



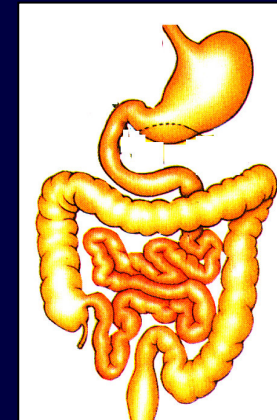
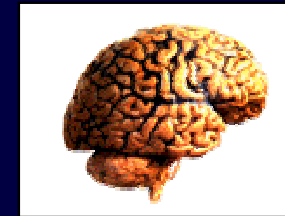
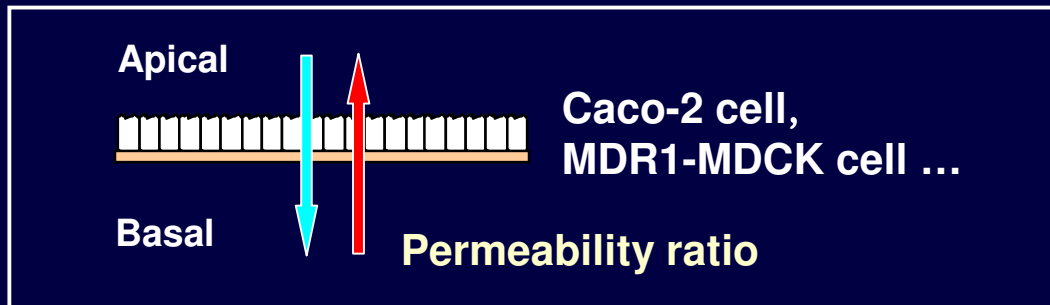
**Quantitative Evaluation of the Effect of
P-Glycoprotein on Oral Drug Absorption
-Assessment of Drug Permeability to Rat Small Intestine-**

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PURPOSE

Transport Experiments by using P-gp Expressing Cell Monolayers



$$\text{Transport Rate} = f (K_m, V_{\max}, P_{\text{passive}})$$

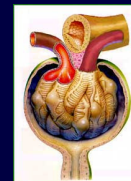
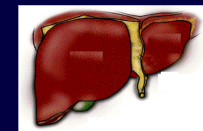
Drug Concentration

P-gp Expression Level

Affinity to P-gp

Passive Membrane Permeability

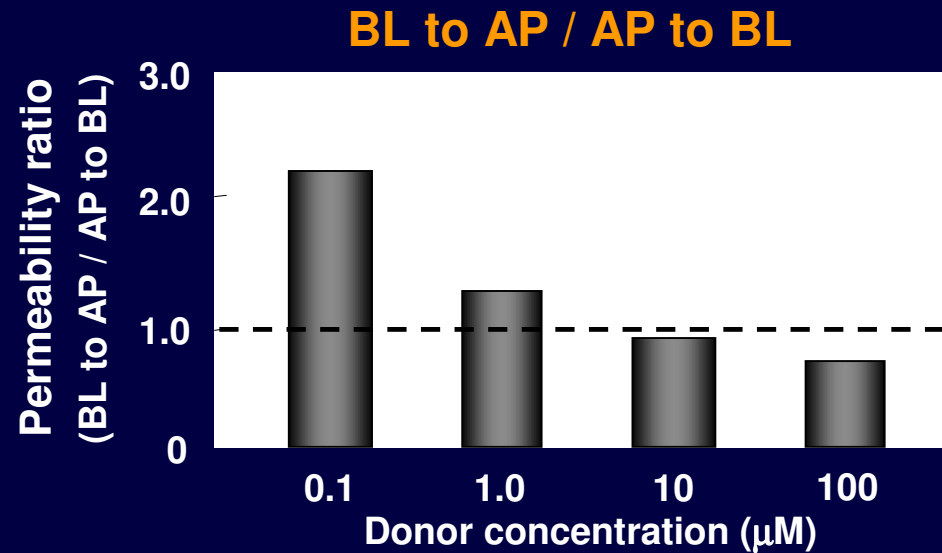
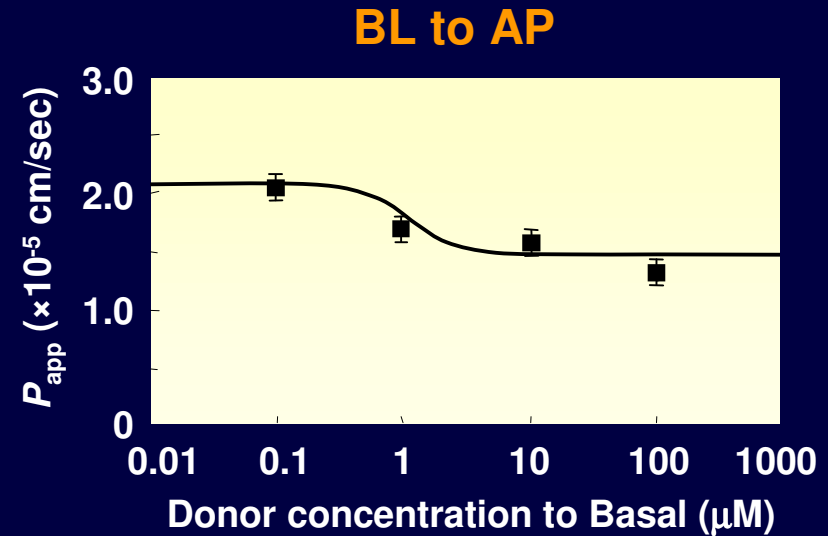
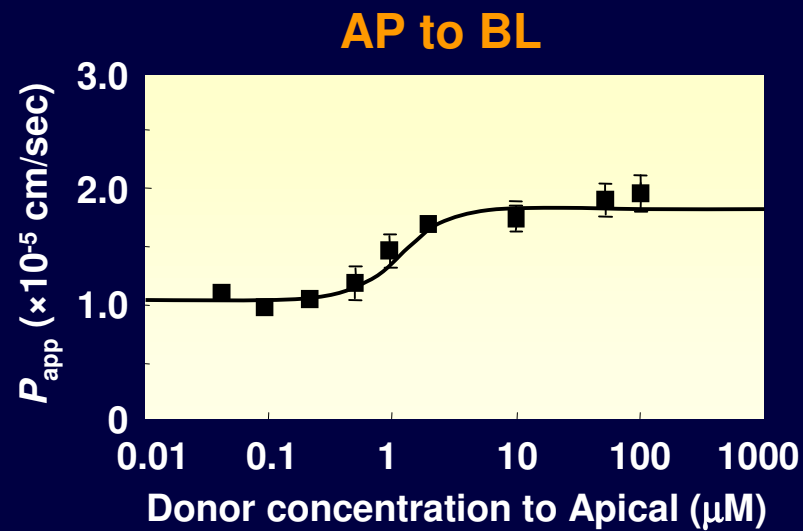
Cellular Accumulation



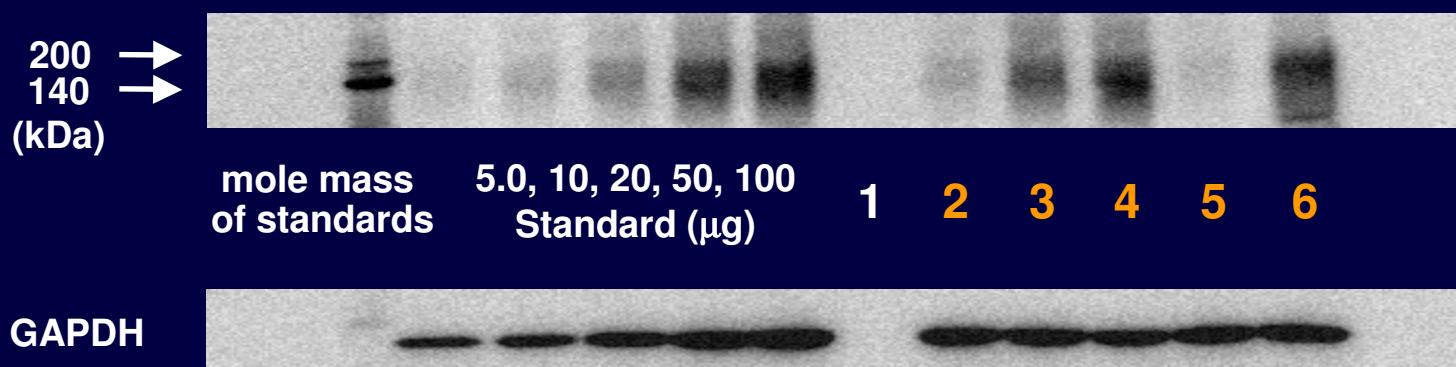
< Impact of P-gp on *in vivo* Oral Drug Absorption >

Development of kinetic model that can predict the *in vivo* absorption of P-glycoprotein substrate drugs from *in vitro* data.

Effect of donor concentration on AP to BL and BL to AP transport of quinidine in Caco-2 monolayers



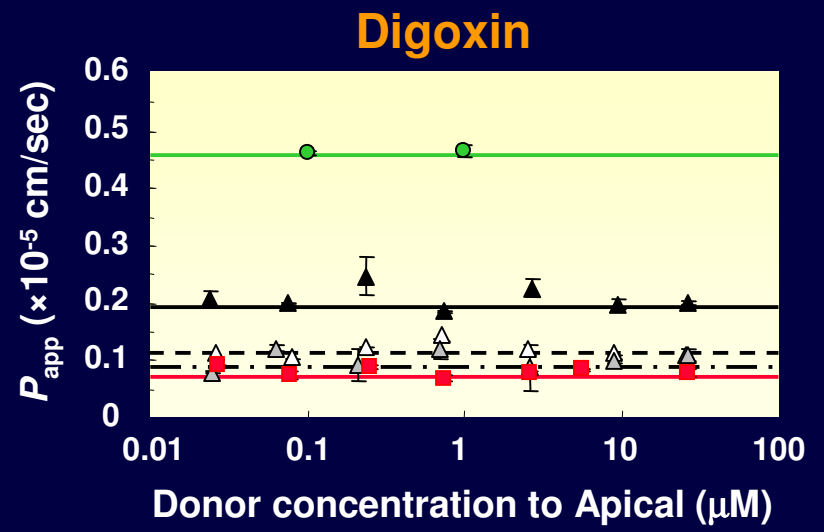
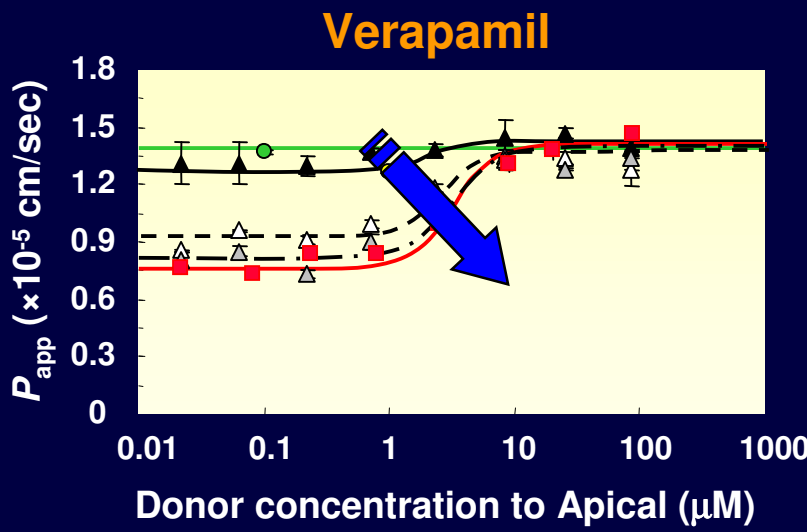
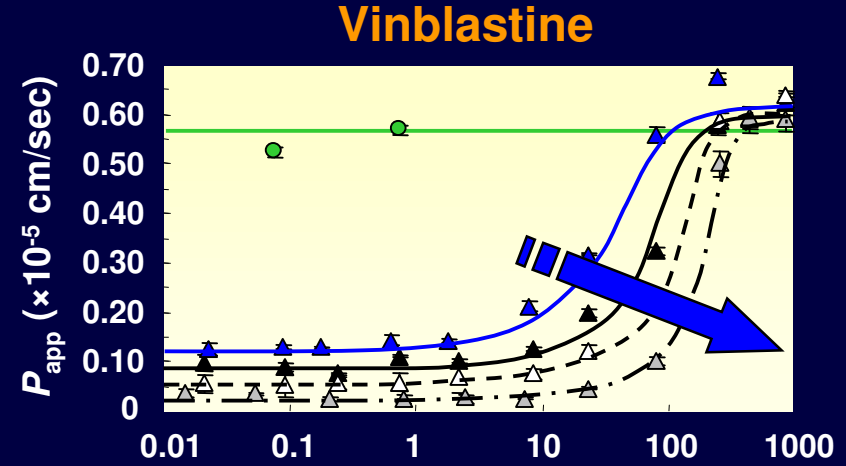
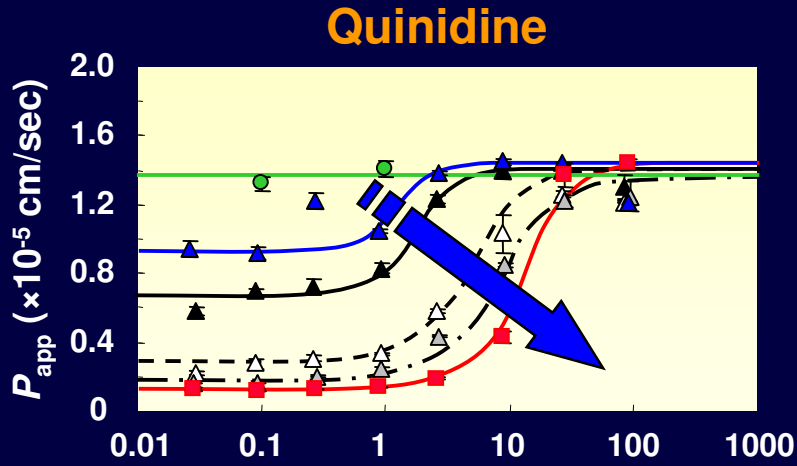
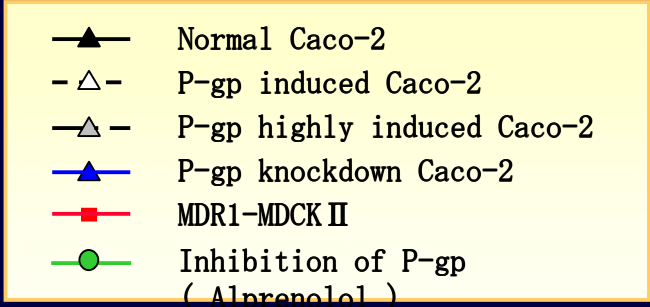
Various cell lines with different expression levels of P-gp



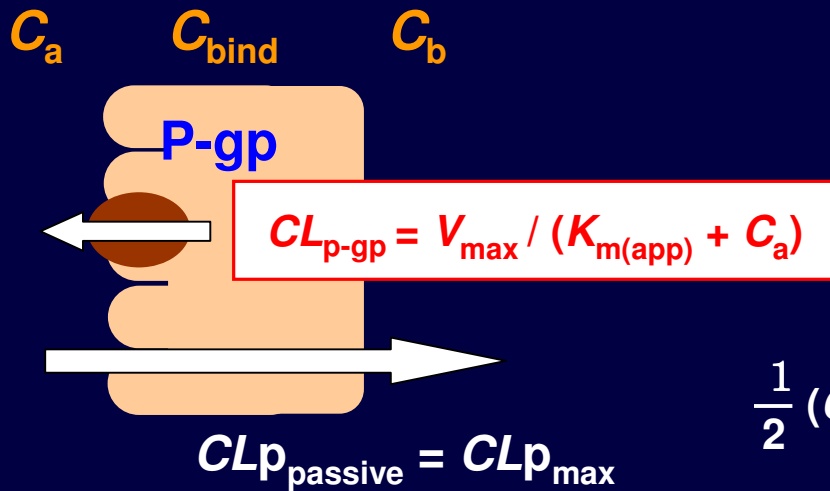
cell line	P-gp expression levels	
	Protein ($\mu\text{g}/\text{cm}^2$)	mRNA ($\times 10^{-5}/\text{GAPDH}$)
1 Non-cell (Blank)	0.00	0.00
2 Normal Caco-2 cells	26.8	3.22
3 P-gp-induced Caco-2 cells	103.7	5.20
4 P-gp-highly induced Caco-2 cells	191.0	9.53
5 MDR1-knockdown Caco-2 cells	8.71	1.32
6 MDR1-MDCKII cells	359.6	144.59

Protein levels were detected and quantified by Western blotting, loading total protein of 50 μg for cells and standard. mRNA levels were determined and quantified by Real-time quantitative PCR.

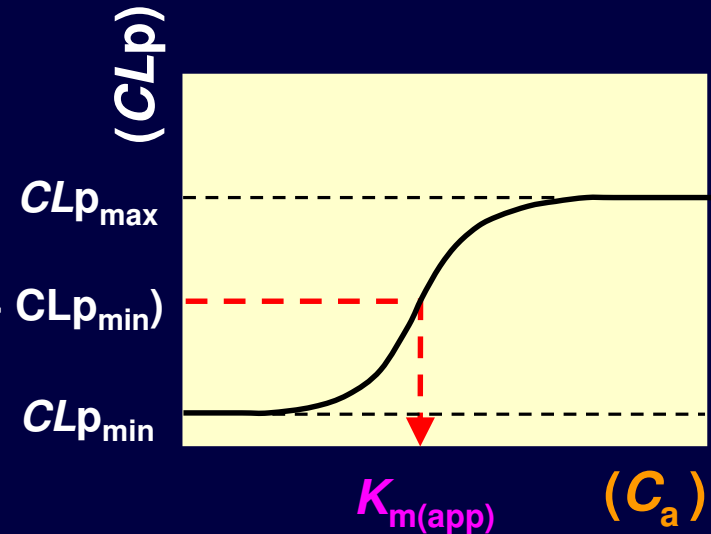
Effect of donor (apical) concentration on AP to BL transport of P-gp substrates in various cell monolayers



Analysis of P-gp function based on “Sigmoid E_{\max} model”



$$\frac{1}{2} (CL_{p_{max}} + CL_{p_{min}})$$




$$CLp = CL_{p_{max}} - CL_{P-gp}$$

Sigmoid E_{\max} model

$$\text{Flux rate} = CL_{p_{max}} \cdot C_a \cdot \frac{V_{max} \cdot C_a^r}{K_{m(app)}^r + C_a^r}$$

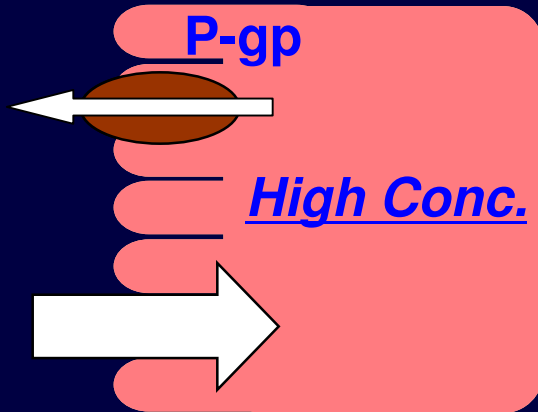
To add the flexibility, Hill Coefficient (r) was introduced to Sigmoid E_{\max} model

Estimation of $K_{m(app)}$, V_{max} and hill coefficient of three P-gp substrates by fitting CLp to Sigmoid E_{max} model

Cell line		MDR1-kd. Caco-2	Normal Caco-2	P-gp id. Caco-2	P-gp highly id. Caco-2	MDR1- MDCK II
P-gp expression level						
Quinidine	$K_{m(app)}$ (μM)	0.61	1.69	6.20	8.13	16.39
	V_{max} ($\times 10^{-7}$ $\mu\text{mol}/\text{min}/\text{cm}^2$)	3.00	7.74	41.0	63.0	132.5
	r (Hill coefficient)	1.16	0.98	1.00	1.01	1.02
Verapamil	$K_{m(app)}$ (μM)	—	1.01	1.66	2.07	2.85
	V_{max} ($\times 10^{-7}$ $\mu\text{mol}/\text{min}/\text{cm}^2$)	—	0.91	5.09	8.40	12.6
	r (Hill coefficient)	—	1.02	1.00	1.01	1.02
Vinblastine	$K_{m(app)}$ (μM)	29.88	80.69	149.01	323.35	—
	V_{max} ($\times 10^{-7}$ $\mu\text{mol}/\text{min}/\text{cm}^2$)	88.3	213.7	498.7	1180.9	—
	r (Hill coefficient)	1.01	1.01	1.00	1.01	—

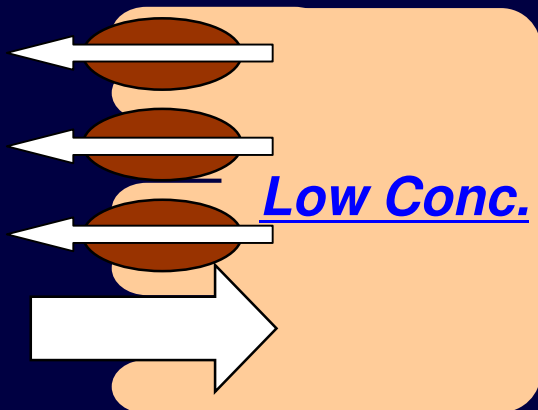
Why does " $K_{m(app)}$ " value fluctuate depending on the expression level of P-gp in cells

Lower expression level

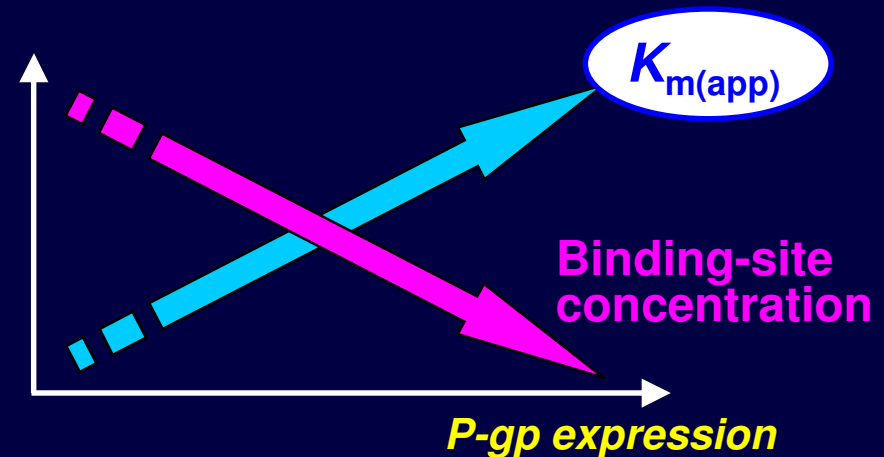


Drug concentration at drug binding-site of P-gp increase easily when the apical concentration become high.

Higher expression level



Drug concentration at drug binding-site of P-gp is kept low even when the apical concentration become high.

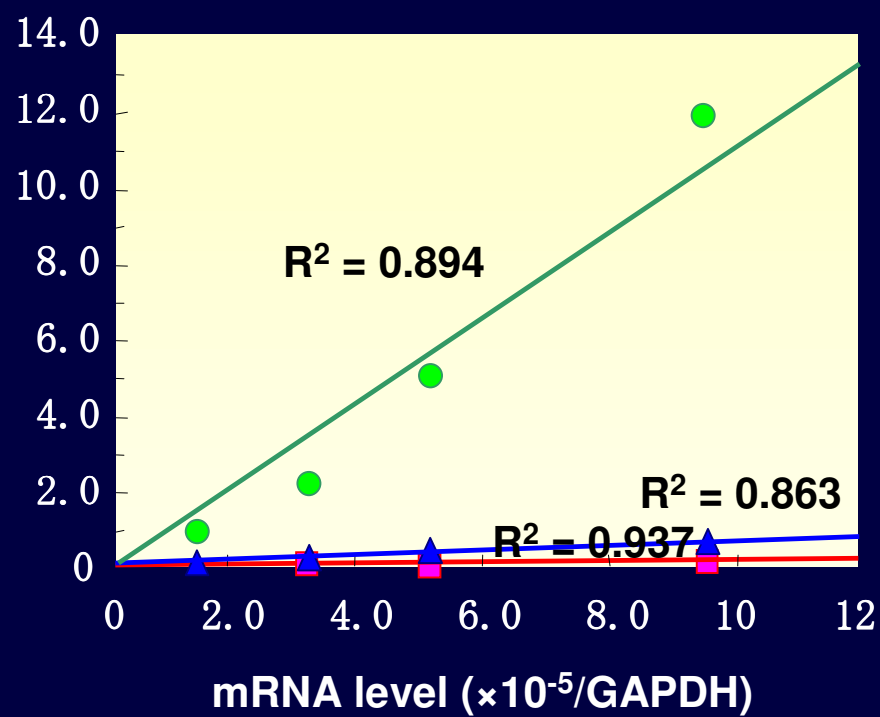
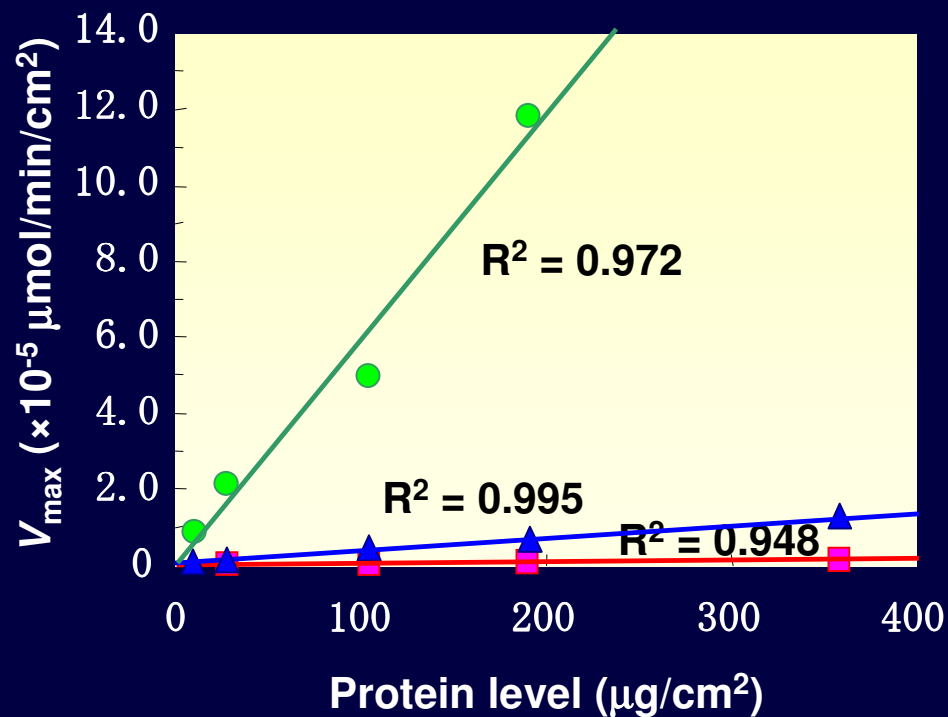


Correlation between P-gp expression level and “ V_{max} ” value of three P-gp substrates

- ▲ Quinidine
- Verapamil
- Vinblastine

V_{max} versus Protein

V_{max} versus mRNA

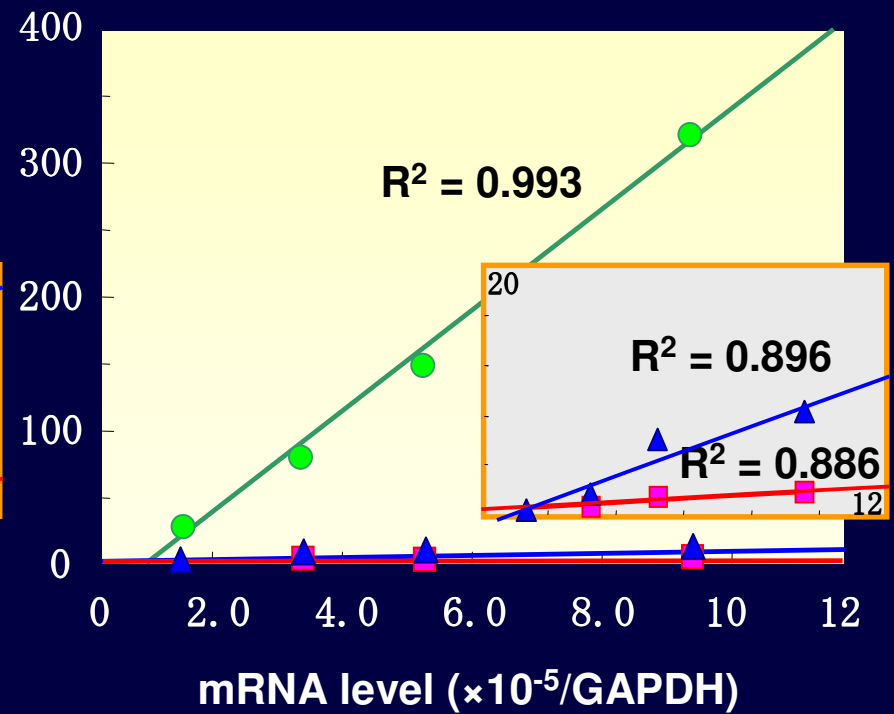
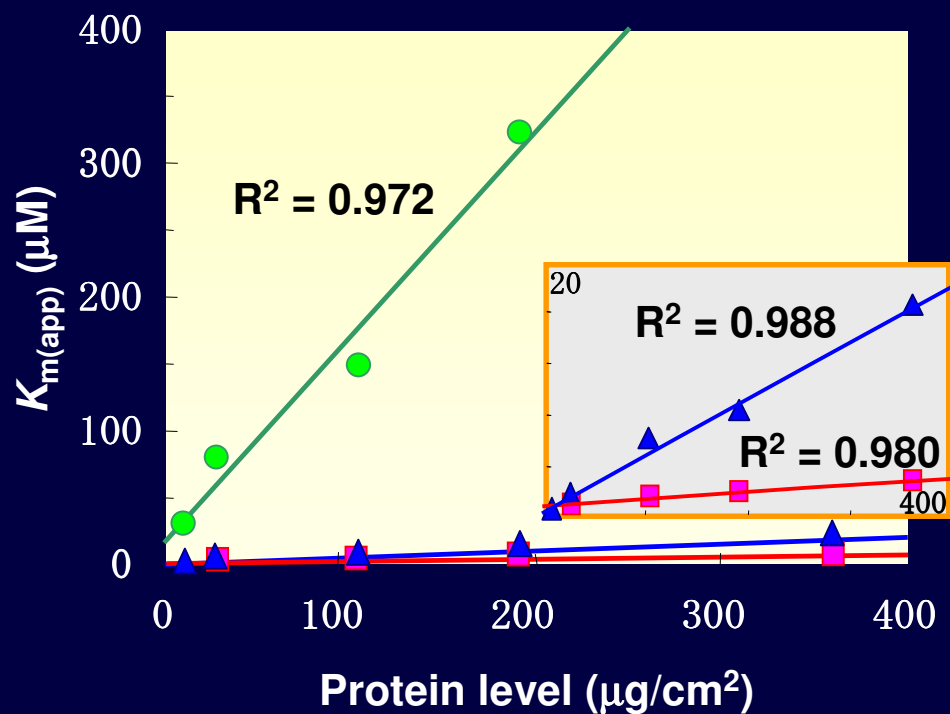


Correlation between P-gp expression level and “ $K_{m(app)}$ ” value of three P-gp substrates

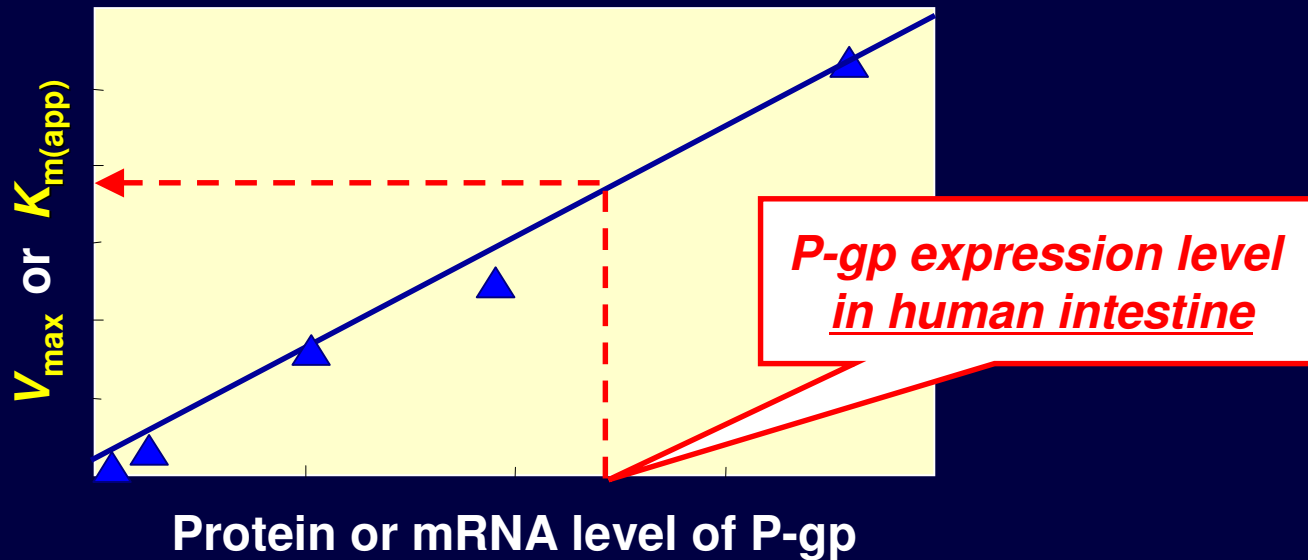
- ▲ Quinidine
- Verapamil
- Vinblastine

$K_{m(app)}$ versus Protein

$K_{m(app)}$ versus mRNA



How to estimate P-gp-mediated efflux in human intestine? How to predict the permeability in human intestine?



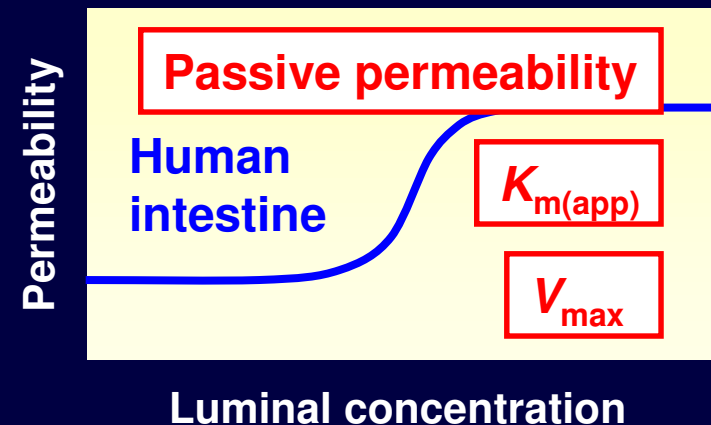
(1) Passive permeability :

Possible to be predicted from Caco-2 data by using P-gp inhibitor.

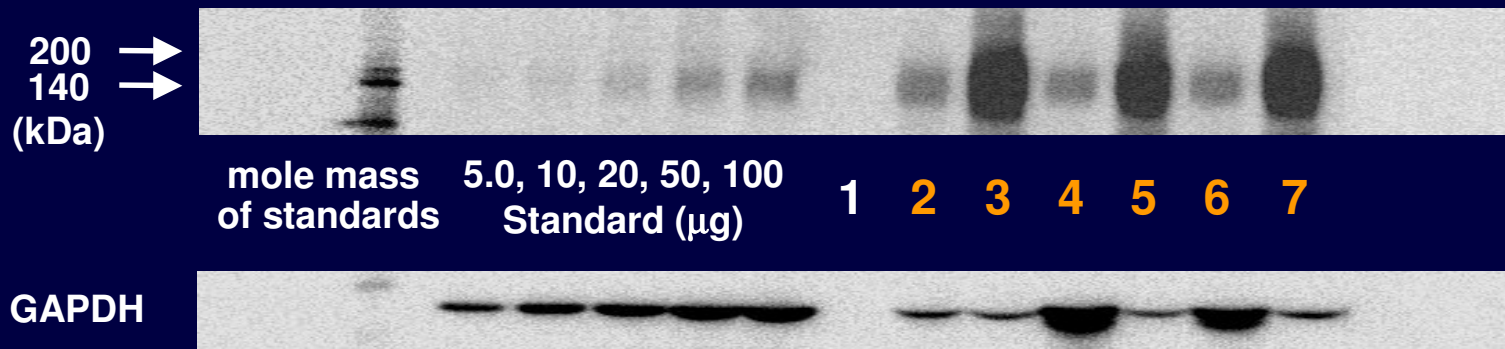
(2) P-gp mediated efflux :

$$CL_{\text{p-gp}} = V_{\max} / (K_{m(\text{app})} + C_a)$$

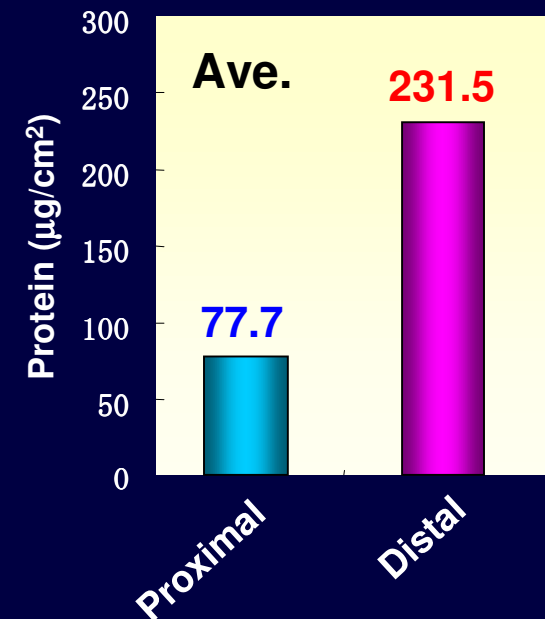
In vivo permeability



Regional difference in P-gp expression levels in rat small intestine (proximal and distal)

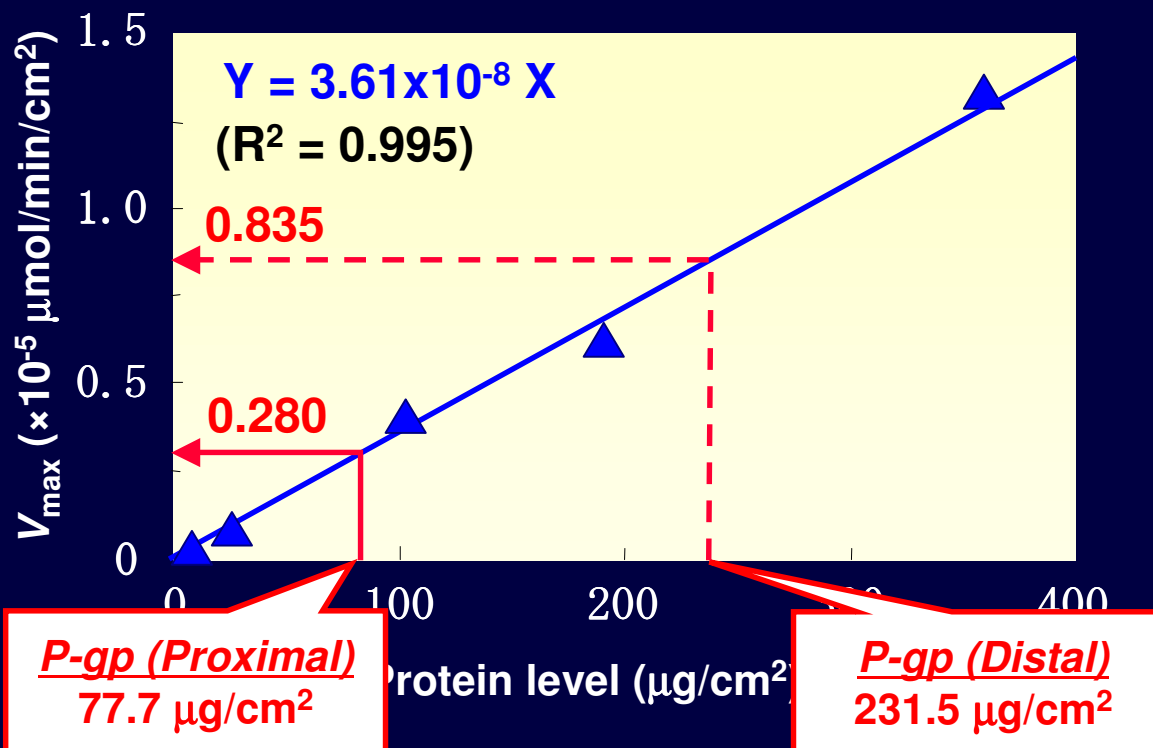


Rat # / Region		P-gp expression levels	
		Protein (µg/cm ²)	
Rat #1	1	Blank	0.00
	2	Proximal	89.8
	3	Distal	291.7
Rat #2	4	Proximal	65.7
	5	Distal	166.8
Rat #3	6	Proximal	77.6
	7	Distal	235.9



Protein levels were detected and quantified by Western blotting, loading 100µg for rat BBM; 50µg for standard.

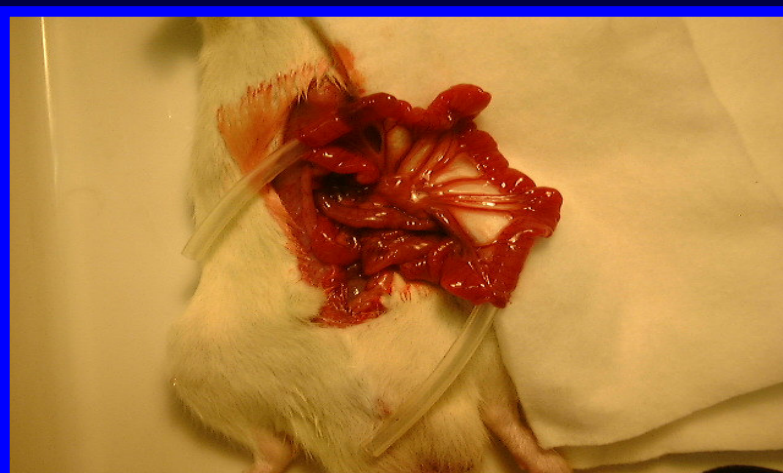
Estimation of “ V_{max} ” and “ $K_{m(app)}$ ” values of quinidine in rat small intestine (proximal and distal)



Quinidine		V_{max} ($\times 10^{-5}$ $\mu\text{mol}/\text{min}/\text{cm}^2$)	$K_{m(app)}$ (μM)
Rat intestine	Proximal	0.280	1.615
	Distal	0.835	3.202

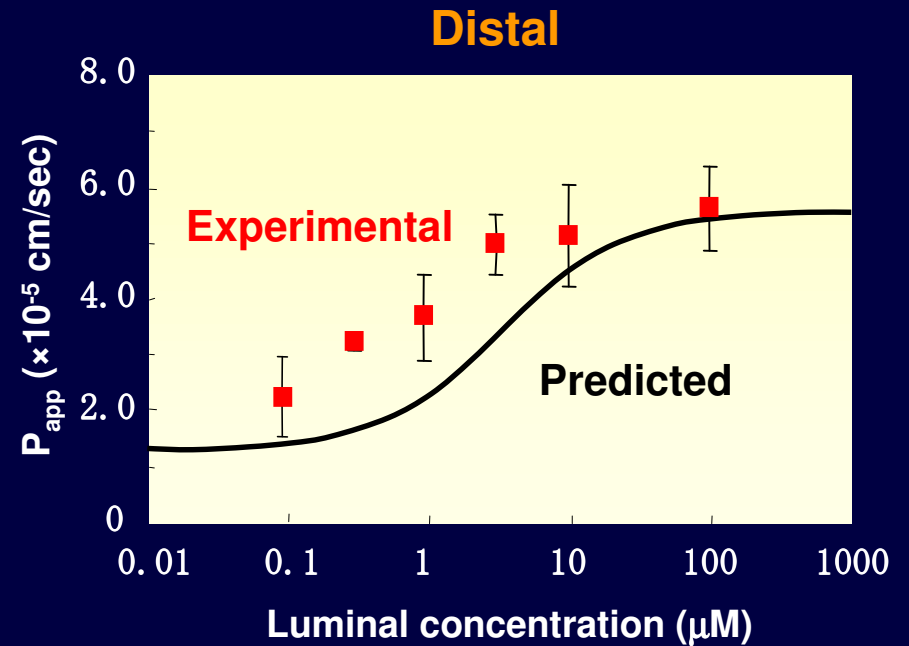
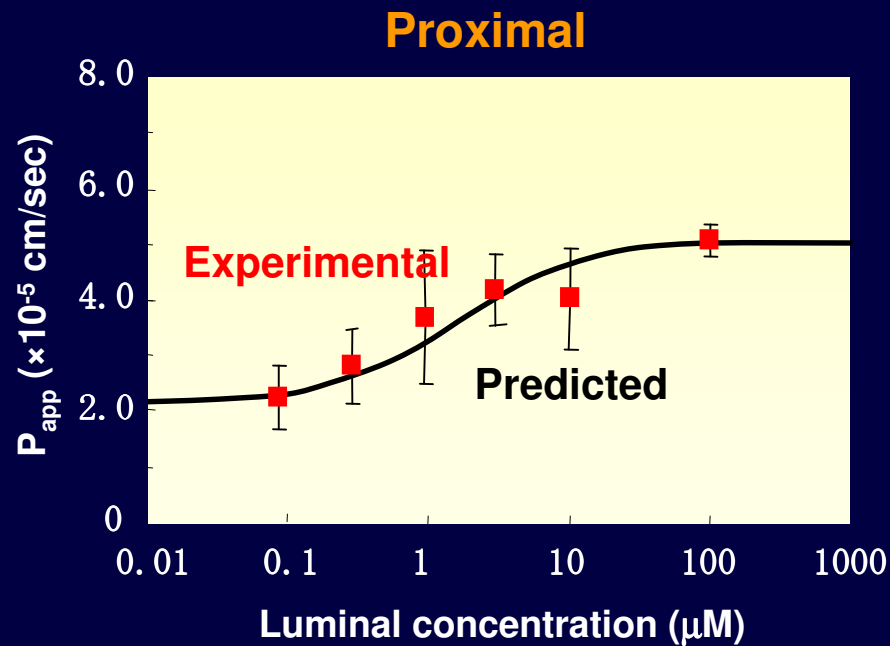
Passive permeability " P_{app} " of **quinidine** in rat small intestine (proximal and distal)

		Quinidine	
		Passive P_{app} ($\times 10^{-5}$ cm/sec)	Ratio (Rat/Caco-2)
Cell monolayers		1.41	—
Rat intestine	Proximal	5.06	3.59
	Distal	5.60	3.97



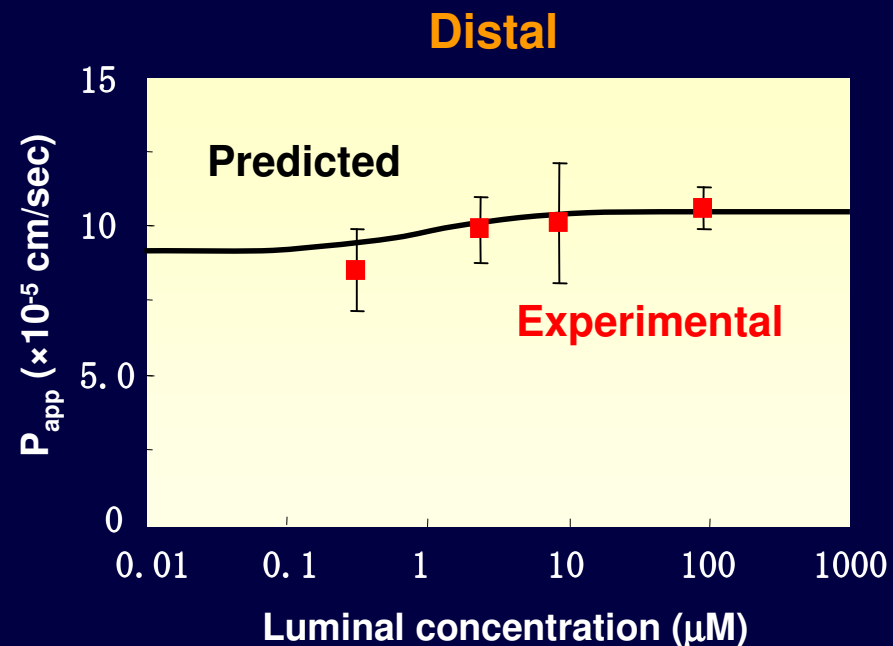
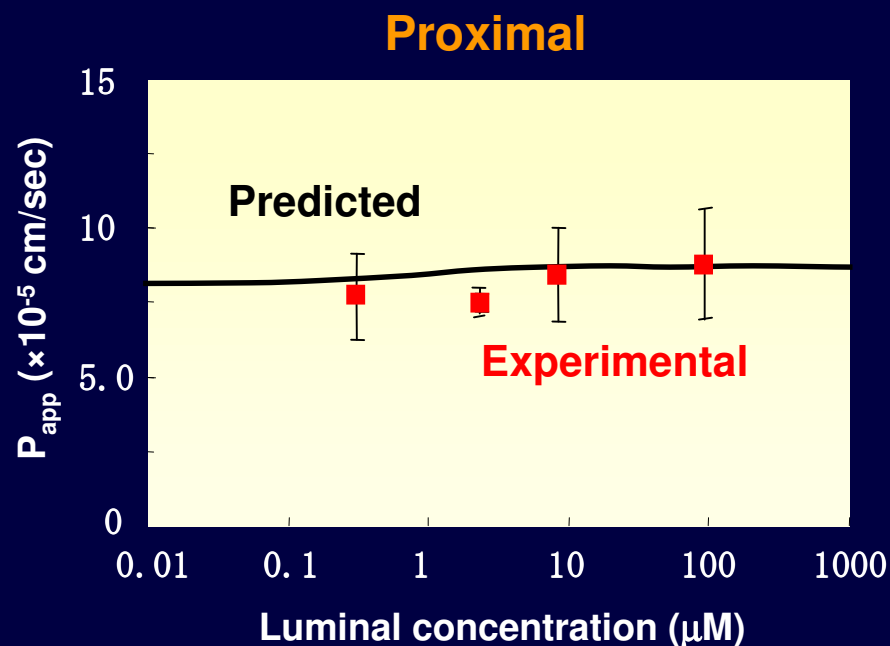
P_{app} of **quinidine** to rat small intestine was measured by *in situ* single-pass perfusion experiments.

Simulation of concentration-dependent permeability of quinidine in rat small intestine (proximal and distal)



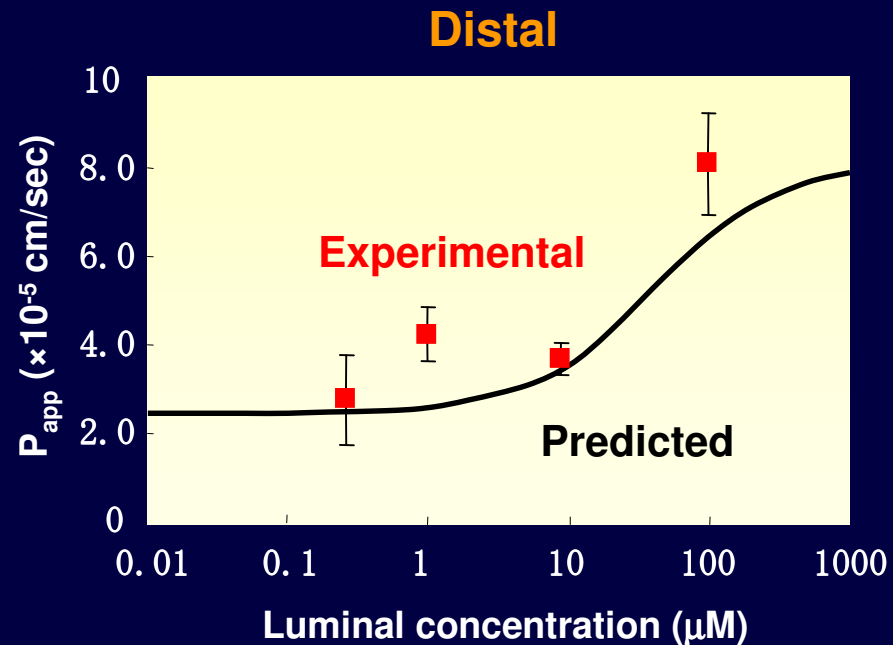
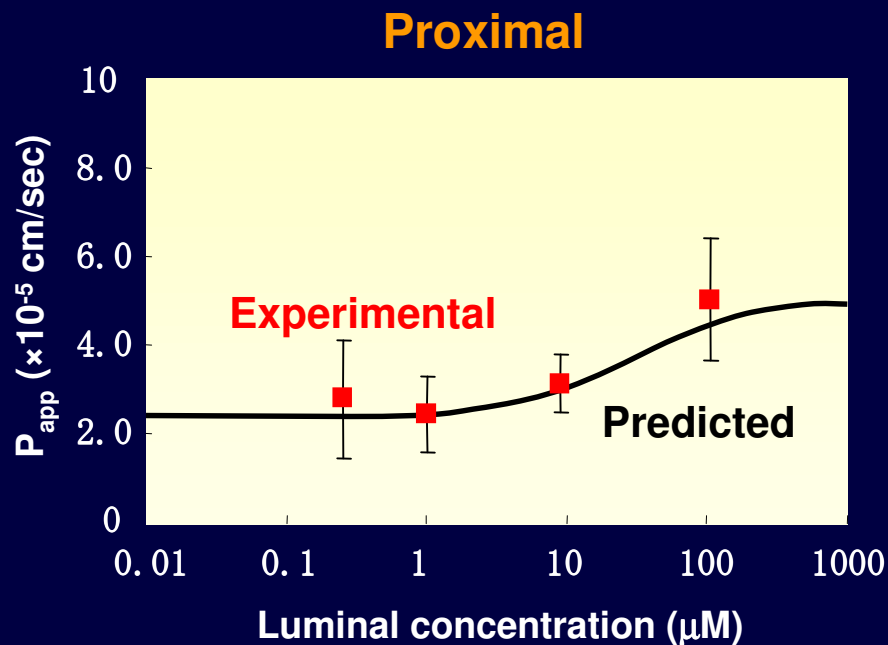
	Proximal			Distal		
	V_{max} ($\times 10^{-5}$ $\mu\text{mol}/\text{min}/10\text{cm gut}$)	$K_{m(\text{app})}$ (μM)	$V_{max}/K_{m(\text{app})}$ ($\times 10^{-5}$ L/min /10cm gut)	V_{max} ($\times 10^{-5}$ $\mu\text{mol}/\text{min}/10\text{cm gut}$)	$K_{m(\text{app})}$ (μM)	$V_{max}/K_{m(\text{app})}$ ($\times 10^{-5}$ L/min /10cm gut)
Predicted	3.13	1.61	1.94	9.35	3.20	2.92
Experimental	1.08	0.60	1.80	1.70	0.70	2.42

Simulation of concentration-dependent permeability of verapamil in rat small intestine (proximal and distal)



	Proximal			Distal		
	V_{max} ($\times 10^{-5}$ $\mu\text{mol}/\text{min}/10\text{cm gut}$)	$K_{m(\text{app})}$ (μM)	$V_{max}/K_{m(\text{app})}$ ($\times 10^{-5}$ $\text{L}/\text{min}/10\text{cm gut}$)	V_{max} ($\times 10^{-5}$ $\mu\text{mol}/\text{min}/10\text{cm gut}$)	$K_{m(\text{app})}$ (μM)	$V_{max}/K_{m(\text{app})}$ ($\times 10^{-5}$ $\text{L}/\text{min}/10\text{cm gut}$)
Predicted	0.33	0.92	0.36	0.98	1.02	0.95
Experimental	0.69	0.87	0.80	1.50	0.82	1.83

Simulation of concentration-dependent permeability of vinblastine in rat small intestine (proximal and distal)

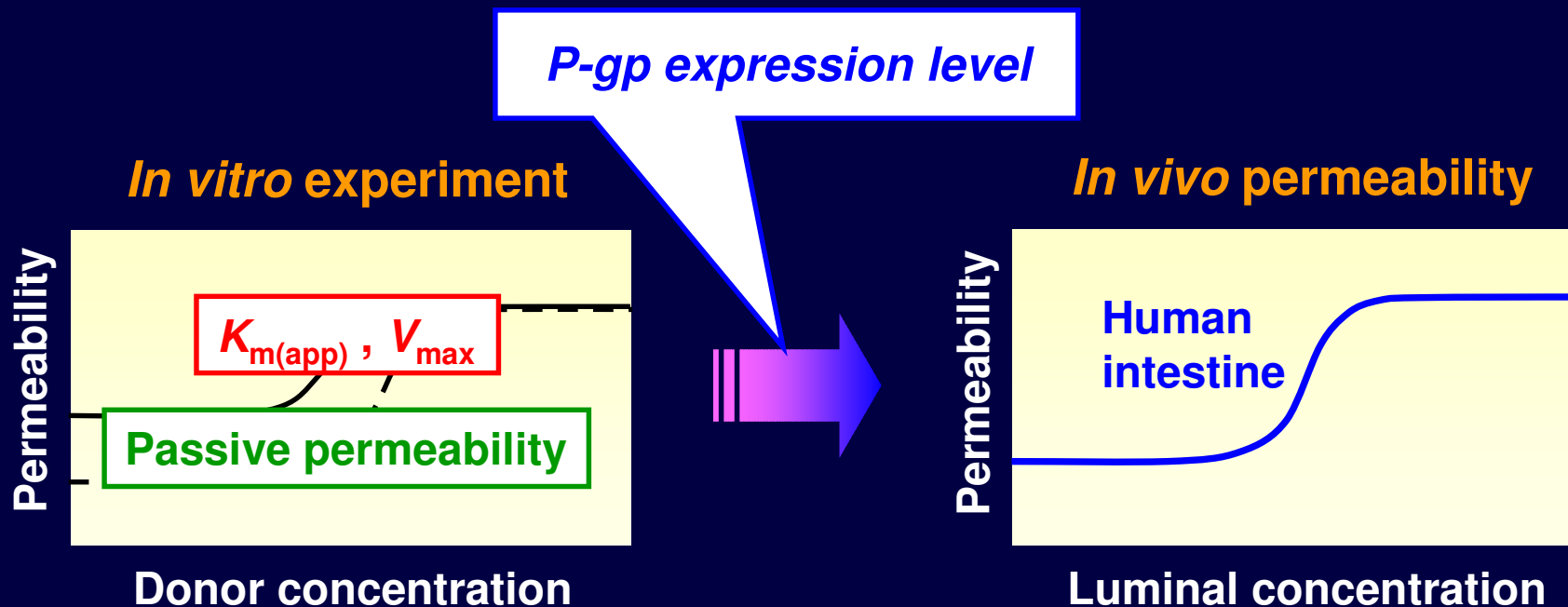


	Proximal			Distal		
	V_{max} ($\times 10^{-5}$ $\mu\text{mol}/\text{min}/10\text{cm}$ gut)	$K_{m(\text{app})}$ (μM)	$V_{max}/K_{m(\text{app})}$ ($\times 10^{-5}$ L/min /10cm gut)	V_{max} ($\times 10^{-5}$ $\mu\text{mol}/\text{min}/10\text{cm}$ gut)	$K_{m(\text{app})}$ (μM)	$V_{max}/K_{m(\text{app})}$ ($\times 10^{-5}$ L/min /10cm gut)
Predicted	51.3	29.3	1.75	153	40.9	3.73
Experimental	38.6	23.1	1.67	111	32.9	3.37

CONCLUSIONS

“Sigmoid E_{\max} model” for permeability-concentration curve of P-gp substrate drugs can provide the fundamental parameters of P-gp-mediated transport, $K_{m(\text{app})}$ and V_{\max} .

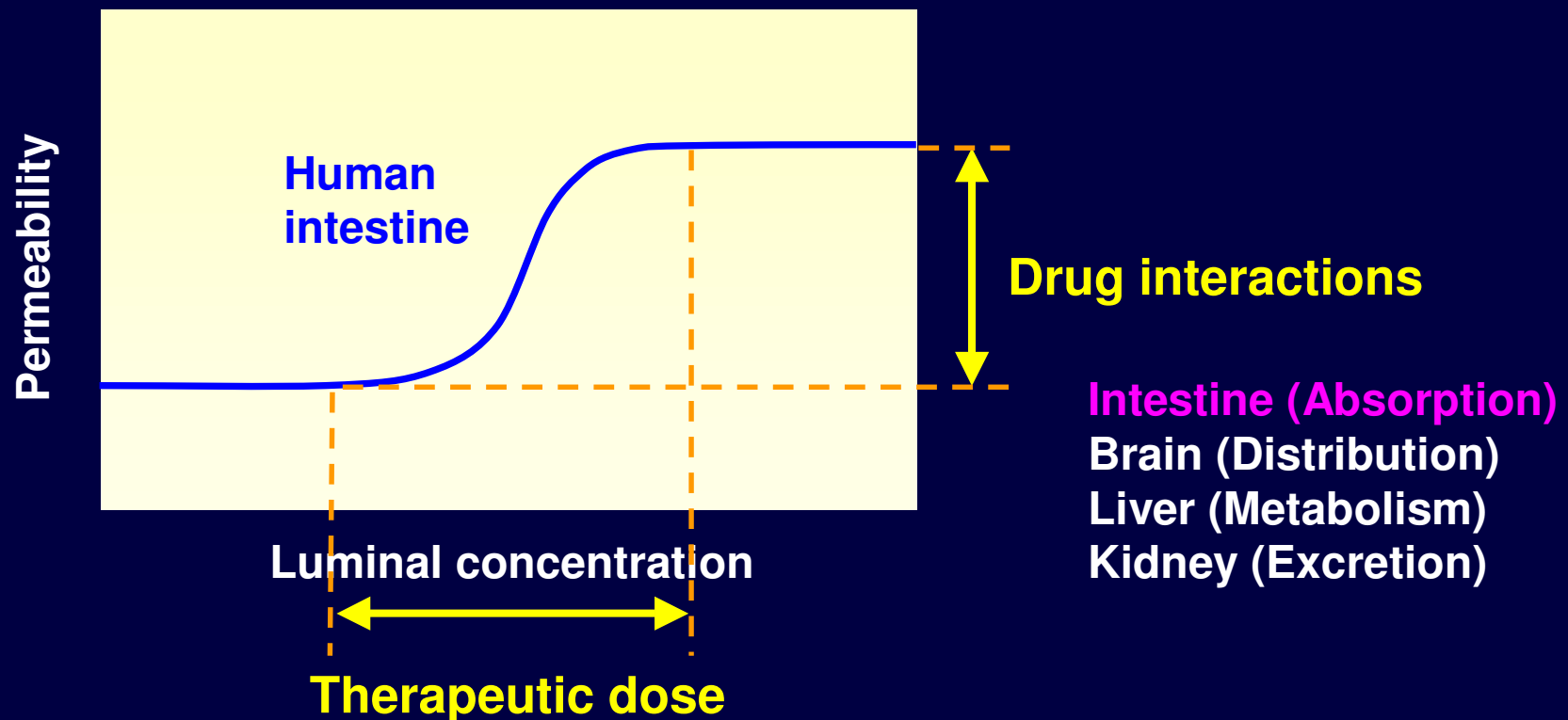
This study clearly demonstrated the possibility to estimate $K_{m(\text{app})}$ and V_{\max} values in human intestine from P-gp expression level and to predict concentration-dependent permeability of P-gp substrate drugs on human intestinal absorption.



SIGNIFICANCE

Permeability-concentration curve of P-gp substrate drugs can predict how P-gp affect the pharmacokinetic profiles of its drugs and how much permeability of its drugs will be changed by dose adjustment and drug-drug interactions in clinical treatment.

In vivo permeability



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