



비임상 DMPK 자료의 생성

신 초 룡

비임상 DMPK 자료의 생성

2022. 11. 30.

개발전략팀
신 초 룡

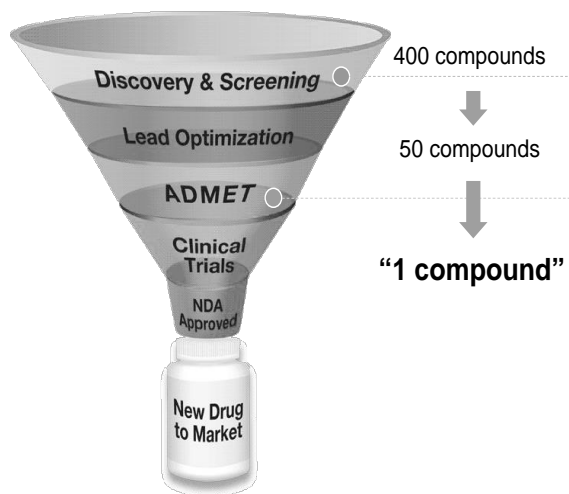
AIMS BioScience
where development meets strategy

AIMS
BioScience

INTRODUCTION

Drug Discovery and Development

AIMS BioScience
where development meets strategy



Discovery stage


- Screening to find druggable compounds



Development stage

- Evaluation of selected compound






Non-clinical DMPK study list

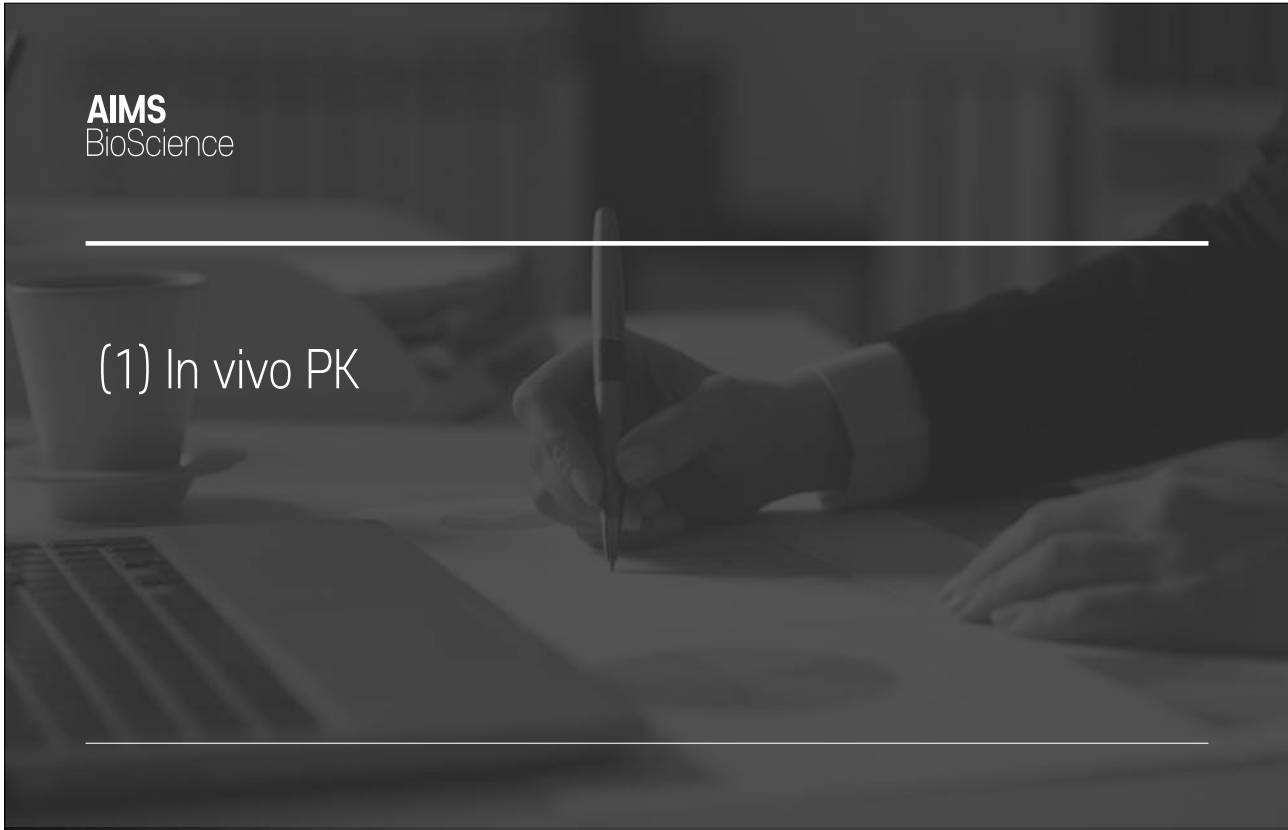
<p>Absorption</p> <ul style="list-style-type: none"> Cell Permeability In vivo PK (C_{max}/T_{max}) Dose proportionality Food effect Gender difference 	<p>Distribution</p> <ul style="list-style-type: none"> Plasma protein binding Microsomal binding Tissue distribution QWBA In vivo PK ($V_{d,ss}$) 	<p>Metabolism</p> <ul style="list-style-type: none"> Plasma stability Metabolic stability In vivo PK (CL) Metabolite ID Reaction phenotyping GSH trapping 	<p>Excretion</p> <ul style="list-style-type: none"> Mass balance Urinary/Fecal excretion Biliary excretion In vivo PK (CL) 	<p>Drug-Drug Interactions</p> <ul style="list-style-type: none"> CYP Direct inhibition CYP Time-dependent inhibition CYP induction Transporter substrate Transporter inhibition
-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------	------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------	------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------	-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------	-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------

4



NON-CLINICAL DMPK STUDIES

- In vivo PK
- Absorption
- Distribution
- Metabolism
- Excretion
- Drug-Drug interaction



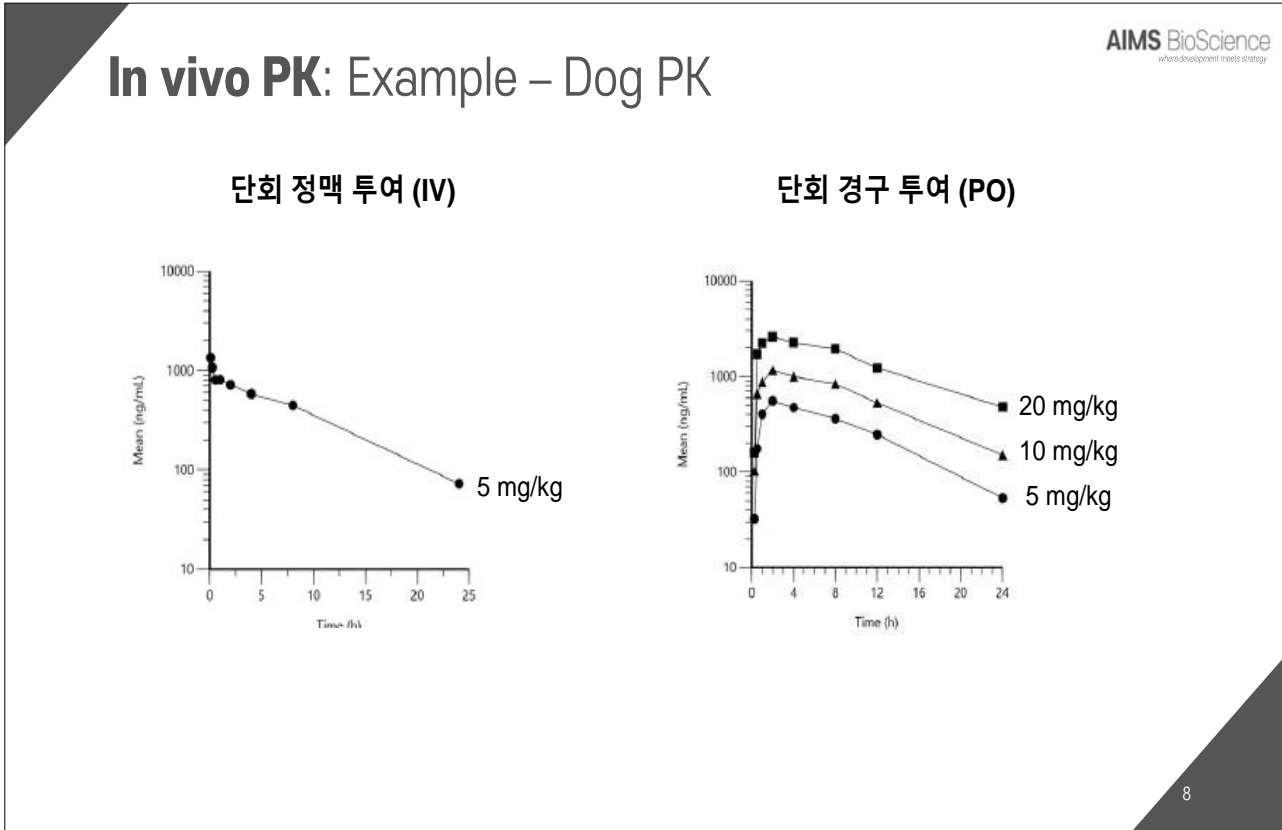
In vivo PK

AIMS BioScience
where development meets strategy

- **시험 목적:** 각 동물 종별 *in vivo* 약물동태 특성 및 dose proportionality 확인
 - Gender difference: male vs. female animals
 - Food effect: fasted vs. fed animals
- **시험 디자인**
 - Species: 3 species 이상에서 평가 권장; Mouse, Rat, Dog (or Monkey)
 - Sampling time points: 7~8 points
 - Group (n=3~6/ group)

Group	Administration Route	Dose (mg/kg)
G1	IV	5 (Low)
G2	PO	5 (Low)
G3		10 (Mid)
G4		20 (High)

7



AIMS BioScience
where development meets strategy

In vivo PK: Example – Dog PK

투여 경로		IV		PO	
Dose (mg/kg)		5	5	10	20
Parameter	Unit				
C_{max}	ng/mL	-	555	1160	2610
C_{min}	ng/mL	1520	-	-	-
$t_{1/2}$	h	-	2	2	2
AUC_{last}	ng·h/mL	8330	6030	13600	32500
$t_{1/2}$	h	6.67	5.52	6.53	7.64
CL or CL/F	mL/h/kg	558	817	669	598
$V_{d,ss}$	mL/kg	5230	-	-	-
F	%	-	72	82	98

[Criteria]

- < 0.3 X 간 혈류량: Low
- 0.3 - 0.7 X 간 혈류량: Moderate
- > 0.7 X 간 혈류량: High

[Criteria]

- 총 체수분량과 비교

- CL: 558 mL/h/kg < Dog's hepatic blood flow (1800 mL/hr/kg)의 30% (= 600 mL/h/kg) → low
- $V_{d,ss}$: 5230 mL/kg > Dog's total body water vol. (1 L/kg) → large, 혈액보다 조직에 더 많이 분포

9

In vivo PK: Example – Dog PK

투여 경로		IV		PO	
Dose (mg/kg)		5	5	10	20
Parameter	Unit				
C_{max}	ng/mL	-	555	1160	2610
C_0	ng/mL	1520	-	-	-
T_{max}	h	-	2	2	2
AUC_{last}	ng·h/mL	8330	6030	13600	32500
$t_{1/2}$	h	6.67	5.52	6.53	7.64
CL or CL/F	mL/h/kg	558	817	669	598
$V_{d,ss}$	mL/kg	5230	-	-	-
F	%	-	72	82	98

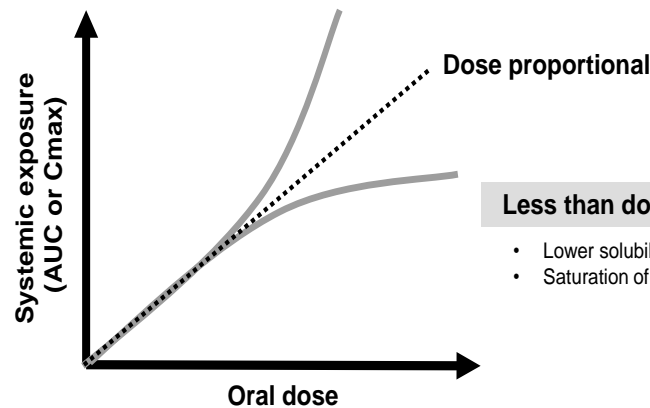
- Systemic exposure: as C_{max} & AUC_{last}
- Dose ratio = 1 : 2 : 4 → C_{max} ratio = 1 : 2.1 : 4.7 / AUC_{last} ratio = 1 : 2.3 : 5.4
 - Systemic exposure was increased dose proportionally at the dose range of 5~20 mg/kg following oral administration.

10

In vivo PK: Dose proportionality

More than dose proportional

- Substrate of efflux transporter
- Saturation of metabolism



Less than dose proportional

- Lower solubility (dissolution process)
- Saturation of uptake transport

11

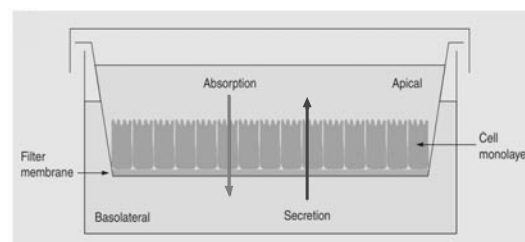
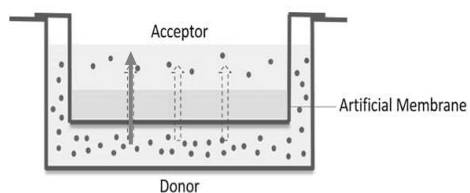
AIMS
BioScience

(2) Absorption

Absorption: Cell permeability

AIMS BioScience
where development meets strategy

- **시험 목적:** 약물의 세포 투과도 평가 (*in vitro*)
 - PAMPA (artificial membrane)
 - Caco-2 cells, MDCK cells...
- **시험 디자인**
 - Test conc. : 1 conc.
 - Caco-2 or MDCK cell assay: efflux ratio 산출 (efflux transporter에 대한 기질성 확인 가능)



1) The Role of Intestinal Permeability in Gastrointestinal Disorders and Current Methods of Evaluation, Tim et al., Sec. Nutritional Immunology.
2) <https://www.orsolyaszahmat.com/post/increased-intestinal-permeability-what-it-is-and-why-you-should-care-about-it>

AIMS BioScience
where development meets strategy

Absorption: Cell permeability

- Caco-2 permeability 시험 결과 (Example)**

[Criteria] ($\times 10^{-6}$ cm/sec)

- High: >10
- Moderate: 1 - 10
- Low: <1

[Criteria]

- Efflux Ratio > 2

Compound	Test conc. (μ M)	Mean P_{app} ($\times 10^{-6}$ cm/sec)		Efflux ratio
		A to B	B to A	
AIMS-001	1	7.44	59.9	8.06

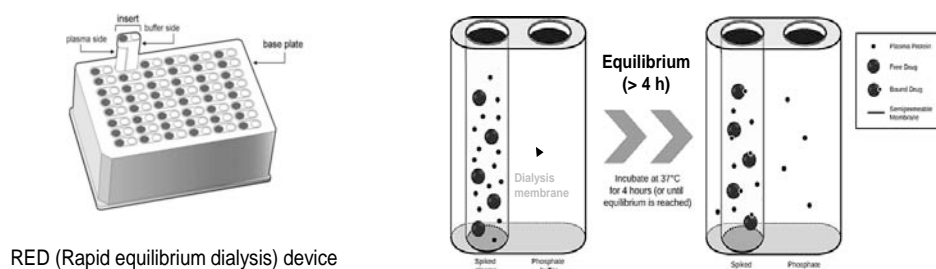
- $P_{app, A \text{ to } B}$: 7.44 ($\times 10^{-6}$ cm/sec) \rightarrow moderate
- Efflux ratio (= $P_{app, B \text{ to } A} / P_{app, A \text{ to } B}$) : 8.06 \rightarrow possible to be a substrate of efflux transporters

14



Distribution: Plasma protein binding

- **시험 목적:** 동물 종의 혈장 내 단백질 결합률 확인 (*in vitro*)
 - Equilibrium dialysis, Ultrafiltration, Ultracentrifugation
- **시험 디자인**
 - Test species: Human + Animals (약효 및 독성 동물 종 포함)
 - Test conc. : 1 conc.



RED (Rapid equilibrium dialysis) device

16

Distribution: Plasma protein binding

- **시험 결과 (Example)**

	Mouse	Rat	Dog	Monkey	Human
fraction bound (% Bound)	0.993 (99.3%)	0.996 (99.6%)	0.993 (99.3%)	0.995 (99.5%)	0.996 (99.6%)
fraction unbound (% Unbound)	0.007 (0.7%)	0.004 (0.4%)	0.007 (0.7%)	0.005 (0.5%)	0.004 (0.4%)

- Fraction unbound ($f_{u,p}$) = 1 - Fraction bound
- Highly bound in the plasma protein of all species

17

Distribution: QWBA

- 시험 목적: 약물의 조직 분포 확인 (*in vivo*) – target organ, eye uveal or pigmented skin
- 시험 디자인
 - Test article: Radio-labeled compound
 - Test species: Rat
 - Dose: 1 dose
 - Animals will be sacrificed at each sampling time points

Group	Species	Sampling time points
G1	SD rats (Albino)	> 5 (up to 96 h)
G2	Long-Evans rats (Pigmented)	> 5 (up to 168 h)

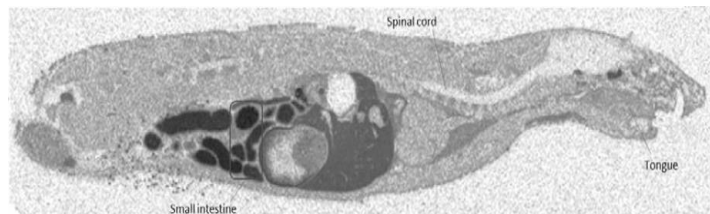


18

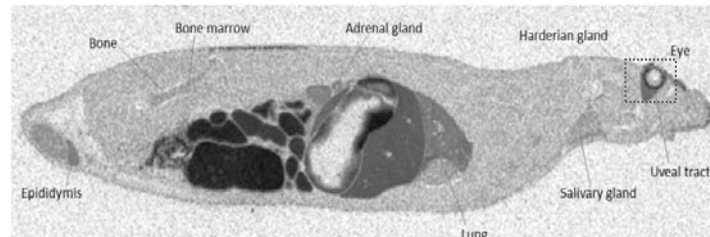
Distribution: QWBA

- 시험 결과 (Example)

G1: SD rats (Albino)



G2: LE rats (Pigmented)



19

Distribution: QWBA

- 시험 결과 (Example)

G2: LE rats (Pigmented)

In the pigmented rats, highly bind to eye uveal and skin and remain longer → potential for melanin binding

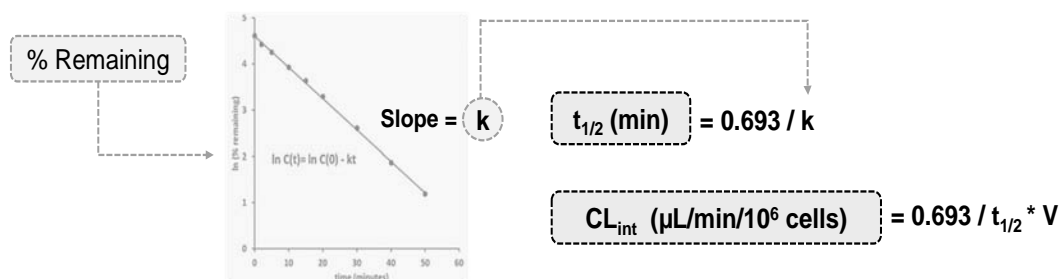
Tissue/Organ	C _{max} (ng-eq/g)	t _{max} (h)	t _{1/2} (h)	AUC _{0-t} (ng-eq ^h /g)	PI	Tissue/Organ	C _{max} (ng-eq/g)	t _{max} (h)	t _{1/2} (h)	AUC _{0-t} (ng-eq ^h /g)	Tissue: Plasma*
Plasma(LSC)	2330	4.0	167	72,557		Lymph gland(neck)	898	4.0	NC	5,917	0.08
Aorta	929	4.0	NC	5,302		Spleen	1433	4.0	NC	9,224	0.13
Blood(heart)	1740	4.0	NC	8,093		Thymus	806	4.0	NC	4,746	0.07
Whole brain	NA	NA	NA	NA	NA	Kidney	3033	4.0	NC	14,985	0.21
Spinal cord	NA	NA	NA	NA	NA	Renal cortex	2657	4.0	NC	27,606	0.38
Eye	3048	8.0	229	241,907	3.33	Renal Medulla	3480	4.0	NC	17,781	0.25
Crystalline lens	NA	NA	NA	NA	NA	Liver	3903	4.0	NC	20,337	0.28
Uvea	8307	8.0	162	1,102,109	15.19	Bladder wall	2974	8.0	NC	10,623	0.15
Non-pigmented skin	835	4.0	NC	3,829	0.05	Heart	1211	4.0	NC	6,584	0.09
Pigmented skin	1454	4.0	NC	23,340	0.32	Muscle (thighbone)	837	4.0	NC	5,743	0.08
Adrenal gland	1908	4.0	NC	11,492	0.16	Prostate	716	4.0	NC	4,130	0.06
Adrenal cortex	2019	4.0	NC	11,673	0.16	Testes	NA	NA	NA	NA	NA
Adrenal medulla	1823	4.0	NC	10,477	0.14	Lung	1444	4.0	NC	7,970	0.11
Hypophysis cerebri	1815	4.0	NC	9,631	0.13	Turbinal	656	4.0	NC	3,795	0.05
Thyroid	1300	4.0	NC	6,744	0.09	Trachea	882	4.0	NC	4,618	0.06
Brown fat	1029	4.0	NC	11,190	0.15	Lacrimal gland	1270	4.0	NC	8,583	0.12
White fat (abdominal cavity)	NA	NA	NA	NA	NA	Harderian gland	1748	4.0	NC	9,188	0.13
Esophagus	3662	1.0	NC	13,866	0.19	Pancreas	1385	4.0	NC	8,781	0.12
Stomach wall (glandular region)	31505	1.0	NC	63,775	0.88	Salivary gland	1703	4.0	NC	9,637	0.13
Stomach wall(Non-glandular region)	33830	1.0	NC	58,641	0.81	Bone marrow (thighbone)	1420	4.0	NC	8,228	0.11
Small Intestine Wall	11336	1.0	NC	31,040	0.43	Bone (thighbone)	NA	NA	NA	NA	NA
Large intestine wall	17894	4.0	35.3	295,349	4.07						

20

(4) Metabolism

Metabolism: Metabolic stability

- **시험 목적:** 간 대사에 대한 약물의 안정성 평가 (*in vitro*)
 - Liver microsome or S9
 - Hepatocyte
- **시험 디자인**
 - Test species: Human + Animals (Mouse, Rat, Dog, Monkey...)
 - Test conc. : 1 conc.
 - Incubation time: > 60 min



22

Metabolism: Metabolic stability

- **시험 결과 (Example)**

Species	Test conc. (uM)	$t_{1/2}$ (min)	CL_{int} ($\mu\text{L/min}/10^6$ cells)	Category
Mouse	1	59.1	23.5	High (>17.8)
Rat		10.8	129	High (>27.5)
Dog		34.2	40.5	High (>10.5)
Monkey		3.59	386	High (>28.3)
Human		39.7	34.9	High (>19.0)

23

Metabolism: Metabolite Identification

- **시험 목적:** 종 간 대사체 프로파일 비교 및 대사체 구조 규명 (*in vitro*)
 - Liver microsome or S9
 - Hepatocyte
- **시험 디자인**
 - Test species: Human + Animals (Mouse, Rat, Dog, Monkey...)
 - Test conc. : 1 conc.
 - Incubation time: > 60 min

24

Metabolism: Metabolite Identification

- **시험 결과 (Example)**

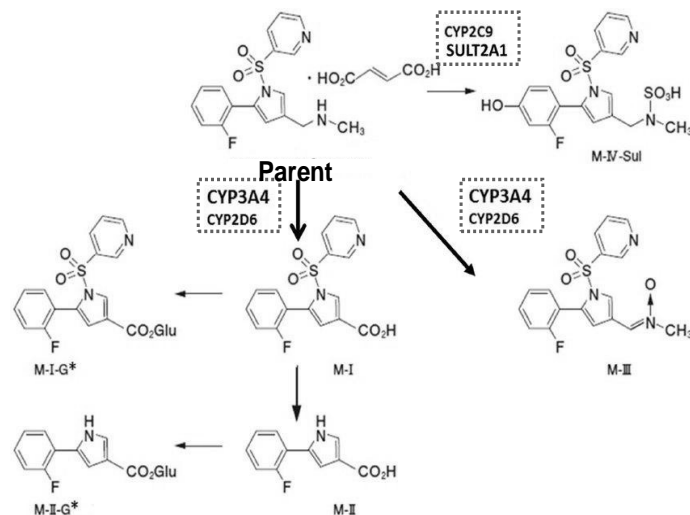
Proposed assignment	Relative Abundance (UV Peak Area %)				
	Human	Mouse	Rat	Dog	Monkey
Parent	80.1	89.1	42.9	89.7	79.3
M1 Oxidation+Desaturation	-	-	1.73	-	-
M2 Oxidation	1.14	7.71	5.29	2.81	3.34
M3 Oxidation	5.79	-	37.8	-	-
M4 Oxidation	2.92	-	2.38	1.37	-
M5 2x Oxidation	-	-	2.71	-	-
M6 Oxidation + Sulfation	-	-	1.65	2.17	-
M7 Oxidation+sulfation	-	-	1.09	-	-
M8 Glucuronidation	2.55	2.16	3.52	3.94	7.05
M9 Oxidation + Glucuronidation	1.17	-	2.87	-	-

- No human unique metabolites
- 사람과 가장 유사한 대사체 프로파일을 보인 동물 종: Rat, Dog

25

Metabolism: Metabolite Identification

Proposed metabolic pathway




26

Metabolism: Reaction phenotyping

- **시험 목적:** 약물의 human CYP (and/or UGT) enzymes에 대한 기질성 확인 (*in vitro*)
 - Human recombinant CYPs/UGTs
 - Human liver microsomes (using chemical inhibitors)
- **시험 디자인**
 - Test Conc: 1 conc.
 - Sampling time points: 4~5 time points

Study item	CYP phenotyping		UGT phenotyping
Isoforms	CYP1A2, 2B6, 2C8, 2C9, 2C19, 2D6, 3A4		UGT1A1, 1A3, 1A4, 1A6, 1A9, 2B7, 2B15
Test system	Individual rCYPs	Liver microsomes (using chemical inhibitors)	Individual rUGTs
Deliverables	% remaining, $t_{1/2}$	Activity remaining (as % of no inhibitor)	% remaining, $t_{1/2}$

27



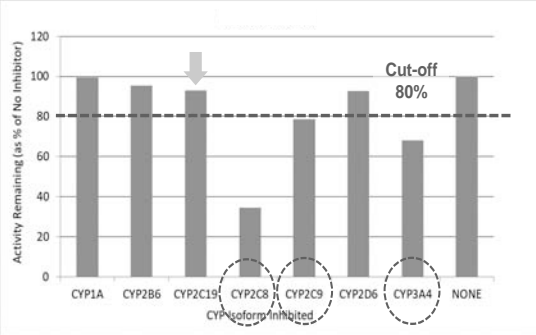
Metabolism: Reaction phenotyping

- CYP phenotyping 시험 결과 (Example)**

1) Recombinant CYPs


Isoforms	% remaining					t1/2 (min)
	0 min	5 min	15 min	30 min	45 min	
CYP1A2	100	80.1	88.8	84.0	79.0	213
CYP2B6	100	93.9	84.5	92.1	87.1	324
CYP2C8	100	0.15	0.15	0.14	0.13	0.54
CYP2C9	100	60.0	25.3	6.72	3.30	7.77
CYP2C19	100	17.7	0.06	0.06	0.06	2.00
CYP2D6	100	91.4	105	99.5	112	N/A
CYP3A4	100	71.0	48.7	29.9	24.3	17.9

2) Using chemical inhibitors

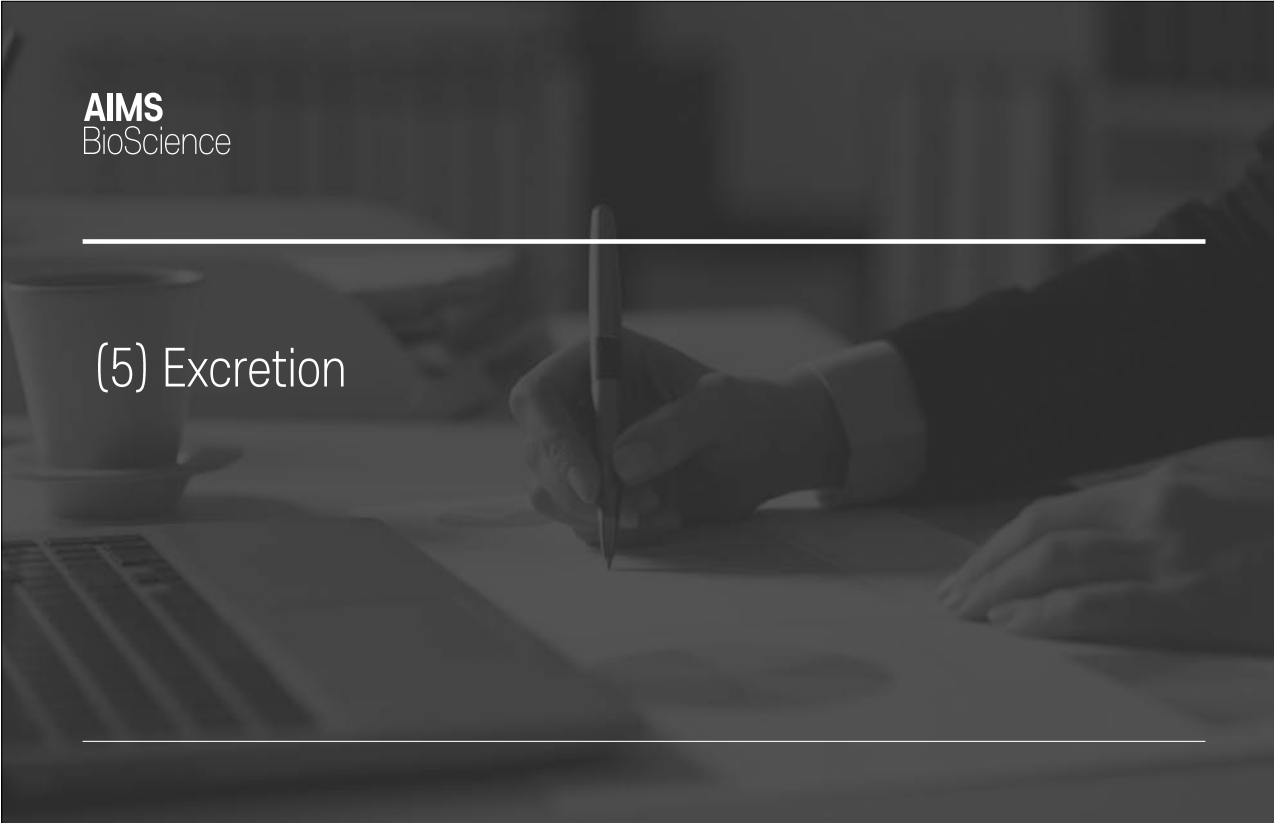


- Can be metabolized by CYP2C8, CYP2C9 and CYP3A4 (as a substrate)

28



(5) Excretion



Excretion: Mass balance

- **시험 목적:** 약물의 주된 배설 경로 규명 (*in vivo*)
 - Excreta (Urine, Feces, Bile) 내 metabolites profiling study 함께 수행할 수 있음.
- **시험 디자인**
 - Test article: Radio-labeled compound
 - Test species: Rat
 - Dose: 1 dose
 - Sampling time points: 7~8 time points

Group		Sampling time points	Collection matrices
G1	Intact rats	up to 96~120 h	Urine, feces (+ Cage rinse, Carcass)
G2	Bile-duct cannulated rats	up to 72 h	Urine, feces, <u>bile</u> (+ Cage rinse, Carcass)

30

Excretion: Mass balance

- **시험 결과 (Example)**

1) Intact Rat

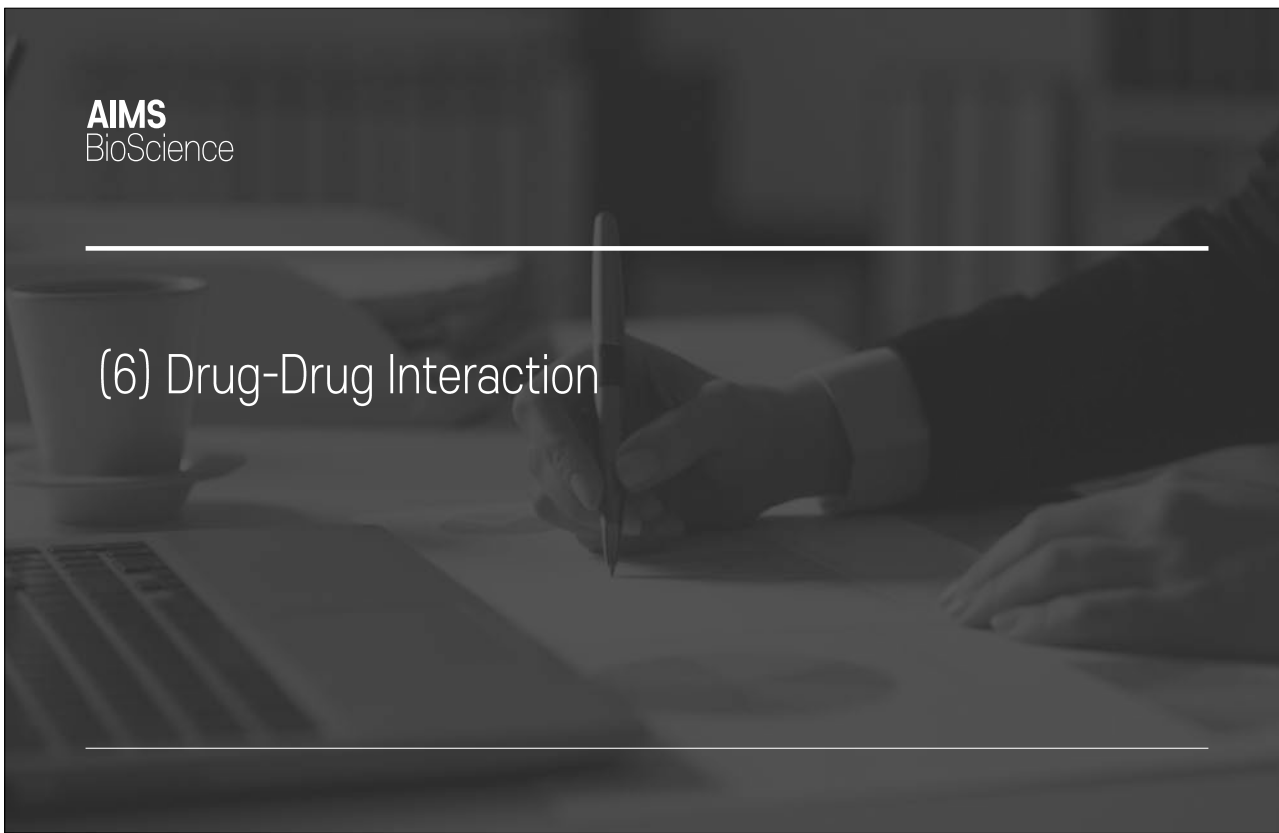
Matrix	Cumulative % of administered dose
Urine	17.3
Feces	82.2
Cage rinse	2.2
Carcass	0.6
Total	102.3

2) BDC Rat

Matrix	Cumulative % of administered dose
Bile	70.0
Urine	19.4
Feces	9.9
Cage rinse	0.2
Carcass	0.6
Total	100.2

- Main route of excretion: Fecal excretion via biliary excretion

31



AIMS
BioScience

(6) Drug-Drug Interaction

AIMS BioScience
where development meets strategy



DDI: *in vitro* Drug Interaction Studies (FDA)

Classification	Metabolism-mediated	Transporters-mediated
Substrate	<ul style="list-style-type: none"> • CYPs: 1A2, 2B6, 2C8, 2C9, 2C19, 2D6, 3A • Use both methods; <ul style="list-style-type: none"> • Chemical inhibitors in pooled HLMs • Human recombinant enzymes 	<ul style="list-style-type: none"> • ABC transporters: P-gp, BCRP • SLC transporters: OATP1B1/OATP1B3, OAT1/OAT3, MATE1/MATE2-K, OCT2
Inhibition	<ul style="list-style-type: none"> • CYPs: 1A2, 2B6, 2C8, 2C9, 2C19, 2D6, 3A (midazolam, testosterone) • Evaluate in both manners; <ul style="list-style-type: none"> • Reversible manner • Time-dependent manner 	<ul style="list-style-type: none"> • ABC transporters: P-gp, BCRP • SLC transporters: OATP1B1/OATP1B3, OAT1/OAT3, MATE1/MATE2-K, OCT2
Induction	<ul style="list-style-type: none"> • CYPs: 1A2, 2B6, 3A4 first • If induce 3A4, should evaluate 2C 	<i>Recommendations are not provided</i>
Note	<ul style="list-style-type: none"> • Preferably before first-in-human studies 	<ul style="list-style-type: none"> • Before clinical studies in patients

33

DDI: CYP inhibition

- **시험 목적:** 약물의 human CYP enzymes에 대한 저해능 확인 (*in vitro*)
 - CYP1A2, 2B6, 2C8, 2C9, 2C19, 2D6, 3A (Midazolam/Testosterone)
 - Reversible inhibition & Time-dependent inhibition
- **시험 디자인**
 - Test system: human liver microsome
 - Test conc: > 6 conc. (to estimate IC₅₀); high enough

Reversible inhibition	Time-dependent inhibition
 + CYP substrates + Drugs (Inhibitor) ↓ Incubation for 15 min	 + Drugs (Inhibitor) / Metabolites ↓ Pre-incubation for 30 min + CYP substrates ↓ Incubation for 15 min

34

DDI: CYP inhibition

- **Reversible inhibition 시험 결과 (Example)**

IC ₅₀ (μM)						
CYP1A2	CYP2B6	CYP2C8	CYP2C9	CYP2C19	CYP2D6	CYP3A4
>50	>50	0.482	28.9	7.87	>50	>50

- Strong inhibition potential on CYP2C8 (IC₅₀ < 1 μM)

35

AIMS BioScience
where development meets strategy

DDI: CYP inhibition

- Time-dependent inhibition 시험 결과 (Example)

Isoforms	IC ₅₀ (µM)		Fold shift
	0 min pre-incub.	30 min pre-incub. (-NADPH) (+NADPH)	
CYP1A2	>50.0	>50.0 >50.0	ND
CYP2B6	>50.0	28.3 37.5	0.76
CYP2C8	<0.200	0.25 0.25	0.99
CYP2C9	17.1	18.8 24.0	0.78
CYP2C19	5.71	7.06 6.74	1.05
CYP2D6	>50.0	>50.0 >50.0	ND
CYP3A4 (MDZ)	>50.0	>50.0 23.0	>2.17
CYP3A4 (TST)	>50.0	>50.0 32.3	>1.55

[Criteria]

- Fold shift ≥ 1.5

- Time-dependent inhibitor of CYP3A4

36

AIMS BioScience
where development meets strategy

DDI: CYP induction

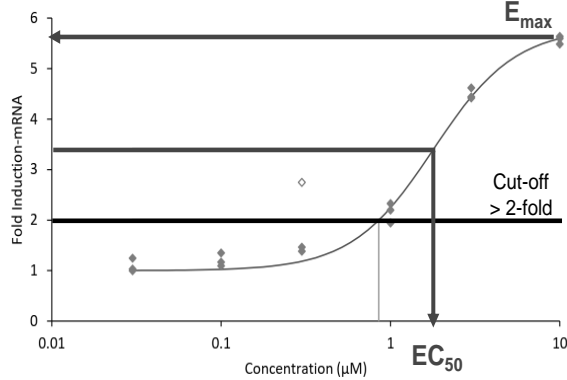
- 시험 목적: 약물의 human CYP enzymes에 대한 유도능 확인 (*in vitro*)
 - CYP1A2, 2B6, 3A4 first
 - If 3A4 inducer → should evaluate induction potential on 2C family
- 시험 디자인
 - Test system: human hepatocytes (from at least 3 donors)
 - Test conc: > 6 conc.
 - Incubation time: 48~72 h
 - Study endpoints: CYPs mRNA levels (and/or enzyme activity)

37

DDI: CYP induction

• 시험 결과 (Example)

Effect on mRNA expression of Cryopreserved Human Hepatocytes cells (CYP3A4)



[Criteria]

- ≥ 2 -fold increase

CYP1A2	mRNA expression		
	donor 1	donor 2	donor 3
Maximum Fold induction	NI	NI	1.53
EC ₅₀ (µM)	NI	NI	NC
E _{max} (µM)	NI	NI	NC
CYP2B6	mRNA expression		
	donor 1	donor 2	donor 3
Maximum Fold induction	NI	5.57	1.93
EC ₅₀ (µM)	NI	1.80	NC
E _{max} (µM)	NI	5.81	NC
CYP3A4	mRNA expression		
	donor 1	donor 2	donor 3
Maximum Fold induction	NI	1.43	NI
EC ₅₀ (µM)	NI	NC	NI
E _{max} (µM)	NI	NC	NI

- Induction potential on CYP2B6

38

DDI: CYP inhibition/induction

• Data interpretation: Basic kinetic model

Category	Reversible inhibition	Time-dependent inhibition	Induction
R-value equation	$R_1 = 1 + \frac{I_{max,u}}{K_{i,u}}$	$R_2 = \frac{K_{obs} + K_{deg}}{K_{deg}}$ $K_{obs} = \frac{K_{inact} \times 50 \times I_{max,u}}{K_{I,u} + 50 \times I_{max,u}}$	$R_3 = \frac{1}{1 + d \cdot \left(\frac{E_{max} \times 10 \times I_{max,u}}{EC_{50} + 10 \times I_{max,u}} \right)}$
Cut-off	$R_1 \geq 1.02$	$R_2 \geq 1.25$	$R_3 \leq 0.8$
Note	I _{max,u} = the maximal unbound plasma concentration of the interacting drug at steady-state (I _{max,u} = C _{max,ss} * f _{u,p}) K _{i,u} = the unbound inhibition constant determined in vitro (K _{i,u} = K _i * f _{u,mic}) K _{I,u} = the unbound inhibitor concentration causing half-maximal inactivation (K _{I,u} = K _I * f _{u,mic}) K _{inact} = the maximal inactivation rate constant E _{max} = the maximum induction effect EC ₅₀ = in vitro concentration causing half-maximum induction d = a scaling factor (assumed to be 1 for the basic model)		

- If $R_1 \geq 1.02$, $R_2 \geq 1.25$ or $R_3 \leq 0.8$ → Mechanistic models or Clinical DDI studies needed.

39

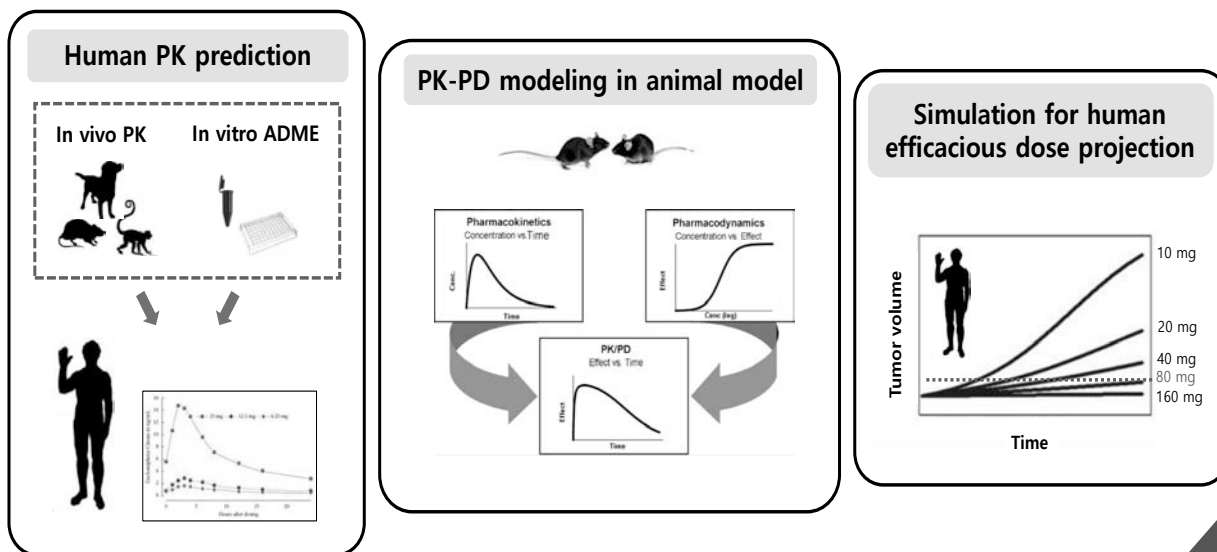
AIMS
BioScience

APPLICATION OF NON-CLINICAL DMPK DATA

Application of Non-clinical DMPK data

AIMS BioScience
where development meets strategy

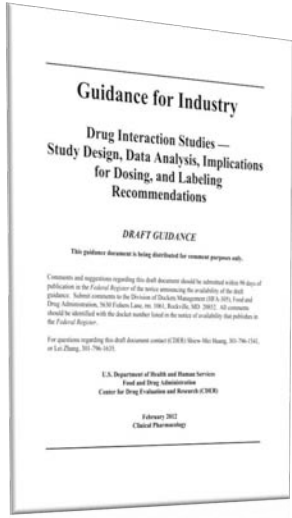
I. Human PK prediction



41

Application of Non-clinical DMPK data

II. Risk assessment for DDI



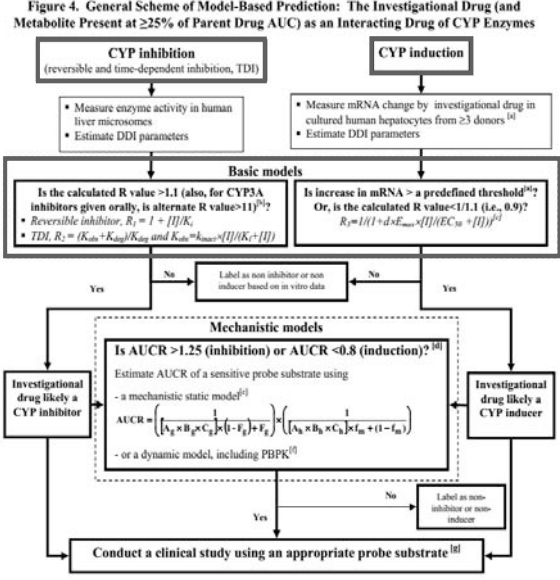


Figure 4. General Scheme of Model-Based Prediction: The Investigational Drug (and Metabolite Present at ≥25% of Parent Drug AUC) as an Interacting Drug of CYP Enzymes

CYP inhibition (reversible and time-dependent inhibition, TDI)

- Measure enzyme activity in human liver microsomes
- Estimate DDI parameters

CYP induction

- Measure mRNA change by investigational drug in cultured human hepatocytes from ≥3 donors^[4]
- Estimate DDI parameters

Basic models

Is the calculated R value >1.1 (also, for CYP3A inhibitors given orally, is alternate R value >11)^[5]?

- Reversible inhibitor, $R_1 = 1 + [I]/K_i$
- TDI, $R_2 = (K_{m,app} + K_{m,app} [I]) / (K_m + [I])$

Is increase in mRNA > a predefined threshold^[6]?

Or, is the calculated R value <1/1.1 (i.e., 0.9)?

$R_1 = 1 / (1 + d \times E_{max} \times [I] / (EC_{50} + [I]))^{Hill}$

Yes/No paths lead to: Label as non inhibitor or non inducer based on in vitro data.

Mechanistic models

Is AUCR >1.25 (inhibition) or AUCR <0.8 (induction)?^[6]

Estimate AUCR of a sensitive probe substrate using:

- a mechanistic static model^[6]
- or a dynamic model, including PBPK^[6]

$AUCR = \left(\frac{[I] \times B_{in} \times C_{p,ss}}{[I] + K_{i,app}} + 1 \right) \left(\frac{[I] \times B_{in} \times C_{p,ss}}{[I] + K_{i,app}} + 1 - f_{in} \right)$

Yes/No paths lead to: Label as non-inhibitor or non-inducer.

Investigational drug likely a CYP inhibitor / **Investigational drug likely a CYP inducer**

Conduct a clinical study using an appropriate probe substrate^[6]

“Clinical DDI studies needed?”

42

경청해 주셔서 감사합니다.

(Question: cherlyn@aimsbiosci.com)