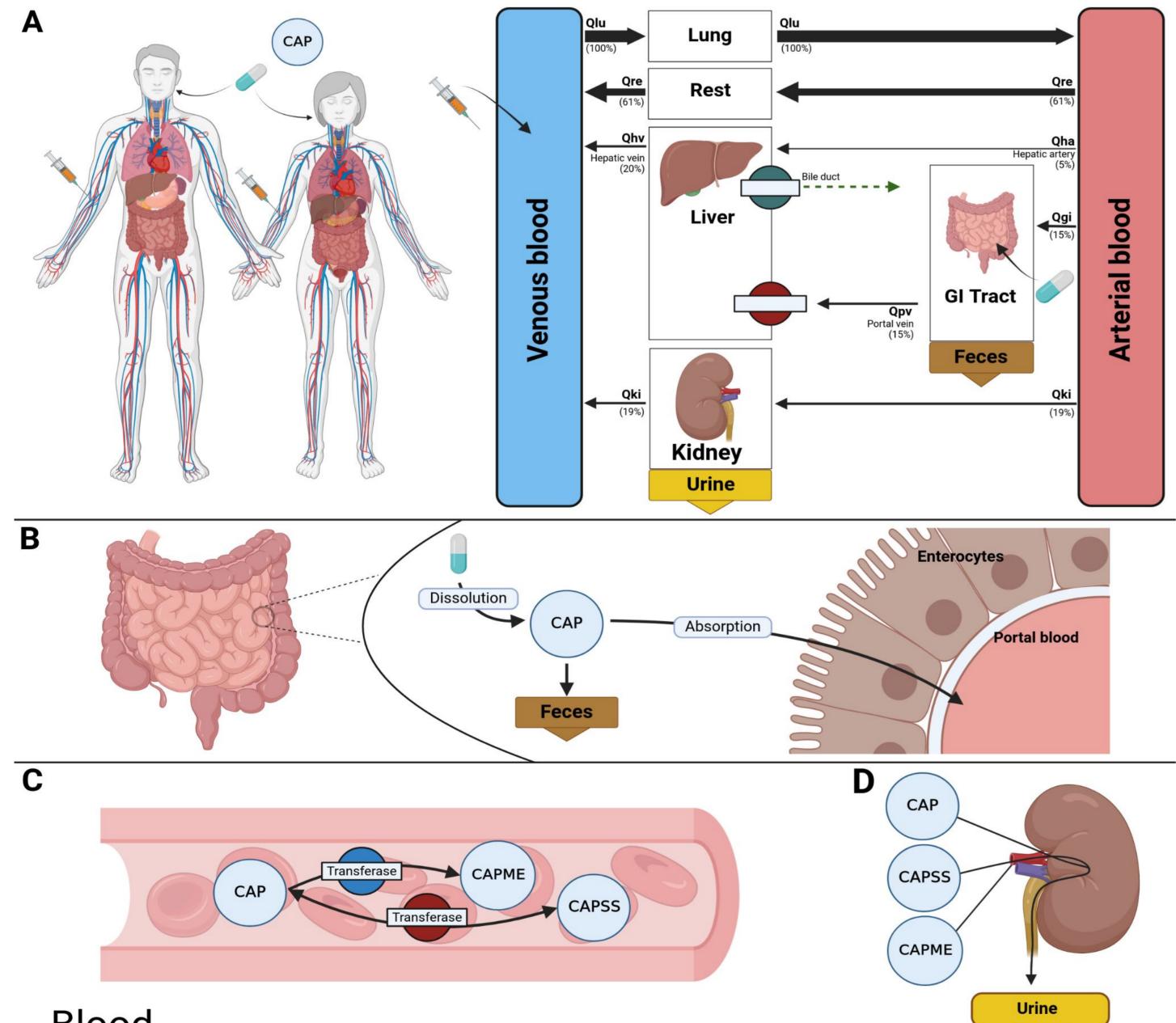


Figure 4. Parameter optimization (goodness of fit) & example prediction.



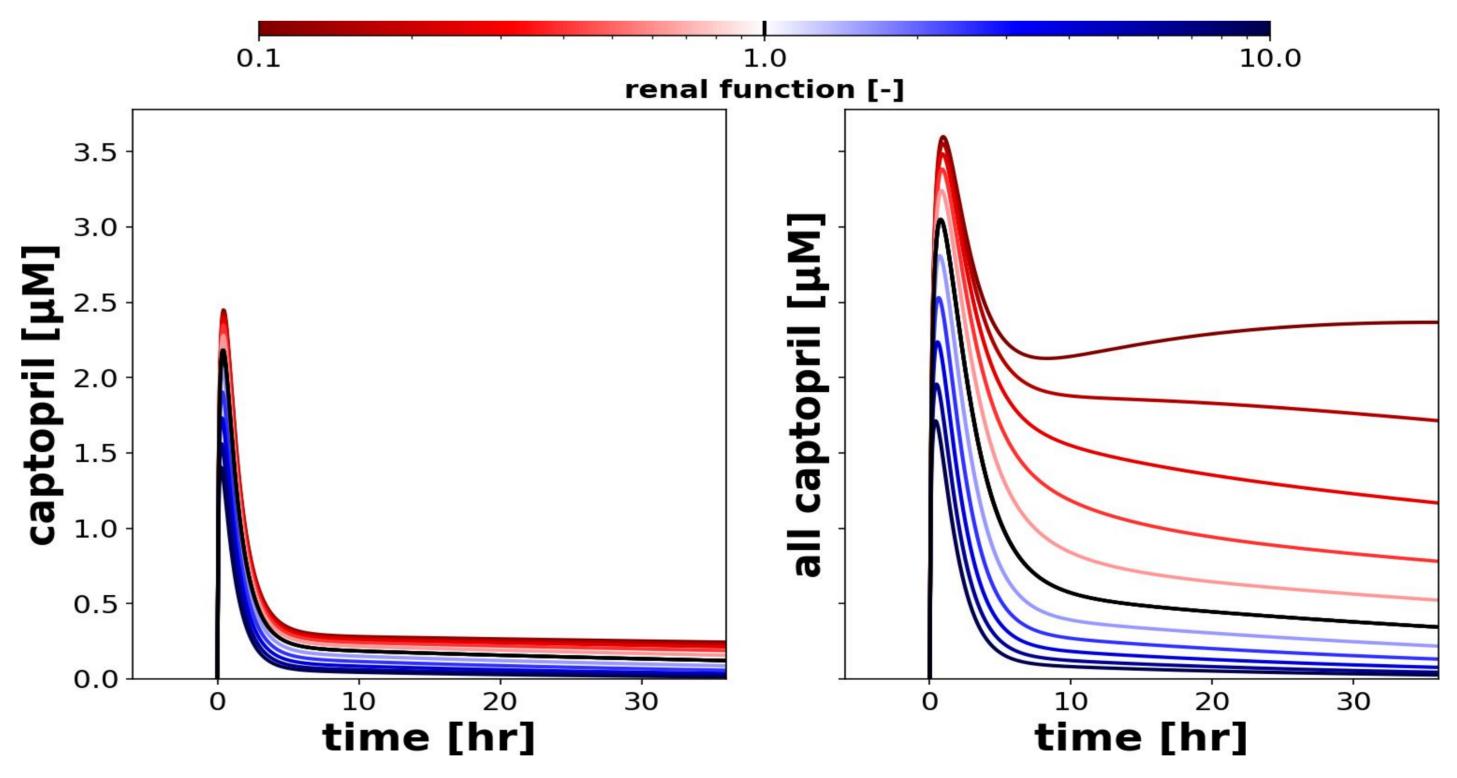
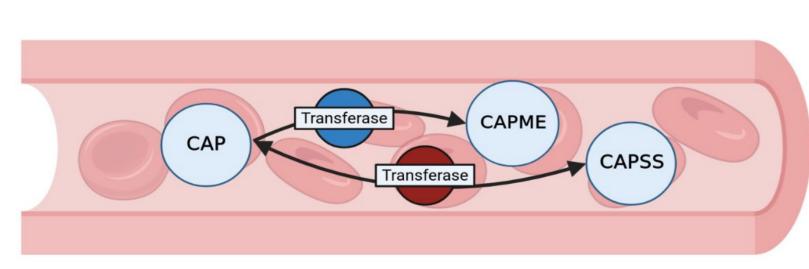


Figure 5. Effect of renal function on captopril pharmacokinetics.

## Conclusions

This study successfully establishes a freely accessible, comprehensive dataset and an open PBPK model of captopril, enhancing our understanding of its pharmacokinetic behavior across different patient groups. Our findings emphasize the



### Blood

Figure 2. A) Overview of PBPK captopril model. B) Intestinal absorption captopril (CAP). C) In blood CAP is converted to disulfides (CAPSS) and S-methyl captopril (CAPME). **D)** All captopril metabolites are excreted by kidneys, while disulfides also undergo hepatobiliary circulation.

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## significance of renal function in captopril elimination.

## References

[1] PK-DB: pharmacokinetics database for individualized and stratified computational modeling. Grzegorzewski et al., Nucleic Acids Res. 2021, <u>10.1093/nar/gkaa990</u>. [2] The FAIR Guiding Principles for scientific data management and stewardship. Wilkinson et al., Scientific Data 2016, <u>10.1038/sdata.2016.18</u>. [3] SBML Level 3: an extensible format for the exchange and reuse of biological models. Keating et al., Mol Syst Biol. 2020, <u>10.15252/msb.20199110</u> [4] Physiologically based pharmacokinetic (PBPK) modeling of the role of CYP2D6 polymorphism for metabolic phenotyping with dextromethorphan. Grzegorzewski et al., Front Pharmacol. 2022, <u>10.3389/fphar.2022.1029073</u>

[5] Pharmacokinetics of caffeine: A systematic analysis of reported data for application in metabolic phenotyping and liver function testing. Grzegorzewski et al., Frontiers in Pharmacology 2022, <u>10.3389/fphar.2021.752826</u>