

Clinical Pharmacology

- [FDA](#)

- [EMA](#)

- [NMPA](#)

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- [PMDA](#)

- [Invalid](#)

FDA

EMA

NMPA

NMPA

????????????????????????????????

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,
,

,
,

Maximum Recommended Starting Dose
MRSD

MRSD
MRSD
No Observed Adverse Effect Level
NOAEL
Human Equivalent Dose
HED
,
Minimal Anticipated Biological Effect Level
MABEL)

,
:
)
MRSD

MRSD
MRSD
MRSD

??????

MRSD

MRSD

????????

??????????????MRSD

NOAEL HED HED MRSD
 NOAEL NOAEL HED
 Safety Factor, SF

MRSD NOAEL MRSD
 ,

NOAEL HED HED
 Pharmacologically Active Dose PAD MRSD

??????????????MRSD

/
 PK/PD

??????????????MRSD

????1?????????NOAEL???

MRSD NOAEL NOAEL
 MRSD
 NOAEL

 NOAEL

NOAEL 1
 2
 3

 NOAEL I

2

3 

4

[illegible]

5□□□

[illegible]

NOAEL □ HED

[illegible]

10

[illegible][illegible][illegible][illegible]

1. ??????

[illegible][illegible]

MRSD [] HED [][][][]

10

[illegible][illegible][illegible][illegible]

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[illegible][illegible][illegible]

HED

[illegible][illegible][illegible][illegible][illegible][illegible][illegible][illegible][illegible]

[illegible]

1.

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 (

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)

[illegible]

☐☐☐☐☐☐) ☐☐☐☐☐☐☐☐☐☐ Cmin ☐ Cmax ☐ AUC ☐☐☐

(

2. NOAEL (NOAEL)

3. ☐ NOAEL (Safety Margin)

4.

[illegible][illegible]

Allometric Interspecies Scaling ☒ Detricks ☐☐☐☐☐☐

[]

[illegible]

]

[illegible][illegible][illegible]NOAEL

1/10

????????????????MRSD

[illegible]

MABEL

[illegible][illegible][illegible]PK/PD ,

????

[illegible]

□□□□ NOAEL□□ HED□□□□□□□□ MRSD

[illegible][illegible][illegible]

1

[illegible][illegible][illegible][illegible]

XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX

??????

XX	XX
MRSDX Maximum Recommended Starting DoseX	XXXXXXXXXXXXXXXXXXXXXXXXXXXX XXXXXXXXXXXX MRSD XXXXXXXXXXXXXXXXXXXX mg/kgX mg/m2 XXXXXX
NOAELX No Observed Adverse Effect LevelX	XXXXXXXXXXXXXXXXXXXXXXXXXXXX XXXXXXXXXXXX NOAEL XXXXXXXXXXXXXXXXXXXX
NOELX No Observed Effect LevelX	XXXXXXXXXXXXXXXXXXXXXXXXXXXX
LOAELX Lowest Observed Adverse Effect LevelX	XXXXXXXXXXXXXXXXXXXXXXXXXXXX X
MTDX Maximum Tolerated DoseX	XXXXXXXXXXXXXXXXXXXXXXXXXXXX
PADX Pharmacologically Active DoseX	XXXXXXXXXXXXXXXXXXXXXXXXXXXX X
HEDX Human Equivalent DoseX	XXXXXXXXXXXXXXXXXXXXXXXXXXXX XXXXXX HEDXXXX NOAEL XXXXXXXXXXXXXXXXXXXX PADXXXX NOAELXXXXXX
BSA-CFX Body Surface Area Conversion FactorX	XXXXXXXXXXXXXXXXXXXXXXXXXXXX mg/kgXXXXXX HEDXXXX - XXXXXXXXXXXXXXXXXXXX
SFX Safety FactorX	XXXXX HEDXXXXX MRSDX
KX	XXXXXXXXXXXXXXXXXXXXXXXXXXXX
KmX	mg/kgXXXX mg/m2 XXXXXX
WX	XXXXX kgX

????

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EMA. Guideline on Strategies to Identify and Mitigate Risks for First-In-Human Clinical Trials with Investigational Medicinal Products. 2007.7

Dedrick RI. Animal Scale-Up. J Pharmacokinet Biopharm 1973; 1: 435-461

Mordenti J. Man versus Beast. J Pharm Sci, 1986; 75: 1028-1040

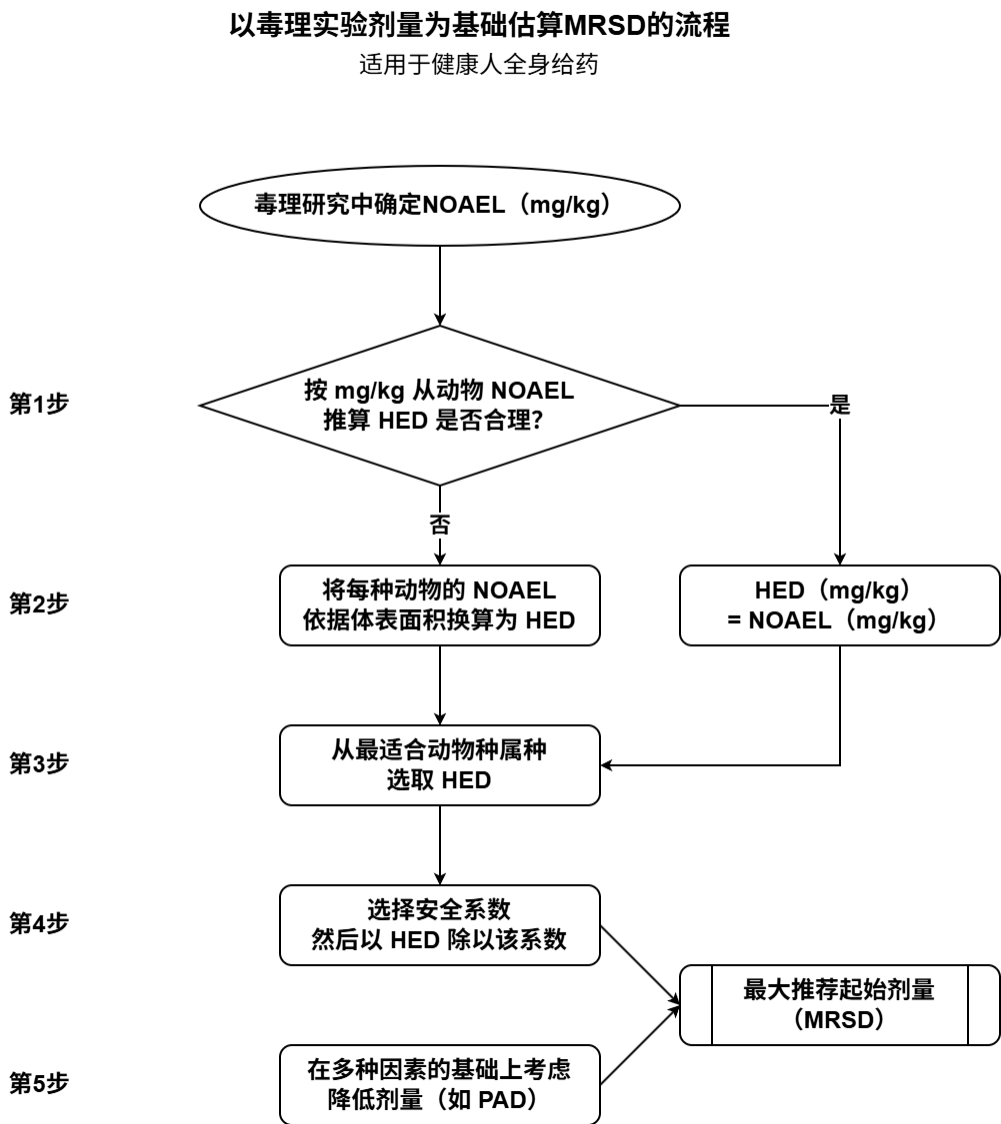
Boxenbaum H. Interspecies Scaling, Allometry, Physiological Time and the Ground Plan of Pharmacokinetics. J Pharmacokineti Biopharm. 1982: 10: 201-207

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Mahmood I. Balian Jd. Interspecies Scaling: Predicting Pharmacokinetic Parameters of Antiepileptic Drugs in Humans from Animals with Special Emphasis on Clearance. J Pharm Scie. 1996: 85: 411-414

R. Scott Obach. Prediction of Human Clearance of Twenty-Nine Drugs from Hepatic Microsomal Intrinsic Clearance Data: an Examination of In Vitro Half-Life Approach and Nonspecific Binding to Microsomes. Drug Metabolism And Disposition. 1999:27:1350-1359

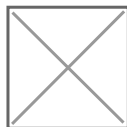
??A ?????????????MRSD???



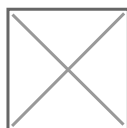
??B ?????(mg/kg)????????HED???

mg/kg

HED

[illegible]

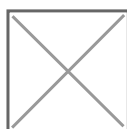
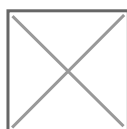
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S

cm² W

g□

[illegible]

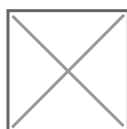
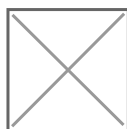
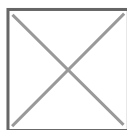
--	--	--	--	--

60kg

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W

3. □ mg/kg□□□□□□□□

 mg/m^2 

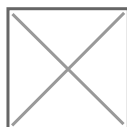
4.

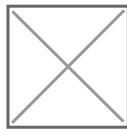
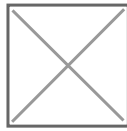
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[illegible]mg/m²

--	--	--	--	--	--

km□





5. Km

	kg	m ²	Km
	60	1.6268	36.88
	20	0.80	26.47
	0.020	0.006086	3.29
	0.080	0.1602	4.99
	0.150	0.02484	6.04
	0.300	0.04029	7.45
	0.300	0.04029	7.45
	0.400	0.404925	8.12
	1.8	0.14073	12.79
	10	0.46580	21.47
^a	3	0.20102	14.92
	20	0.7557	26.47
	40	1.2259	32.63

a

EXCEL

=10*W/POWER(10,(LOG10(W)*0.698+0.8762))

6. NOAEL mg/kg HED

NOAEL		HED
	mg/kg ÷ (km ₁ /km ₂)	
15 mg/kg 10kg,	15 mg/kg ÷ (36.88/21.47) =	8.7 mg/kg
50 mg/kg 150g,	50 mg/kg ÷ (36.88/6.04) =	8.2 mg/kg
50 mg/kg 200g,	50 mg/kg ÷ (36.88/6.6) =	8.9 mg/kg

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Diagram illustrating a 1D lattice structure with 16 sites. The top row shows all sites. The bottom row shows sites 1, 3, 5, 7, 9, 11, 13, and 15, with double quotes between sites 7 and 9, indicating a specific configuration or state.

The diagram illustrates a sequence of rows of rectangular blocks. The first row at the top contains 25 blocks. Each of the following six rows contains one fewer block than the row immediately above it. The rows are arranged such that the blocks are aligned to the left, creating a triangular shape. The total number of blocks is 100.

????????????????

Pharmacokinetics, PK

Model-informed Drug Development, MIDD

2/28

Dose-exposure-response relationship, D-E-R relationship

?????????????

Mechanism, PoMProof of Concept, PoC

Proof of

PoC

PoM

?????????????

PK/

-

-

PKPK

????????????

Physiologically-based Pharmacokinetic, PBPK



6. ?/???????PK??



PK/PD

The diagram shows two horizontal bars. The top bar is a single continuous rectangle divided into 20 equal vertical segments, representing 100%. The bottom bar is a single continuous rectangle divided into 10 equal vertical segments, representing 50%.

2. ???????

Diagram illustrating a 400-bit bus structure. The bus is divided into four segments: a top segment of 40 boxes, a second segment of 32 boxes, a third segment of 32 boxes, and a bottom segment of 16 boxes. To the right of the bottom segment, the text "PK/PD" is followed by 8 boxes.

Diagram illustrating a block cipher structure. The input is a 64-bit block, split into two 32-bit halves. The top half is processed by a function F , and the bottom half is processed by a function G . The outputs are then swapped and combined to form the 64-bit output block. The diagram uses rectangles to represent the 32-bit halves and arrows to show the flow of data.

QT

--	--	--	--	--	--	--	--

--	--	--	--	--

ICH []QT/QTcQT []TQT []QTc []CQT[]

[]

[]

QT/QTc []CQT

[]TQTQT/QTc []CQT

[]

?????-??????

1. ????

[]

[] -

[]

[] -

[]PoM []PoC []

[]-[]/[]-[]PK

[] -

[]/[]/

[]

[] -

[] -

[]

[]-[]

2. ????

[]-[]-[]/

[]

[] -

[]

[] -

[]

???????

??????????

PK
PK
PK
PK

PK
DDI /
PK
PK PD

??????????

PBPK
PBPK

PK PD

PK PD

-

PK

DDI

??????????

-

Pharmacology, QSP Machine Learning

Quantitative Systems
Artificial Intelligence, AI

??????????

??????????

PK PD -

??????????

/

DDI

CQT

FcRn
 PK

Fc

FcRn

PK

??????

???????

PD

 PK PK/PD

 PD

Mechanism of Action MoA PK
DDI
/ PK PD PK/PD
PK PD

-

PK PD

-

-
PK

-

-

PK

??????

???????



???????

□□	□□
Concentration QT (CQT)	□□□ QT □□□□□□□ QTc □□□□□□□□□□□□□□□□
Corrected QT interval (QTc interval)	□□□□□ QT □□□ QTc □□□□□□□□□□□□
Dose-Exposure-Response relationship (D-E-R relationship)	□□ -□□ -□□□□□□□□□□□□□□□□
Drug-Drug Interaction (DDI)	□□ - □□□□□□□□□□□□□□□□□□ □
Mechanism of Action (MoA)	□□□□□□□□□□□□□□□□□□ □
Model-informed Drug Development (MIDD)	□□□□□□□□□□□□□□□□□□ □□□□□□□□□□□□□□□□
Multiple ascending doses (MAD)	□□□□□□□□□□□□□□
Physiologically-based Pharmacokinetics (PBPK)	□□□□□□□□□□□□□□□□□□ □□□□□□□□□□□□□□□□□□ □□□□□□□□□□□□□□□□□□ □□□□□□□□□□
Proof of Concept (PoC)	□□□□□□□□□□□□□□□□
Proof of Mechanism (PoM)	□□□□□□□□□□□
Quantitative systems pharmacology (QSP)	□□□□□□□□□□□□□□□□□□ □□□□□□□□□□□□□□□□□□ □□□□□□□□□□□□□□
QT interval	QT □□□□□□□ QRS □□□□□ T □□□□□□□□□□□□□□□□□□
Single ascending dose (SAD)	□□□□□□□□□□□□□□
Thorough-QT/QTc study (TQT)	□□ QT □□□□□□□□□□ QT □□□□□□□□□□□□□□□□□□
Torsades de pointes (TdP)	□□□□□□□□□□□□□□□□□□ □□□□□□□□ QRS □□□□□□□□□□□□□□ QT □□□□□

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[1] ICH M4E: The Common Technical Document on Efficacy. 2016.

[2] ICH E4: Dose-Response Information to Support Drug Registration. 1994.

[3] ICH E14: The Clinical Evaluation of QT/QTc Interval Prolongation and Proarrhythmic Potential for Non-Antiarrhythmic Drugs. 2005.

[4] ICH E14: The Clinical Evaluation of QT/QTc Interval Prolongation and Proarrhythmic Potential for Non-Antiarrhythmic Drugs. Questions and Answers (R3). 2015.

[5] 国家药品监督管理局. 药品注册管理办法. 2020年12月.

[6] 国家药品监督管理局. 药品注册管理办法. 2020年12月.

[7] 国家药品监督管理局. 药品注册管理办法. 2020年12月.

[8] 国家药品监督管理局. 药品注册管理办法. 2014年7月.

[9] 国家药品监督管理局. 药品注册管理办法. 2021年1月.

[10] 国家药品监督管理局. 药品注册管理办法. 2021年2月.

[11] 国家药品监督管理局. 药品注册管理办法. 2021年12月.

[12] 国家药品监督管理局. 药品注册管理办法. 2021年12月.

[13] 国家药品监督管理局. 药品注册管理办法. 2021年12月.

[14] 国家药品监督管理局. 药品注册管理办法. 2021年12月.

????????????????????????????????
 ???????

Pharmacokinetics PK

PK

PK

PK

Diagram illustrating a 3D genome structure with interactions between a top loop and a bottom loop. The top loop contains a PK (pink) and a blue box. The bottom loop contains a PK (pink), a blue box, and a green box. Arrows indicate interactions between the loops.

[illegible]

PK

PK
PK

PK/PD
PK

1. /
- 2.
3. PK PD
4. PK PD
- 5.
6. PK PD /
- 7.
- 8.
9. /

??????

PK / PD
PK / PD

PK / PD

???????

PK
PK
PD

XXXXXXXXXXXXXXXXXXXX

PK XX

???????

XXXXXXXXXX

PK XXXXXXXXXXXXXXXXXXXX

PK

XXXXXXXXXXXXXXXXXXXXXXXXXXXX

/XXXXXXXXXXXX

1. ????

XXXXXXXXXXXXXXXXXXXX

pH

XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX

XX -XXXXXXXXXXXXXXXXXXXXXXXXXXXX

XXXXXXXXXX

PK

XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX

2. ????

XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX

XXXXXXXXXXXXXXXXXXXXXXXXXXXX

PK

XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX

XXXXXXXXXXXXXXXXXXXX

PK XXXXXXXXXX

PK X /X PD XXXX

XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX

PK

XXXXXXXXXX

PK XXXXXXXXXX

PK XXXXXXXXXXXXXXXXXXXX

3. ????

XXXXXXXXXXXXXXXXXXXX

PK

XXXXXXXXXXXXXXXXXXXXXXXXXXXX

PK

XXXXXXXXXXXXXXXXXXXX

XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX

XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX

XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX

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PK XXXXXXXXXXXXXXXXXXXXXXX

PKXXX

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XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX

XXXXXXXXXX

PK XXXXXXXXXXXXXXXXXXXXXXX

PK

XXXXXXXXXXXXXXXXXXXX

4. ????

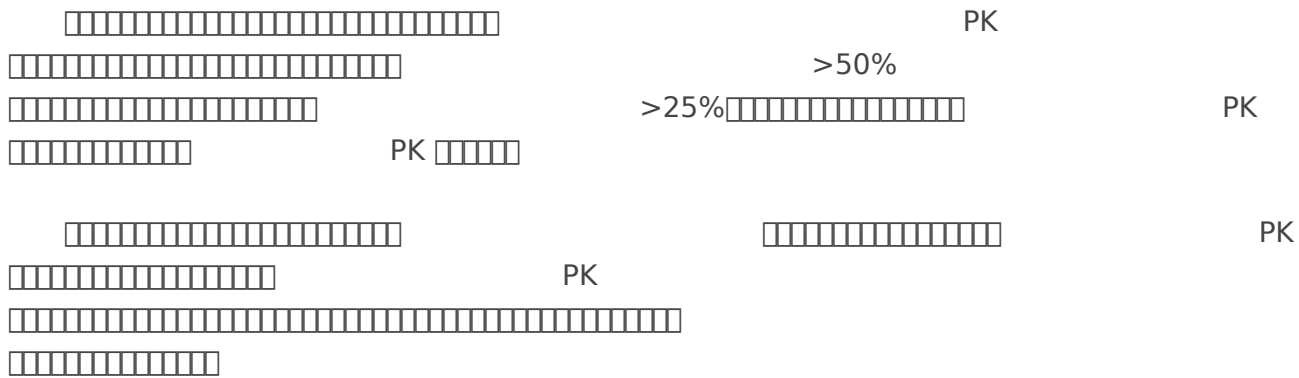
XXXXXXXXXXXXXXXXXXXX

PK XXXXXXXXXX

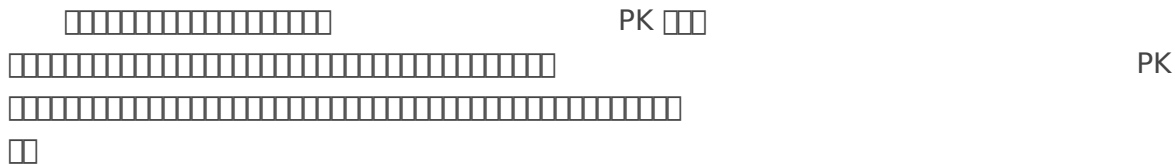
PK

XXXXXXXXXX

5. ????????



6. ???????



7. ????

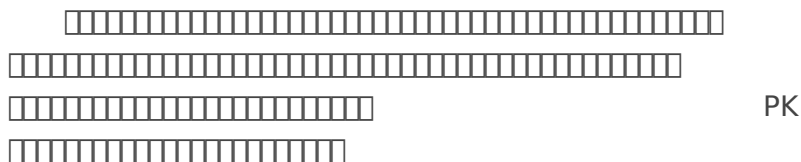
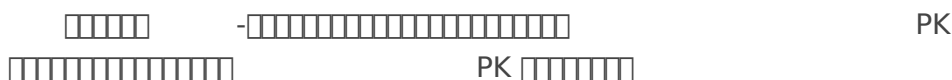


8. ??????

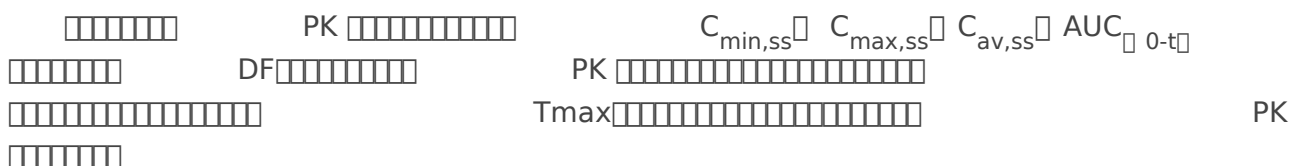


??????

??????????????



$t_{1/2}$ PK T_{max} C_{max} AUC_{0-t} $AUC_{0-\infty}$ V_d V_d/F K_{el} t
 MRT CL CL/F / PK



?????-??-??????

[] -PK [] PK
 [] PK [] Power
 Model [] PK [] - []
 [] PD [] - []

????????????

[] PK []
 []
 []

?????

[]

??????

[] PK
 []
 [] - [] PD [] PK/PD
 [] PK/PD []

[] PK
 [] /
 []

[] PK [] -
 [] - []

????

[1] [] . [] . 2005.

[2] [] . [] . 2021.

[3] European Medicines Agency. Guideline on strategies to identify and mitigate risks for first-in-human and early clinical trials with investigational medicinal products, EMA/CHMP/SWP/28367/07 Rev.1/ [] Committee for Medicinal Products for Human Use (CHMP) 20 July 2017.

[4] European Medicines Agency. Guideline on the use of pharmacogenetic methodologies in the pharmacokinetic evaluation of medicinal products [] EMA/ CHMP/37646/2009 Committee for Medicinal Products for Human Use (CHMP) 12 December 2011.

[5] [] . [] 9012 [] .2020.

NMPA

????????????????

????

Antibody drug conjugate ADC

ADC pH

ADC Pharmacokinetics PK “ ” Pharmacodynamics PD

ADC International Council for Harmonisation of Technical Requirements for Pharmaceuticals for Human Use ICH

ADC ADC

?????

ADC ADC ADC ADC ADC

The diagram shows a 1000-word memory array organized into 10 rows of 100 words each. The array is divided into four 250-word segments. The first segment (rows 1-4) is labeled 'DLT' and the second segment (rows 5-8) is labeled 'ADC'. The third segment (rows 9-12) is unlabeled, and the fourth segment (rows 13-16) is also unlabeled.

?2?PK ??

3 PD

PD

PK

3. ????

ADC [|||||] PK [] PD [|||||] PK [] PD

ADC [|||||] PK [] PD [|||||] PK [] PD

ADC [|||||] PK [] PD [|||||] PK [] PD

ADC [|||||] PK [] PD [|||||] PK [] PD

ADC

ADC

ADC

ADC

ADC

1

/

/

③

2.

ADC
ADC
ADC
ADC
P450 2D6
Antibody-dependent Cell-mediated
Cytotoxicity
ADCC
IgG
FcγRs
ADCC

3.

ADC
DDI
DDI
DDI
ADC

DDI
DDI
DDI
DDI

DDI
DDI
DDI
DDI

DDI
CYP
ADC

ADC
PK
DDI
ADC
IgG Fc
ADC
FcRn
ADC
PK

DDI

DDI

ADC

DDI

CYP

DDI

?????-????

ADC

ADC

E-R

-

ADC

E-R

ADC

E-R

ADC

E-R

ADC

ADC

ADC

E-R

PK/PD

???QT/QTc???

QT/QTc

QT/QTc

???????

Diagram illustrating the structure of a 16-bit ADC output. The output is shown as a 16-bit bus. The first 4 bits are labeled 'ADC'. The next 12 bits are labeled 'ADC'. The last 4 bits are labeled 'PK', 'PD', and 'ADC'.

[illegible]

Diagram illustrating the structure of various antibody-based therapies. The diagram shows five horizontal bars representing different molecules. The first bar is labeled "ADC" and consists of a single segment. The second bar is labeled "ADC" and consists of a single segment. The third bar is labeled "ADC" and consists of a single segment. The fourth bar is labeled "ADC" and consists of a single segment. The fifth bar is labeled "ADC" and consists of a single segment.

Diagram illustrating the 1000 Genomes Project data structure. The data is organized into a grid with columns for PK, PD, and ADA, and rows for ADC and PK. The grid is divided into sections for each chromosome (1-22, X, Y). The ADA column is the largest and most prominent, indicating a high density of data points for this variable across all chromosomes and individuals.

???????

[illegible]

Diagram illustrating the placement of ADCs (Analog-to-Digital Converters) in a 100-unit array:

- Row 1: ADC at the end (100 units).
- Row 2: ADC in the middle (50 units).
- Row 3: ADC at the beginning (0 units).
- Row 4: ADCs at both ends (0 and 100 units).

1.  /  ADC

2. ☐ QTc

?????

Input	Output
Drug to antibody Ratio [DAR]	[]
Antibody drug conjugate [ADC]	DAR [] 1 []
[]	DAR [] 0 []
[]	ADC []
Linker []	[]
[]	[]
Conjugated Payload []	[]
[]	[]

NMPA

????????????????????

1.

max

$T_{\max}$ T_{lag}

PK

2.


3.

4. [] / [] CRO

[]

[]

5. ISR ISR

6. 

[illegible][illegible][illegible][illegible][illegible]

1 PK BE -

C

AUC

PK

BQL

CRO

ISR

PMDA

Invalid