Endogenous Biomarkers for OATP1B: Preclinical to Clinical Translations

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Outlines

- Current strategies to assess clinical DDIs for OATP inhibitor
- Endogenous probes for OATP inhibition
 - Preclinical evaluation
 - Clinical studies
- **G** Summary

OATP inhibition decision tree



Variability in Ki or IC50 of Inhibitors



Vaidyanathan, et al. (2016) J. Clin. Pharmacol. 56 Suppl 7, S59-72

Relationships between observed AUCR and Imax(free)/ Ki or IC50.



Vaidyanathan, et al. (2016) J. Clin. Pharmacol. 56 Suppl 7, S59-72

Summary of the challenges in assessing potential OATP inhibition DDIs

 The potential for a high rate of false-positive (and negative) prediction has been a particular concern

(Prueksaritanont et al., 2013; Tweedie et al., 2013)

- Other challenges
 - Dilemma: the timing for human DDI studies and the selection of does
 - Selection of probe substrates
 - Extrapolating DDIs to different population or ethnic group
 - Complex DDIs

Supplemental material to this article can be found at: http://jpet.aspetjournals.org/content/suppl/2016/02/23/jpet.116.232066.DC1.html

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Coproporphyrins I and III as Functional Markers of OATP1B Activity: In Vitro and In Vivo Evaluation in Preclinical Species^S

Hong Shen, Jun Dai, Tongtong Liu, Yaofeng Cheng, Weiqi Chen, Chris Freeden, Yingru Zhang, W. Griffith Humphreys, Punit Marathe, and Yurong Lai *Pharmaceutical Candidate Optimization, Bristol-Myers Squibb Company, Princeton, New Jersey* Received January 12, 2016; accepted February 12, 2016

> Supplemental material to this article can be found at: http://jpet.aspetjournals.org/content/suppl/2016/06/17/jpet.116.234914.DC1.html

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Coproporphyrins in Plasma and Urine Can Be Appropriate Clinical Biomarkers to Recapitulate Drug-Drug Interactions Mediated by Organic Anion Transporting Polypeptide Inhibition^{SI}

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Learning from "Rotor Syndrome"

Rotor Syndrome: complete deficiency of OATP1B1 &1B3



Genotype	Urinary Ratio of CP I (%)	Urinary Ratio of CP III (%)
Reference Subjects	~30	~70
Rotor Syndrome	~70	~30

Wolkoff et al., *Am J Med* 1976, 60: 173

Van de Steeg et al., JCI 2012, 122:519

Coproporphyrin I and III (CP I and III)

- CP I and CP III are byproducts of heme synthesis.
- They are not further metabolized by the liver and excreted in bile and urine. CP I predominates (70%) over CP III in bile whereas the reverse is found in urine.



Substrate specificity of CPs

- CP-I and CP-III are substrates of OATP1B1 and 1B3
- Both CP-I and III are <u>not</u> substrates for OAT1/2/3/4, OCT2, MATE1 and MATE2k
- CPs are metabolically stable in human, monkey and mouse hepatocytes

Deletion of Oatp1a/1b genes increased plasma conc. and urinary excretion of CPs in mice



• Oatp1a/1b cluster genes deleted: *Slco1a1, 1a4, 1a5, 1a*6 and *1b2*.

Shen et al., JPET 2016, 357: 382

CsA increased plasma CPs and urinary excretion of CP-I in monkeys



Fold Change	CP I	CP III
C _{max}	3.3	6.9
AUC _(0-24 h)	2.6	5.2

RIF significantly increased plasma conc. and urinary excretion of CPs in monkeys



Clinical study on CPs as probes for OATP inhibition

Open label, three-treatment, three-period, single dose crossover study in twelve healthy male adult subjects without carrying <u>OATP1B1*5, *15, *17 polymorphisms</u>.



RIF Pharmacokinetics

- C_{max} and $C_{24 h}$ of ~30 and 1 μ M, respectively.
- IC50 values of OATP1B1/1B3-mediated CP transport were
- Systemic exposure of RIF was comparable to literature reports
- Similar exposure when dosed alone or in combination with RSV



Paramatar	RIF	
	RIF alone	RIF + RSV
Cmax (µM)	26.7 ± 10.7	30.6 ± 7.8
Tmax (h)	2.5 ± 1.7	2.2 ± 0.9
AUC _(0-24 h) (µM*h)	206.8 ± 70.9	219.5 ± 57.9
AUC _{tot} (µM*h)	215.4 ± 79.4	224.5 ± 61.9
T1/2 (h)	4.2 ± 1.4	3.5 ± 1.2

RIF significantly increased plasma conc. of CP-I and CP-III, and urinary excretion of CP-I



Fold Change	CP-I (RIF, RIF+RSV)	CP-III (RIF, RIF+RSV)	RSV	
C _{max}	5.7, 5.9	5.4, 6.5	13.2	
AUC _(0-24 h)	4.01, 3.8	3.4, 3.3	5.0	
X _{e (0-24 h)}	3.6, 3.4	2.1, 1.34*	1.6, 1.4	*NS

Urinary excretion of CPS



CP-I appears to be passively filtered, but CP-III is actively secreted with a large inter-individual variability

Plasma conc. of CPs in OATP1B1*1b heterozygous (9) and homozygous (3)



20% reduction of CP-I in plasma was detected in *1b homozygous, but there is no statistically significant difference (likely small N?)

RIF increased (marginally) the levels of total- and indirect-, but not direct-bilirubin in plasma



Plasma conc. of bile acids—with statistical significance



Fold changes	GDCA (RIF, RIF+RSV)	TDCA (RIF, RIF+RSV)
C _{max}	7.6, 4.3	4.4, 3.4
AUC _(0-24 h)	6.5, 4.0	3.6, 3.1

Endogenous probes for DDI assessment

- Specific
 - A specific endogenous substrate of the transporter
 - Not a biomarker of a disease or a dietary constituent—
 - rejecting bile acids and bilirubin as probes
 - Reflect instant response, v.s. delayed (compensatory) effects

• Predictive and Translational

- Correlate with the extent of transporter inhibition, e.g fold increases of statins?
- Reflect site of inhibition, e.g Imax/IC50

Accessible

- Non-invasive sampling from either blood or urine
- Can be monitored during early drug development phase, e.g Phase I does finding trials

Reproducible

• Rapid, accurate and reproducible detection, e.g LC/MS

Cost effective

Manageable lists

Summary

- Plasma CP-I and CP-III are relatively consistent, with minimum fluctuation over the three study periods.
- CPs are determined by LC/MS which is is amendable to high throughput, GLP setting
- OATP inhibitor, RIF, significantly increased plasma conc. of CP-I and CP-III.
- CPs in plasma or CP-I in urine could serve as endogenous probes for OATP inhibition, and be monitored during phase I dose finding trials
- Changes of CPs with less potent OATP inhibitors, and the impacts of genetic polymorphisms, gender/age, disease state, and organ impairments etc need to be further investigated

Integration of endogenous probes into OATP DDI decision trees



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