



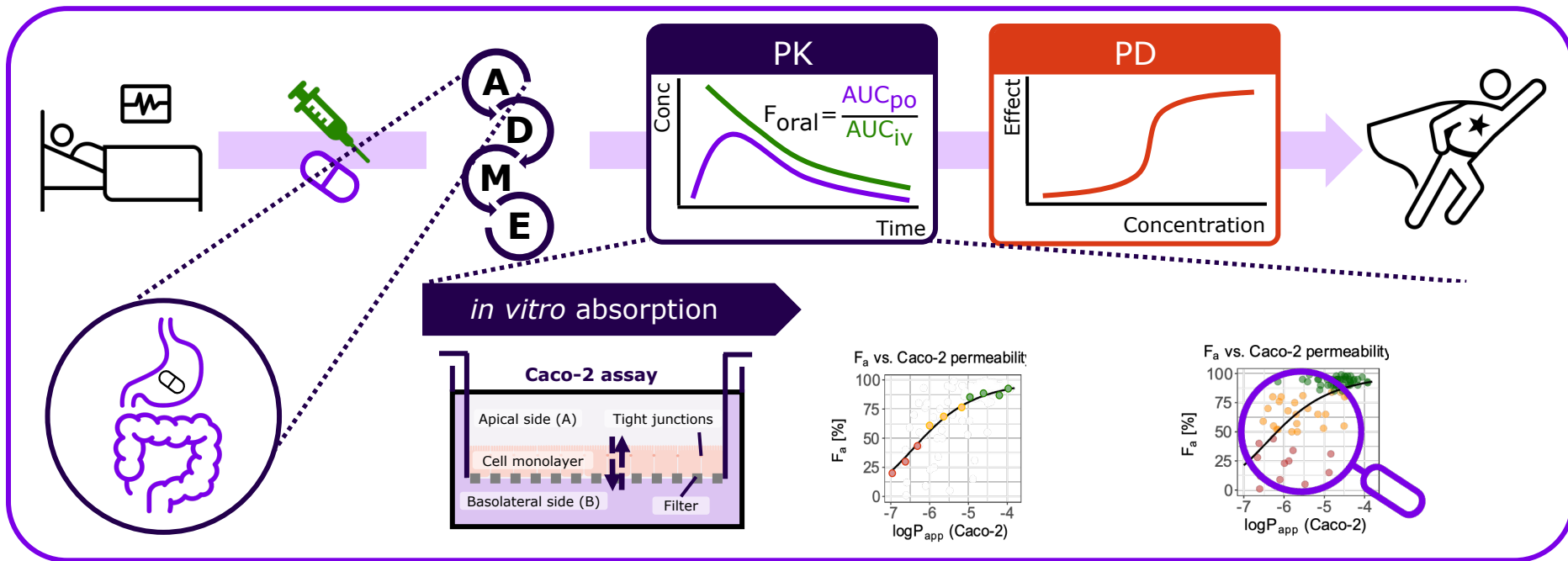
Prediction of human fraction absorbed from in vitro Caco-2 permeability – are we there yet?

OSP Community Conference, Basel, Oct 7-8, 2024
Session II: In Vitro-In Vivo Extrapolation

Denise Feick, DMPK Modeling & Simulation, Sanofi, Frankfurt



From **compound** to **cure**



Let's take a closer look!

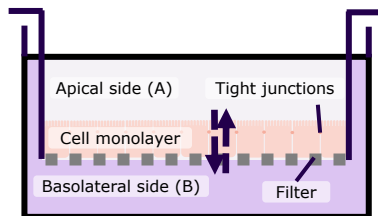


How can we inform intestinal absorption for MID3?

Examples for in vitro, in situ and in silico models



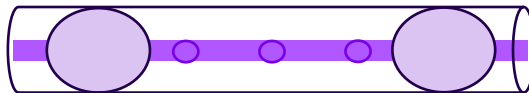
in vitro
- Cell systems -



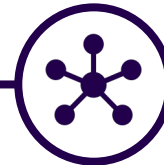
Caco-2 permeability assay^{1,2}



in situ
- Intestinal perfusion studies -



Loc-I-Gut technique³



in silico
- Regression models -



Quantitative Structure Property Relationship (QSPR) models⁴

¹ Sun et al. Expert Opin Drug Metab Toxicol. 2008;4(4):395-411

² van Breemen and Li. Expert Opin. Drug Metab. Toxicol. 2005, 1, 175-185

³ Lennernäs. Xenobiotica. 2007;37(10-11):1015-51.

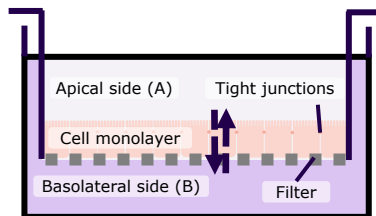
⁴ O'Shea et al. Eur. J. Pharm. Sci. 2022, 1:170:106098

How can we inform intestinal absorption for MID3?

In vitro Caco-2 permeability assay



in vitro - Cell systems -



Caco-2 permeability assay^{1,2}

SETUP

- Human colon epithelial cell line
- Cell monolayer with tight junctions, transporters and enzymes
- Compound added to apical (A) or basolateral (B) compartment (donor) and determined in opposite compartment (receiver)

COMMON READOUTS

1. Apparent permeability coefficient (P_{app})

Calculated from steady-state flux, surface area of the filter and initial compound concentration

2. Efflux ratio ($P_{app}B-A / P_{app}A-B$)

3. Absorptive or secretory quotient

Permeability in presence vs. in absence of specific inhibitor

4. Recovery (mass balance)

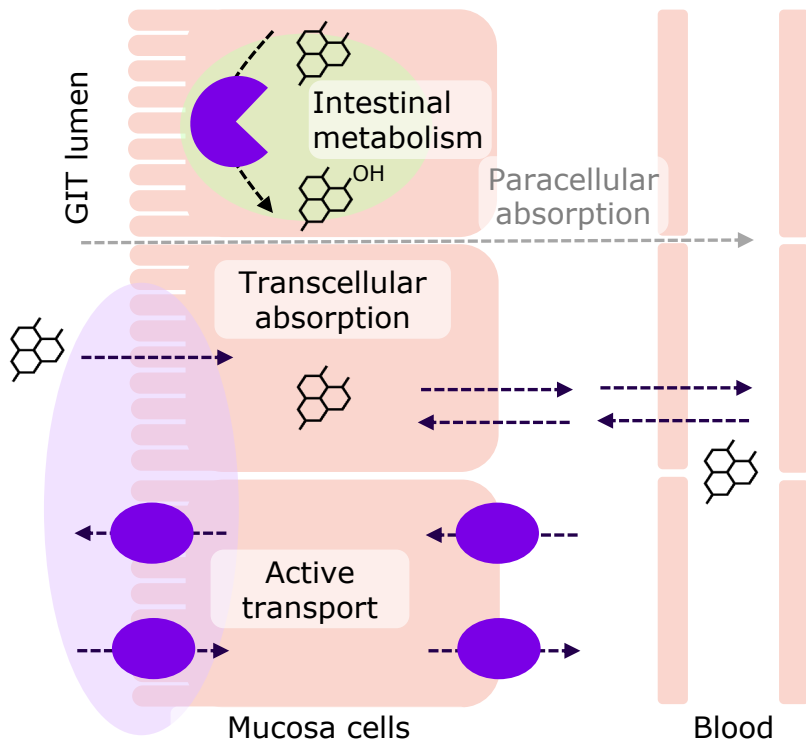
Compound in donor and receiver at end of experiment

¹ Sun et al. Expert Opin Drug Metab Toxicol. 2008;4(4):395-411

² van Breemen and Li. Expert Opin. Drug Metab. Toxicol. 2005, 1, 175-185

Intestinal absorption

High physiological complexity



$$F_{\text{oral}} = F_a \times F_g \times F_h$$

F_a

Fraction of a dose permeated from gut lumen into enterocytes

F_g

Fraction of a dose that escaped intestinal metabolism

F_h

Fraction of a dose that escaped hepatic metabolism

Key assumptions

- 1 Fraction absorbed \neq oral bioavailability
- 2 Fraction absorbed is the sum of **passive** and **transporter-mediated** membrane permeation

Intestinal absorption

Affected by various conditions

How is this represented in OSP Suite?



Physiological properties

- pH in lumen
- Feeding state
- Location of intestinal segment
- Intestinal transit time
- ...



Formulation properties

- Formulation dissolution
- Drug release
- Excipient effects
- ...



Drug properties

- Molecule size
- Solubility
- Lipophilicity
- Binding
- Metabolism
- Active transport
- ...

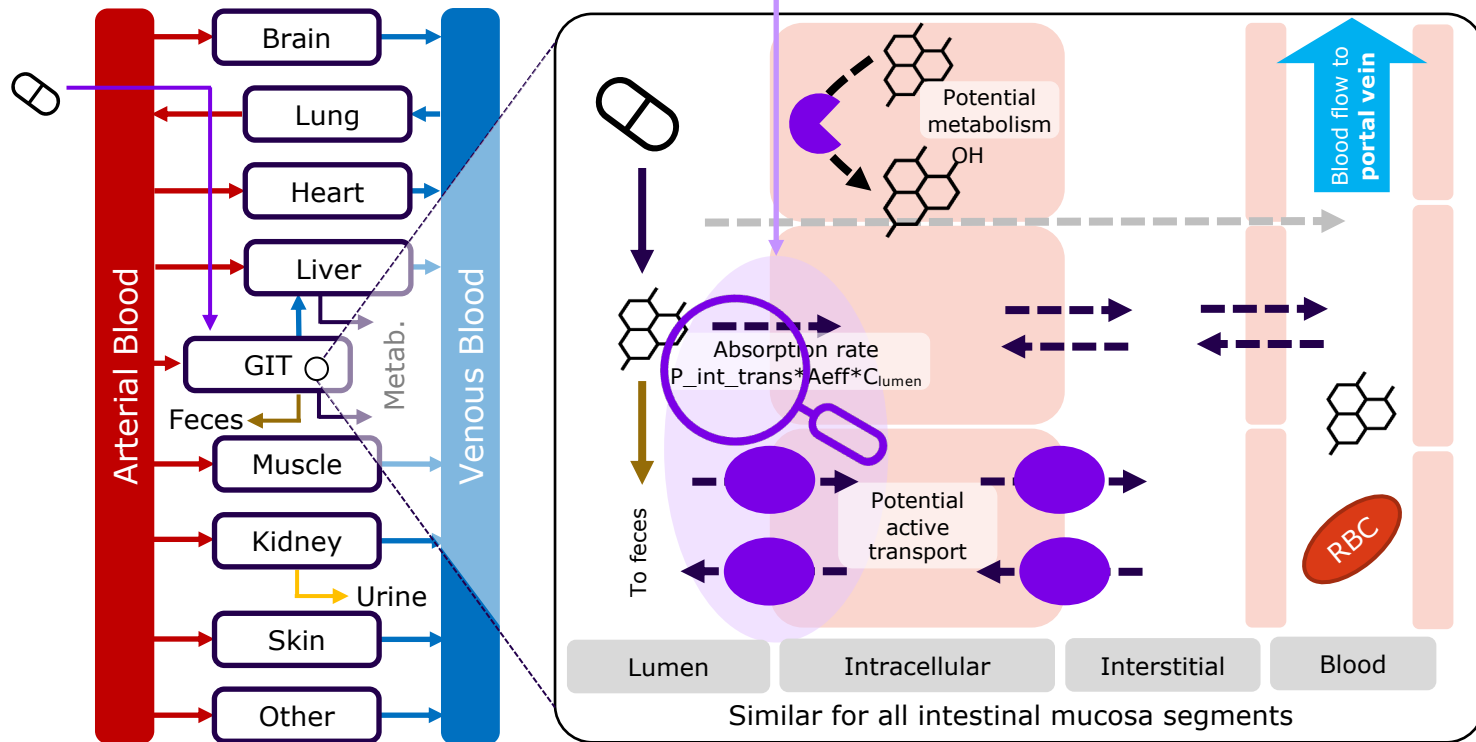
O'Shea et al. Eur. J. Pharm. Sci. 2022, 1:170:106098

Intestinal absorption

Representation in OSP Suite

$$F_{\text{oral}} = F_a \times F_g \times F_h$$

Fraction absorbed = Σ oral mass absorbed per mucosa segment divided by oral drug mass



Intestinal absorption

Representation in OSP Suite

PK-Sim Simulation

Simulation: 'Quinidine Example Simulation'

Parameters Reaction Diagram Time Profile Analysis

Filter

Scale 1.00 Reset

Molecule

Name	Value	Value Origin	Favorites
3-Hydroxyquinidine			
Quinidine			
Intestinal permeability (paracellular)	0 cm/min	Publication-Kuepfer, L., Niederalft, C., Wendt, T., Schl...	
Intestinal permeability (transcellular)	2.59E-5 cm/min		
P (blood cells->plasma)	7.00E-3 cm/min		

Description

Intestinal permeability via transcellular route

Formula

$$10^{(m * \log_{10}(P_{\text{int_InVitro}}) + b)}$$

References

b is defined as: Quinidine Example Simulation|Quinidine|Parameter b for correlation of intestinal permeability (transcellular)
m is defined as: Quinidine Example Simulation|Quinidine|Parameter m for correlation of intestinal permeability (transcellular)
P_int_InVitro is defined as: Quinidine Example Simulation|Quinidine|Specific intestinal permeability (transcellular)

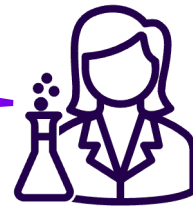
Default settings: $b = 0$, $m = 1$
 $\rightarrow P_{\text{int_trans}} = P_{\text{int_InVitro}}$

Defined in Compound building block
Default: Calculated from physicochemical properties according to Thelen et al. J Pharm Sci. 2011;100(12):5324-45

Intestinal absorption

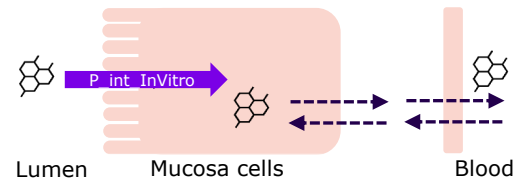
Representation in OSP Suite

Can we inform P_{int_trans} with *in vitro* permeability from Caco-2?



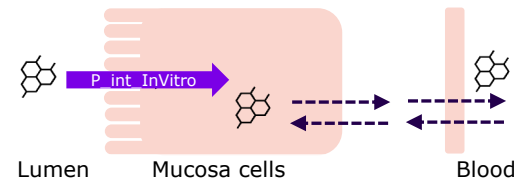
Scenario I: Default
 $P_{int_InVitro}$ calculated from physico-chemical properties

Intestinal permeability based on physico-chemical properties equals *in vivo* permeability
→ Unsaturable
→ *Only passive absorption considered*



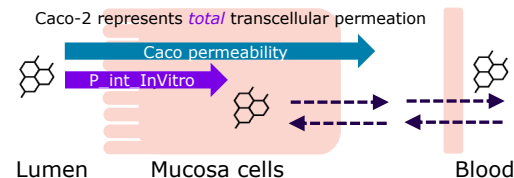
Scenario II: Common practice
User defined input for $P_{int_InVitro}$ → fitted

Intestinal permeability fitted, leaving out physico-chemical properties and active processes
→ Unsaturable
→ *Does not support bottom-up approach*



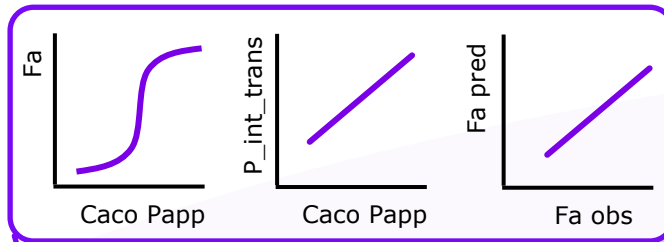
Scenario III: IVIVE
User defined input for $P_{int_InVitro}$ → Caco-2 based

Linear correlation *in vitro* permeability and *in vivo* permeability
→ Unsaturable
→ *Postulates that Caco-2 is representative for F_a*



Intestinal absorption

In Vitro-In Vivo Extrapolation



**Raw data
regression**

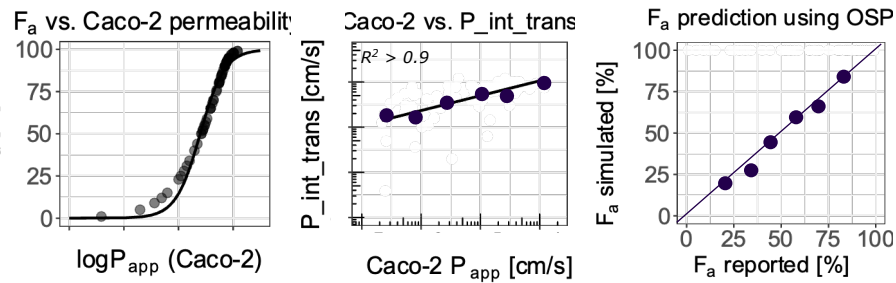
**Dataset
generation**

- *In vivo*, *in vitro* and *in situ* absorption
- Physicochemical and pharmacokinetic properties
- Calculated permeability (OSP, Thelen)
- Optimal permeability (OSP, estimated)

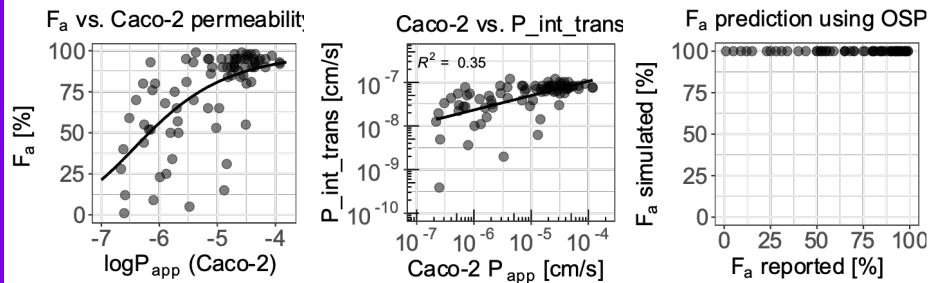
Correlation of Caco-2 permeability and fraction absorbed

Expectation vs. Reality

Expectation



Reality

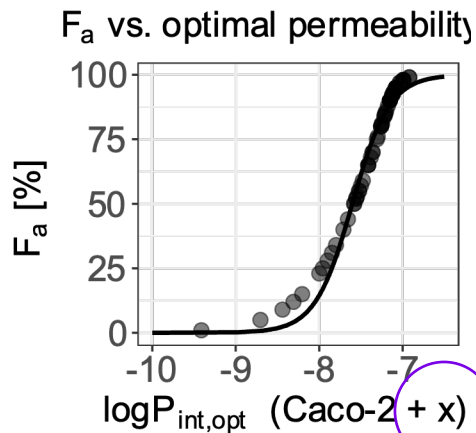


Be careful with literature statements about “perfect” correlation of Caco-2 permeability and fraction absorbed

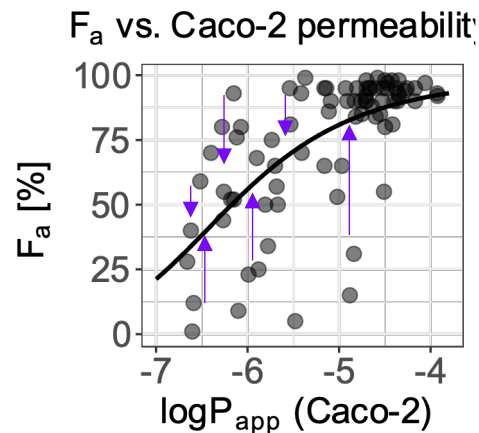
Correlation of Caco-2 permeability and fraction absorbed

Path to an ideal correlation using OSP Suite

Goal



Status quo

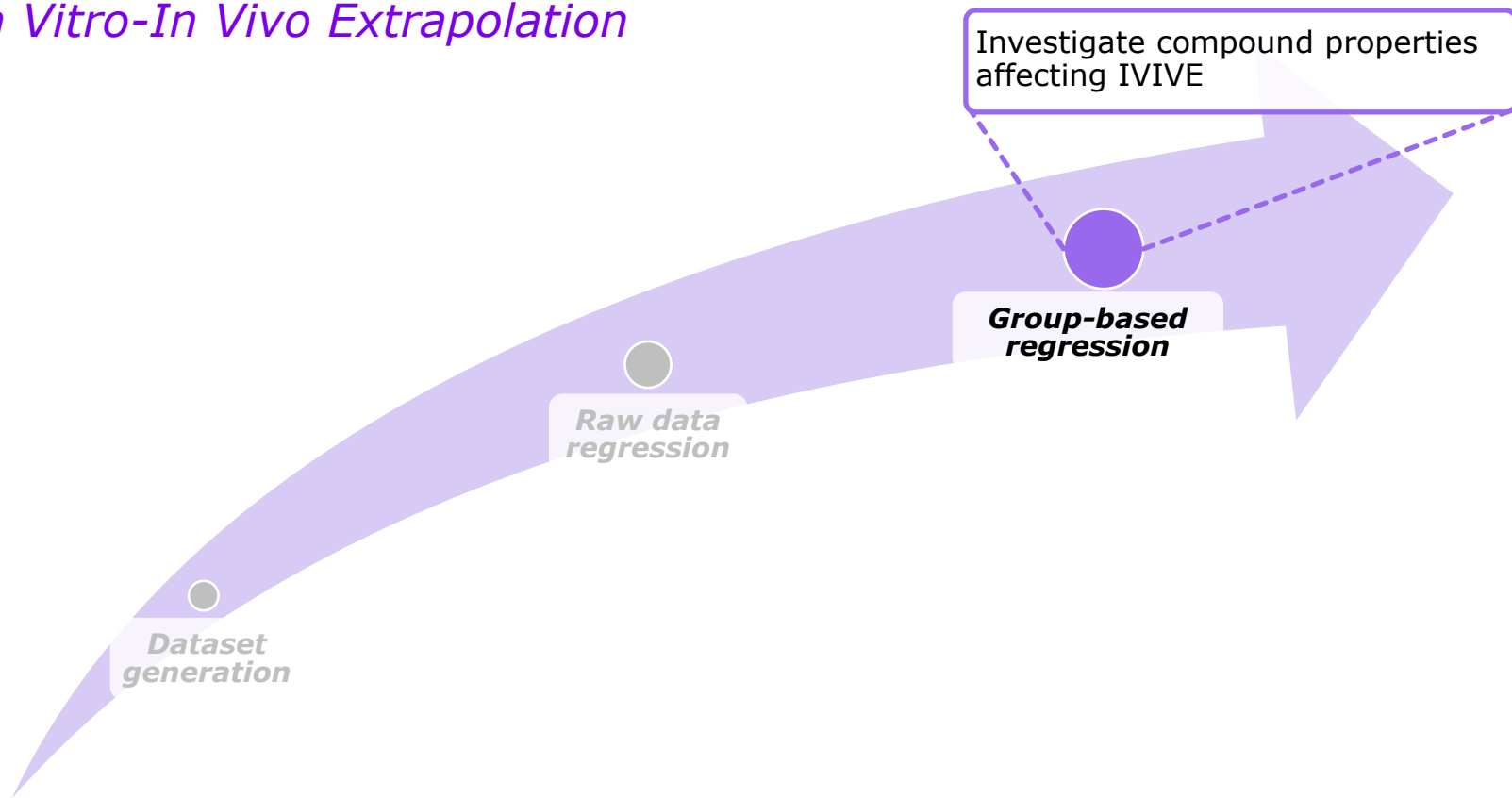


Can we systematically assess "X"?



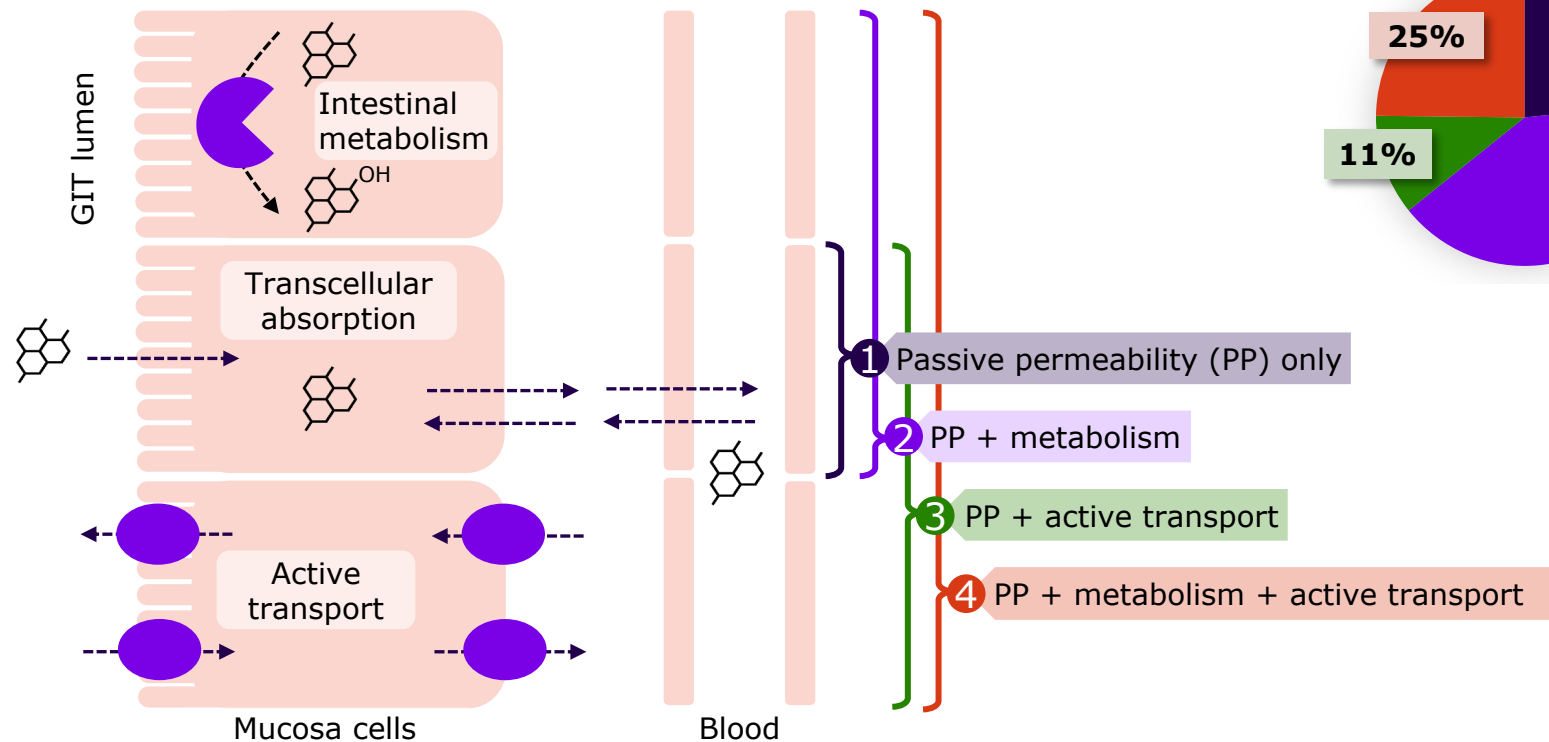
Intestinal absorption

In Vitro-In Vivo Extrapolation

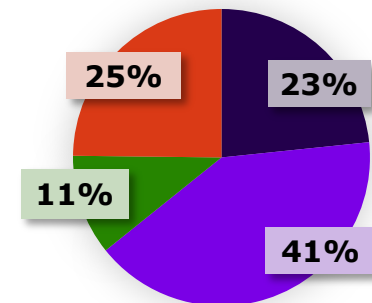


Benchmark dataset

Compound assignment to four ADME-related groups

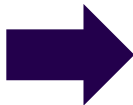
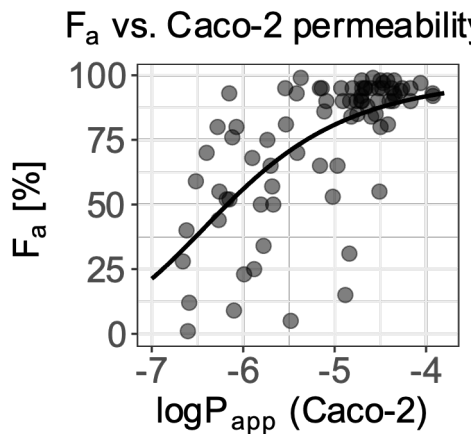


Distribution of ADME groups in dataset

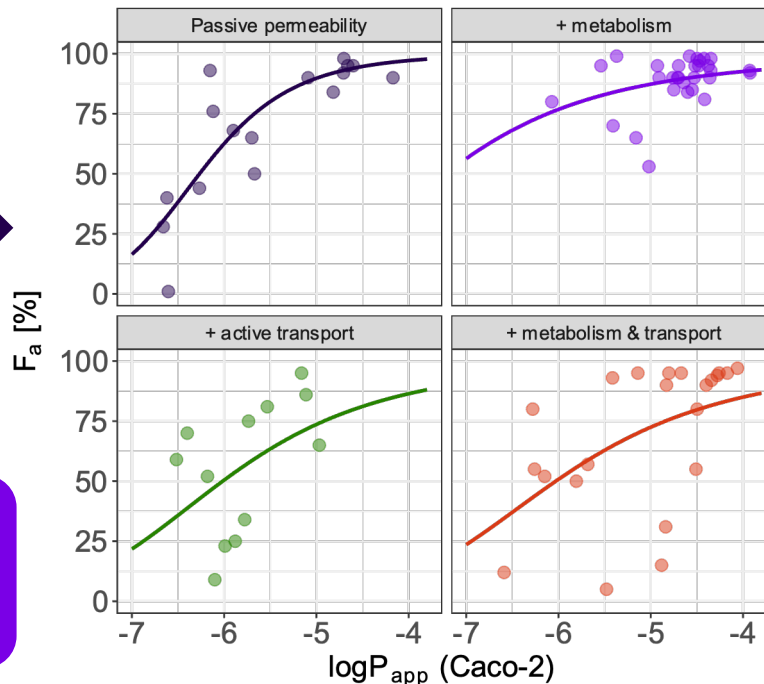


Benchmark dataset

Compound assignment to four ADME-related groups



F_a vs. Caco-2 permeability



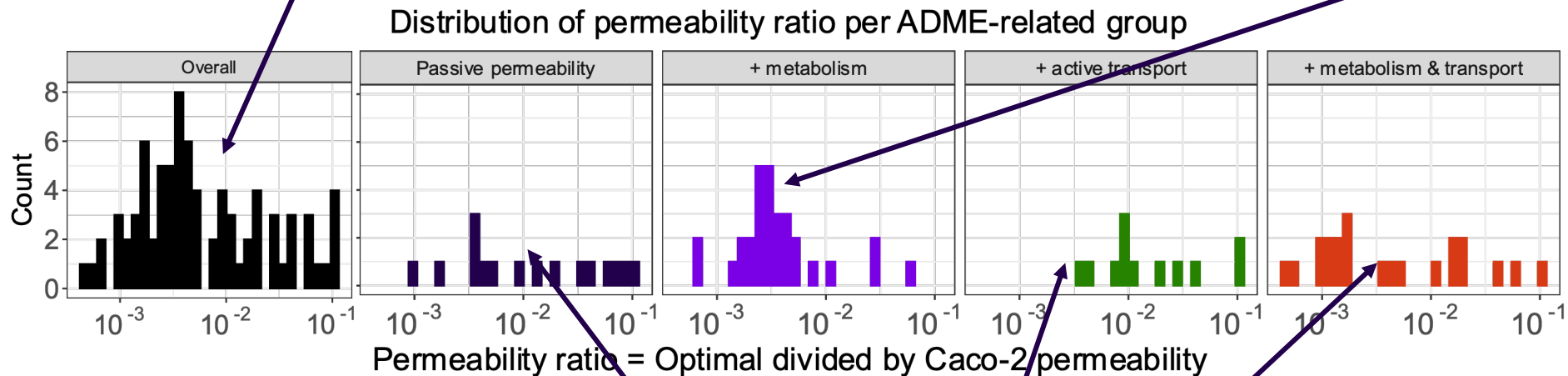
Knowledge about compound properties with respect to metabolism and active transport can help to improve prediction of fraction absorbed

In Vitro-In Vivo Extrapolation

Determination of an ADME-group specific correction factor

Same correction factor not appropriate for all compounds

Narrow distribution could indicate one correction value per group



Flat distribution could indicate that rather a correction function than a correction value is appropriate

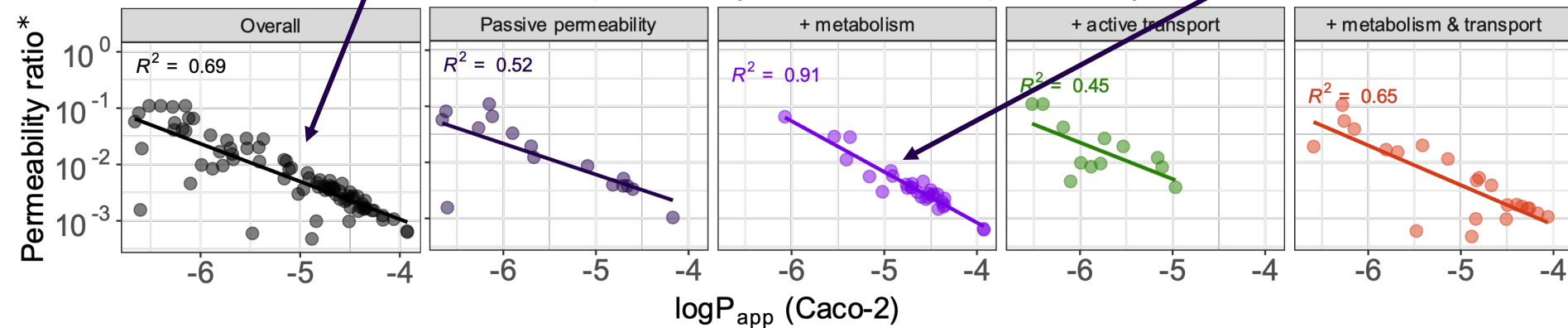
In Vitro-In Vivo Extrapolation

Determination of an ADME-group specific correction function

Overall larger gap between $P_{\text{int,opt}}$ and Caco-2 P_{app} identified for higher permeable compounds

Best correlation for metabolized compounds

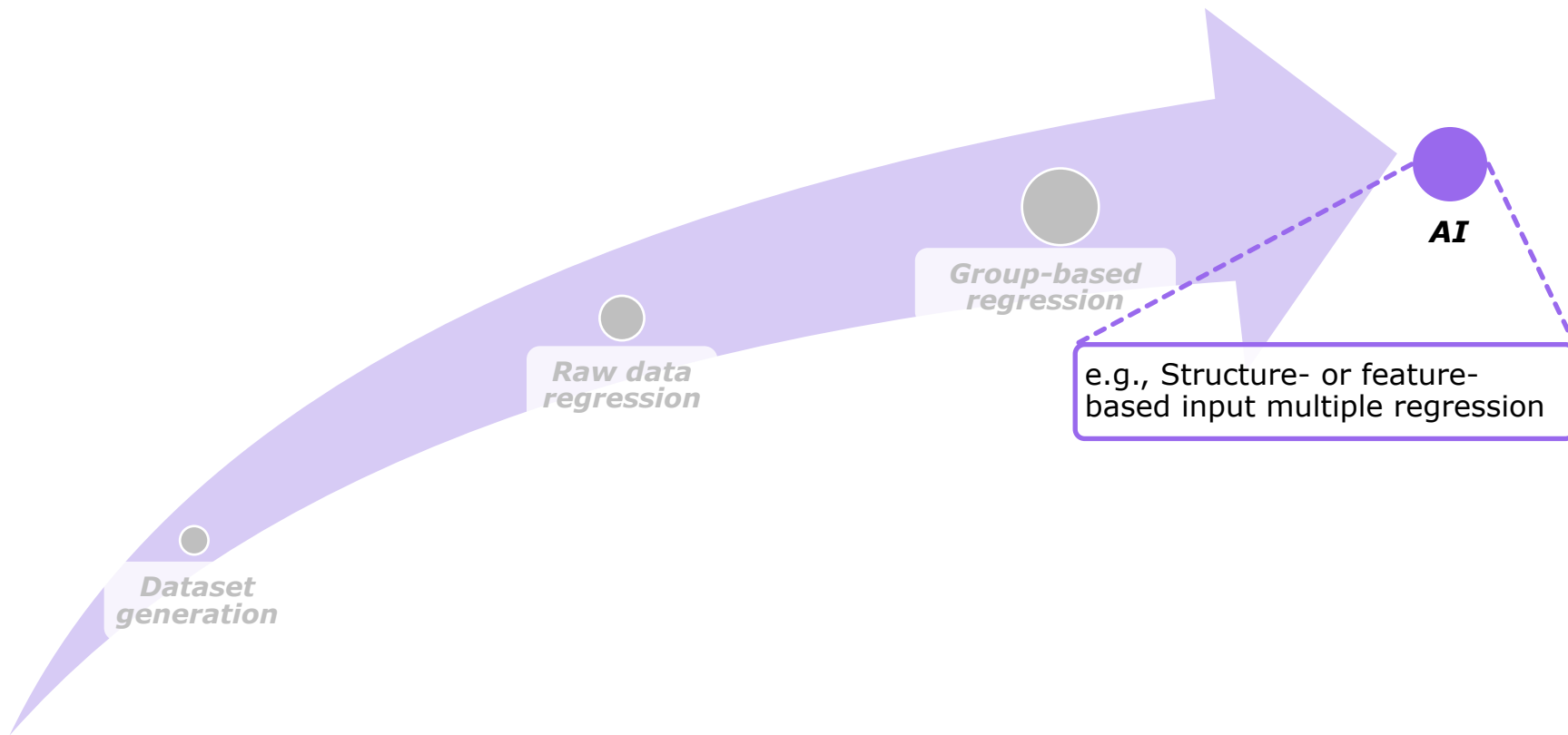
Correlation permeability ratio and Caco-2 permeability



Improved $P_{\text{int_trans}}$ and Caco-2 P_{app} correlation by incorporating function!

* Optimal divided by Caco-2 permeability

What comes next?



Prediction of human fraction absorbed from in vitro Caco-2 permeability – are we there yet?

Our analysis shows that **correction factors** and **functions** based on **active processes information** can **improve utilization of Caco-2 permeability** for IVIVE

How can we determine compound individual correction factors/functions?

- Can we integrate other DMPK assays to be more predictive?
- Can we systematically integrate non-linear ADME processes?
- Can we generate an optimal permeability with AI?

What does the OSP-Community think?



sanofi

Acknowledgements

OSP Focus Group IVIVE

Sanofi

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Thank you for
your attention!