



Navigating the Transporter Wave of 2013: A Review of the Seven Recent ITC Publications on Drug Transporters

> Greg Loewen, Andrea Wolff, Amanda Hays, Ph.D. XenoTech Drug Transport Group

XenoTech – brief introduction

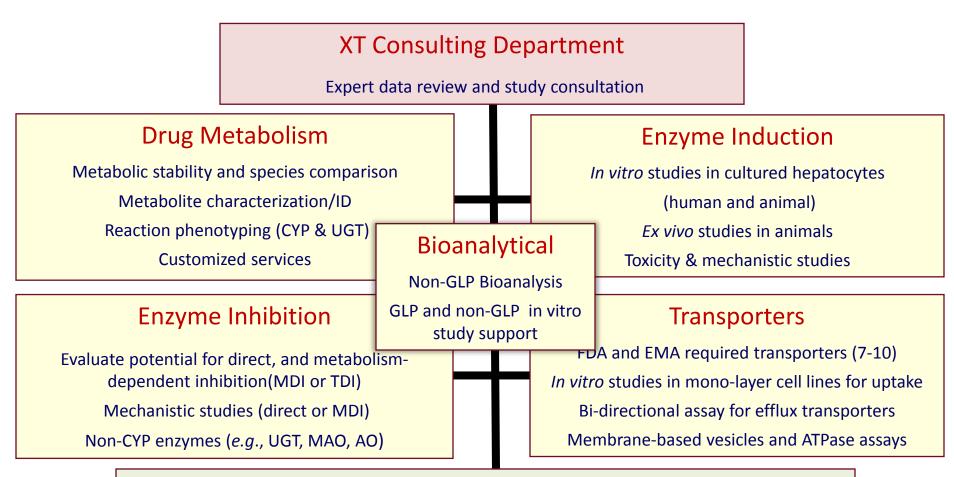
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Main focus: In Vitro products and services to predict clinical DDI as outlined in the FDA and EMA guidance documents on DDI

- Located in Lenexa, KS
- GLP compliant CRO, founded in 1994 by Dr. Andrew Parkinson
- ~110 employees
- Global: Distributors in Europe, Japan, Korea, India, China and Singapore
- Partners include Sekisui
 Medical (Japan), Cypex
 (Scotland), Xenometrics (USA)
- July 2008 XenoTech acquired by Sekisui (SMD)
- •NEXT IND, Partnership of small CROs





Discovery and Late-Preclinical Support

Medium-throughput ADME screening studies

Definitive ADME and DDI studies to support design and necessity of clinical studies

XenoTech support groups

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XenoTech is big and diverse enough to have independent support groups such as our analytical group, knowledge management and QA which add value to your study but we are small enough the groups work seamlessly to provide a high quality final product



- Follow recommended methods outlined in FDA and EMA guidance documents
- Review and discuss physiochemical properties and pk data to suggest most relevant studies
- GLP compliant facility high standards of documentation and traceability
- Strong in house LC-MS/MS support samples analyzed the same day
- Collaborate with our parent company in Japan – doubling our size and knowledge base
- Widest range of transporter assays available
- Fully licensed to use all cell lines and transporters
- Been involved in the transporter field since 2006

Drug Transport services at XenoTech



All transporter assays requested by the FDA and EMA and we help you with study design

MDCKII, LLC-PK1 and Caco-2 cells for transwell assays (P-gp and BCRP)

Vesicles for efflux transporters (MRPs, BSEP, BCRP, P-gp etc)

Transfected HEK293 and S2 cells for SLC (uptake) transporters, (Oocytes capability)

Hepatocyte uptake and Lysosomal trapping assays

Partner with Sekisui on certain assays and study design



Efflux

- MDR1 (P-gp) (human, mouse 1a/1b, rat 1a/1b, monkey, dog)
- BCRP (human and mouse)
- BSEP
- MRP2 (and MRP1, MRP3, MRP4, MRP5)
- MATE1
- MATE2K

Uptake

- OATP1B1 1B3 2B1 -1A2
- OAT1 2 3 4 7
- OCT1 2 3
- OCTN1 N2
- NTCP1 NTCP2 (ASBT)
- NPT1
- PEPT1 2
- URAT1
- OST α/β
- LAT



Thanks to the ITC!



Lead Author	Title	Presenter
KM Giacomini	International Transporter Consortium Commentary on Clinically Important Transporter Polymorphisms	Amanda
KM Hillgren	Emerging Transporters of Clinical Importance : An Update From the International Transporter Consortium	Amanda
D Tweedie	Transporter Studies in Drug Development: Experience to Date and Follow-Up on Decision Trees From the International Transporter Consortium	Amanda
KLR Brouwer	<i>In Vitro</i> Methods to Support Transporter Evaluation in Drug Discovery and Development	Greg
X Chu	Intracellular Drug Concentrations and Transporters: Measurement, Modeling, and Implications for the Liver	Greg
MJ Zamek- Gliszczynski	ITC Recommendations for Transporter Kinetic Parameter Estimation and Translational Modeling of Transport-Mediated PK and DDIs in Humans	Andrea
JC Kalvass	Why Clinical Modulation of Efflux Transport at the Human Blood–Brain Barrier Is Unlikely: The ITC Evidence-Based Position	Andrea



International Transporter Consortium Commentary on Clinically Important Transporter Polymorphisms

KM Giacomini¹, PV Balimane², SK Cho³, M Eadon⁴, T Edeki⁵, KM Hillgren⁶, S-M Huang⁷, Y Sugiyama⁸, D Weitz⁹, Y Wen¹⁰, CQ Xia¹¹, SW Yee¹, H Zimdahl¹² and M Niemi¹³; on behalf of the International Transporter Consortium

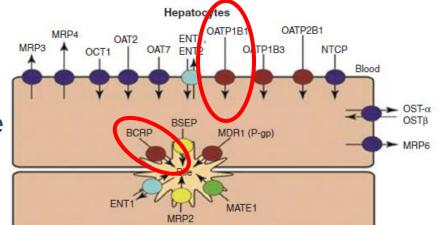
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OATP1B1 and BCRP



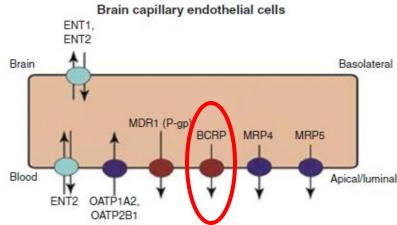
• OATP1B1 (*SLCO1B1*)

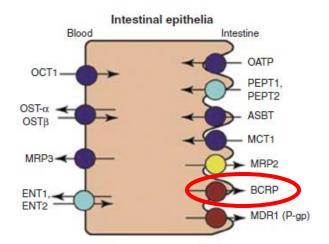
- ➢ c.521T>C, p.V174A
- Increased statin plasma exposure



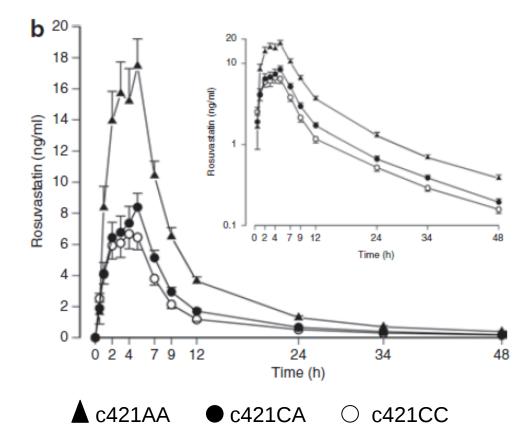
• BCRP (ABCG2)







Clinically relevant transporter polymorphisms



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- 1. Conduct *in vitro* studies to determine whether a drug is a <u>substrate</u> of key transporters (i.e., OATP1B1 and BCRP).
- 2. Consider conducting a DDI study with a transporter <u>inhibitor</u> as recommended in the FDA guidance.
 - Small therapeutic window, AUC > 2
- 3. If a pharmacogenomic study is suggested, design well-powered studies that consider <u>variants</u> in drug transporters that have been shown to interact with the drug in *in vitro* studies.
- 4. Collect DNA samples from subjects enrolled in studies. Include sufficient <u>numbers of subjects</u> with <u>variant alleles</u> representative of various <u>race/ethnic</u> groups.
- 5. Associate SNPs with drug concentrations, response and safety.
- 6. Apply a <u>multiscale</u> systems pharmacology approach that incorporates all data (*in vitro* and *in vivo* human studies).



KM Hillgren¹, D Keppler², AA Zur³, KM Giacomini^{3,4}, B Stieger⁵, CE Cass⁶ and L Zhang⁷; on behalf of the International Transporter Consortium

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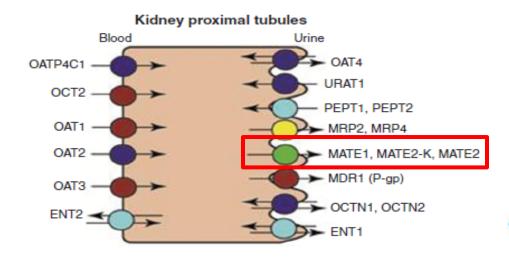
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P-gp (MDR1, ABCB1)

- BCRP (ABCG2)
- > OATP1B1 (SLCO1B1)
- > OATP1B3 (SLCO1B3)
- ➢ OCT2 (SLC22A2)
- > OAT1 (SLC22A6)
- > OAT3 (SLC22A7)

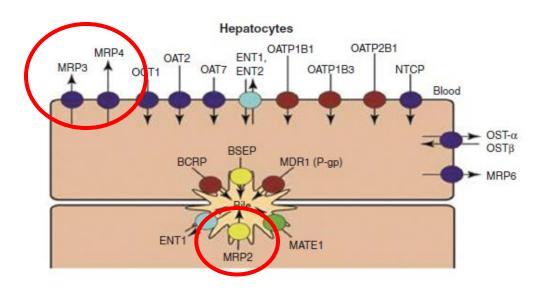


- MATE1, MATE2, MATE2K.
- Expressed at the apical membrane of renal proximal tubules.
- Secrete cations and zwitterions into urine.
- Cation/H⁺ antiporter in tandem with OCT2 (Metformin)
- Current evidence for clinical DDIs is strongest for MATE1 and MATE2K.
- MATE SNPs have been linked to clinical effects in metformin-treated subjects.
- Suggest in vitro models





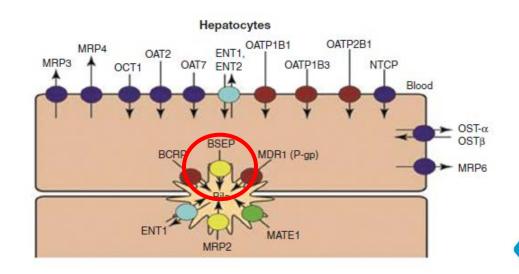
- MRP2, MRP3, MRP4.
- Unidirectional ATP-dependent efflux pumps.
- Hepatocytes, intestinal epithelia, kidney tubules.
- Anionic substrates, drugs and conjugates.
- A SNP with complete loss of MRP2 functionality contributes to hyperbilirubinemia (Dubin-Johnson Syndrome).
- Inside-out membrane vesicles



BSEP and hepatotoxicity



- Rate-limiting step of bile salt transport across hepatocytes.
- No backup system.
- SNPs cause progressive cholestatic liver disease.
- Unlikely involved in drug disposition, inhibition may contribute to adverse cytotoxic events.
- Isolated membrane vesicles.





• For certain drug classes, should consider:

- hENT1 (anticancer nucleoside analogs)
- Pept1 (oral dosed peptide-like drugs)

• Model systems:

- Oocytes
- Cell lines
- Knock-out mice





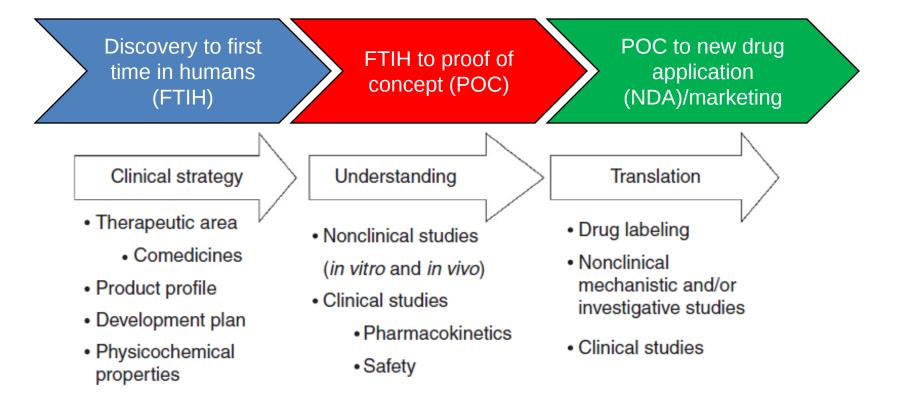
- NMEs that are actively renally secreted are recommended for *in vitro* investigation as MATE1 or MATE2K substrates in addition to being investigated as OAT1, OAT3 or OCT2 substrates.
- If signs of cholestasis are seen, retrospective analysis of inhibition of MRPs and BSEP can help determine the mechanism of toxicity.
- hENT1 and Pept1 are examples of transporters that do not need to be routinely screened but can contribute to efficacy and distribution.

Transporter Studies in Drug Development: Experience to Date and Follow-up on Decision Trees From the International Transporter Consortium

D Tweedie¹, JW Polli², E Gil Berglund³, SM Huang⁴, L Zhang⁴, A Poirier⁵, X Chu⁶ and B Feng⁷; On Behalf of the International Transporter Consortium

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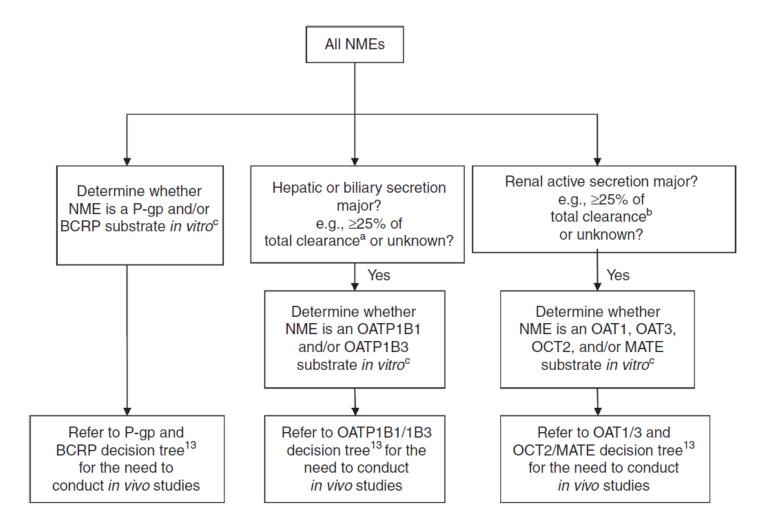


Central tenet is the clinical plan, which considers the therapeutic area, comedicines, and the patient population

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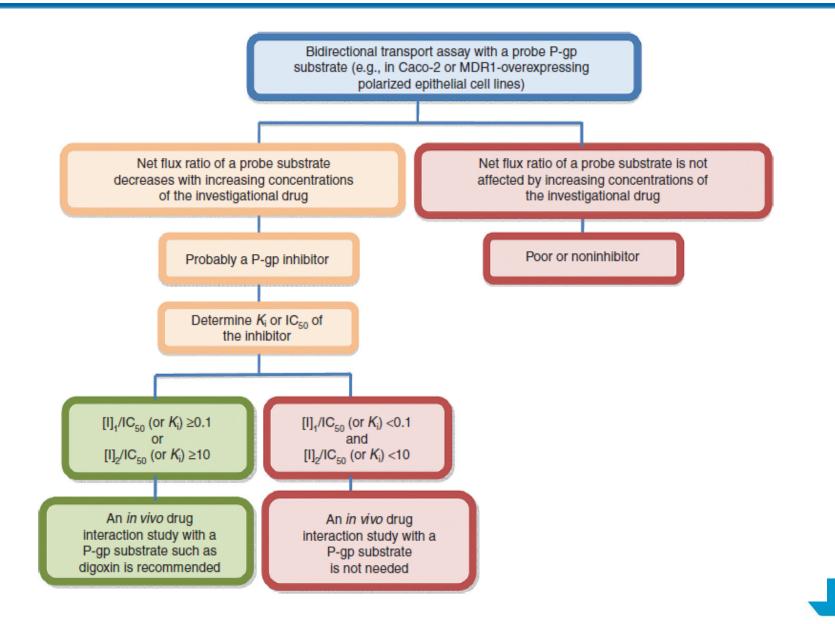
Evaluation of investigational drugs as substrates of transporters





P-gp/BCRP inhibition tree





Summary



- Flow chart presented for timing of Transport studies
- Discussion of the integration of transport studies into the clinical plan (*e.g.*, therapeutic area, co-meds, patient population, physicochemical properties, PK)
- Three decision trees presented
- Summary of approved compounds that 'failed' the FDA cutoff for clinical DDI study (table 1)
- Summary of PMR and PMC (table 2)
- Three case studies presented
 - Applying P-gp/BCRP inhibition decision trees
 - Complexity of DDI involving OATP transporters
 - Assessment of transporter-mediated DDIs for livertargeting compounds



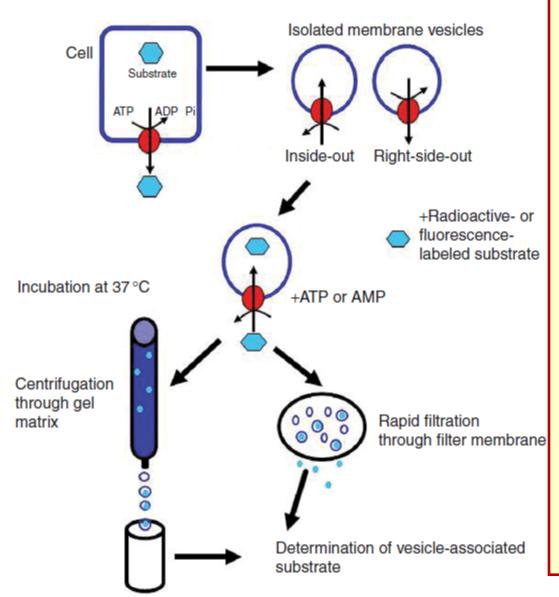
In Vitro Methods to Support Transporter Evaluation in Drug Discovery and Development

KLR Brouwer, D Keppler, KA Hoffmaster, DAJ Bow, Y Cheng, Y Lai, JE Palm, B Stieger and R Evers; on behalf of the International Transporter Consortium

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Vesicle assays





- ABC transporter expressing vesicles are prepared
 - From Spodoptera frugiperda insect cells (Sf9 or Sf21) infected with a cDNA containing baculovirus
 - cDNA-transfected mammalian cell lines; HeLa, V79 hamster, HEK293, MDCK, and MDCKII
 - Tissue including kidney and liver
- Homogenization method is very important
- Substrates and inhibitors listed in the article
- MRP2, BSEP most common, also MRPs, P-gp and BCRP



- Cells are transfected with full length cDNAs encoding a transporter of interest
 - Plasmids introduced into the host cell line either chemically, physically, or by retroviral transduction to create a stable transfection

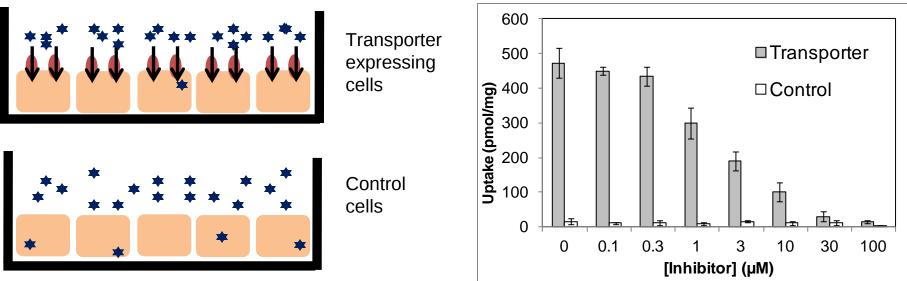
(and transiently transfected but not preferred)

- Uptake transporters (SLC):
 - HEK293, CHO, MDCKII, S_{2} , monolayer on a the bottom of cell culture plate
 - Oocytes: in suspension
- Efflux transporters (ABC)
 - LLC-PK1, MDCKII
 - Bidirectional transport or transwell assay

Uptake (SLC) transporter assays



- Test system: transporter transfected cells grown in cell culture plates
- Inhibition: measure the effect of the test article on the accumulation of a probe substrate
- Substrate: measure the accumulation of the test compound in transfected and control cells
- Hepatic: OATP1B1, OATP1B3, OCT1
- Renal: OAT1, OAT3, OCT2, MATE1, MATE2K



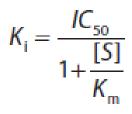


Accumulation is rapid and kinetics should be studied in the linear range of time and protein concentration

Solubility and non-specific binding should be considered – solvents and BSA

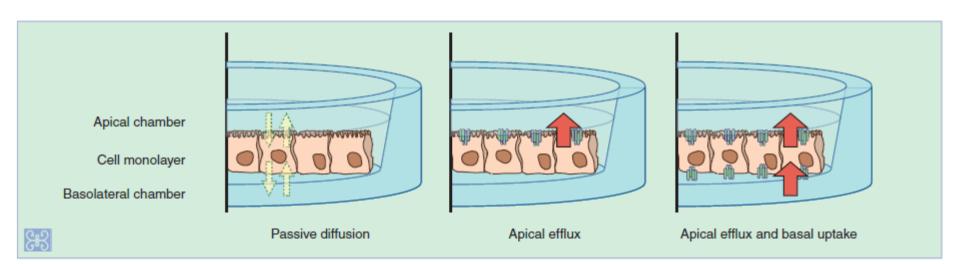
No consensus on the best way to calculate inhibition constants for transporters **yet**

- 1. For competitive inhibitors, the IC₅₀ value depends on the substrate concentration the *Ki does not*
- 2. IC₅₀ values will approach *Ki* if a substrate concentration far below the *Km* is used
- *3. If the* Cheng–Prusoff equation is used to estiamte the Ki the **probe substrate concentrations within twofold of the** *Km value.*
- 4. The Cheng–Prusoff equation assumes that the inhibition is competitive in nature



Bidirectional transport (transwell)

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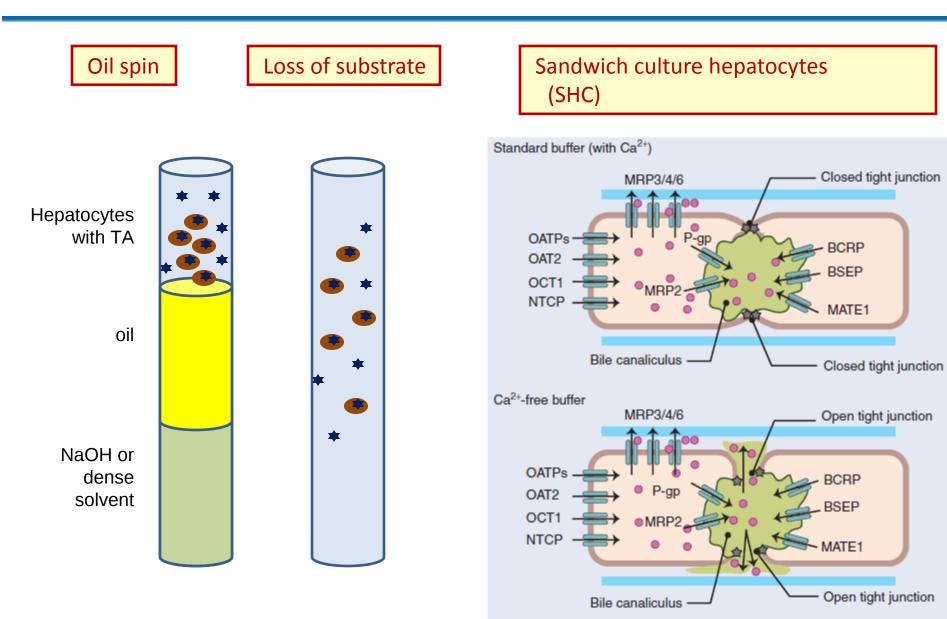


- Typically for P-gp and BCRP
- MCKDII, LLC-PK1 or Caco-2 cells
- Compound is loaded on the Apical (A) or Basolateral (B) side and permeability across the monolayer is measured and efflux ratio (ER) determined (BA ÷ AB)
- ER > 2 typically suggests active transport
- Difficult to correlate ER to extent of active transport *in vivo*

- Binding and metabolism can complicate data interpretation, recovery should be > 70%
- High permeable compounds; saturation Low permeable compounds uptake transporter
- Caco-2 can be used as a model for oral absorption
- Inhibition: several IC₅₀ calculations available (Bentz-Lee paper)

Hepatocytes







- **Table 1:** Vesicle substrates and inhibitors
- Table 2: Cell system substrates and inhibitors
- **Table 3:** Commonly used equations for calculation of kinetic parameters in vesicles, cell lines, and hepatocytes
- **Table 4:** Applications, strengths, and limitations of various in vitro transporter assay systems
- **Table 5:** Integration of in vitro and in vivo data to determine the role of transporters in compound absorption, distribution, clearance, and DDIs
 - Scientific question:
 - Observations to support In vitro transporter investigations:
 - In vitro tools to address scientific hypothesis:
 - Outcome of in vitro experiments:
 - Potential follow-up studies:



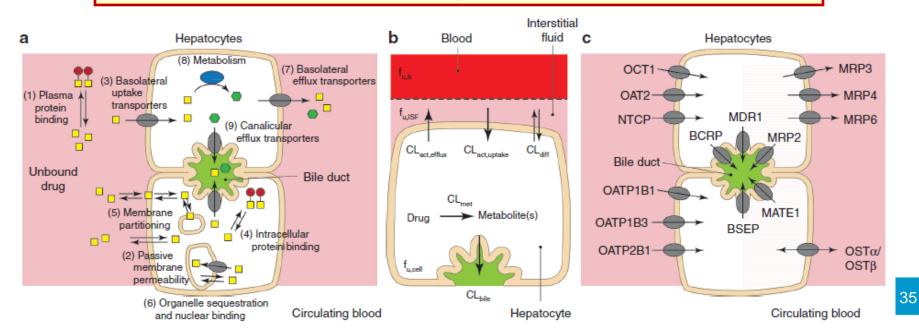
X Chu, K Korzekwa, R Elsby, K Fenner, A Galetin, Y Lai, P Matsson, A Moss, S Nagar, GR Rosania, JPF Bai, JW Polli, Y Sugiyama and KLR Brouwer; on behalf of the International Transporter Consortium

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Factors that influence intracellular drug disposition in the liver

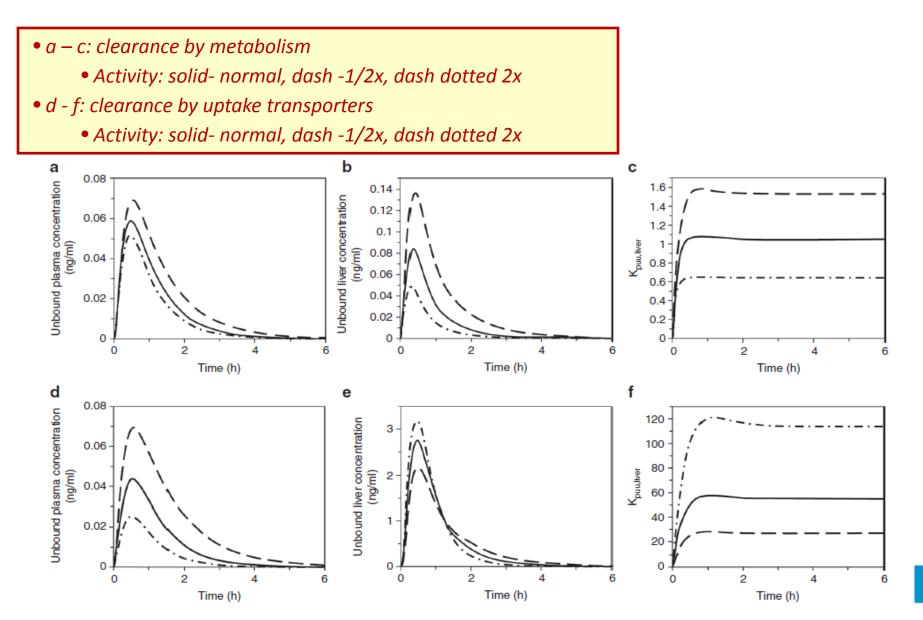


- Blood flow, mixing of portal and venous blood
- K_{puu,liver} is defined as the steady-state liver-to-sinusoidal blood partition coefficient for unbound drug
- function of CL_{diff}, CL_{act,uptake}, CL_{act,efflux}, CL_{bile}, and Cl_{met}
- High permeability compunds: K_{puu,liver} =1
- Metabolized or effluxed: K_{puu,liver} <1
- Active uptake: K_{puu,liver} >1
- Organelle or membrane distribution: K_{puu,liver} >1



PBPK model

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Summary of tables



- **Table 1:** Examples of studies in which membrane transporters have been shown to alter hepatocellular drug concentrations
 - MRP2, MRP3, BCRP, OCT1, OATP1B1, MATE1
- **Table 2:** *In vitro, in situ,* and *in vivo* models to estimate the intracellular concentrations of drugs and metabolites in the liver
 - Systems, Advantages, Disadvantages and examples of parameters estimated
 - Vesicles, recombinant proteins, hepatocytes, SCH, perfused liver, animal models, KO animals, Human *in vivo* studies
- **Table 3:** Direct and indirect methodologies for the estimation of intracellular drug concentrations
 - Direct or indirect method, analyte/matirx, detection method, utility/limitations/assumptions
 - Capillary electrophoresis, MSI: Nano-SIMS, MIMS, Raman microscopy, Nuclear microscopy, Microautoradiography, PET/SPECT imaging, PET imaging with simultaneous microdialysis, Bulk analysis
- **Table 4:** Summary of methodologies available to estimate intracellular fraction of unbound drug in cells
 - Method, Predictive equations developed for f_{u,cell}, Comments
 - Physiochemical properties, monolayer permeability, hepatoctye uptake and binding, transport kinetics, PBPK models



- Our understanding of intracellular drug disposition is rudimentary
 - e.g., trapping within the cell
- Currently available *in vitro and in vivo models have limited* capability to quantitatively predict the impact of transporters on intracellular drug concentrations
- Currently, there are no standardized, accepted methods to directly measure unbound intracellular drug concentrations
- Significant progress has been made in using modeling approaches to predict the effect of drug-metabolizing enzymes and transporters on the systemic exposure of drugs in preclinical species and humans

ITC Recommendations for Transporter Kinetic Parameter Estimation and Translational Modeling of Transport-Mediated PK and DDIs in Humans

MJ Zamek-Gliszczynski¹, CA Lee², A Poirier³, J Bentz⁴, X Chu⁵, H Ellens⁶, T Ishikawa⁷, M Jamei⁸, JC Kalvass⁹, S Nagar¹⁰, KS Pang¹¹, K Korzekwa¹⁰, PW Swaan¹², ME Taub¹³, P Zhao¹⁴ and A Galetin¹⁵; on behalf of the International Transporter Consortium

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Uptake transporters: 2 Step Method (Conventional)



- Single low substrate concentration, single incubation time point
- Cell systems
 - Linear rate conditions and time range, 4 and 37°C
 - Active uptake = Uptake at 37°C Uptake at 4°C
 - Transporter mediated uptake determined with inhibitors
 - Limitations
 - Limited membrane fluidity at 4°C
 - Lack of specific of inhibitor
 - Substrate dependent inhibition (OATPs)
- Expression systems
 - Active uptake = Uptake in transfected cells Uptake in control cells
 - Limitations
 - Assumes transport is an isolated process, Does not account for the bidirectional nature of passive diffusion, intracellular binding, metabolism, or active efflux
 - Data transformation required for parameter estimation
- Applicable for low permeable compounds

$$v = \frac{V_{max} \times C_{med}}{K_m + C_{med}} + CL_{diff} \times C_{med}$$



- Plated or suspended hepatocytes
 - Active transport, passive diffusion, and intracellular/ extracellular binding
- Multiple sub substrate conc. and times, 37°C
- Time points beyond time-linear range used to attain steady-state intracellular conditions
 - more accurate estimation of intracellular binding
- Assumes intracellular binding is not saturated
 - overestimation of the fraction unbound in cell $(f_{u,cell})$
- Active efflux not considered; also internalization of efflux transporters
- Metabolism not incorporated; uptake data obtained in the presence of ABT should be used for CYP substrates
- Applicable for compounds that are not metabolized or only metabolized by CYP enzymes

$$\frac{dC_{cell}}{dt} = \frac{\frac{V_{max} \times C_{med,u}}{K_{m,u} + C_{med,u}} + CL_{diff} \times C_{med,u} - CL_{diff} \times C_{cell} \times f_{u,cell}}{V_{cell}} \qquad \qquad \frac{dC_{med,u}}{dt} = \frac{-\frac{V_{max} \times C_{med,u}}{K_{m,u} + C_{med,u}} - CL_{diff} \times C_{cell} \times f_{u,cell}}{V_{med}}$$



- Plated or suspended hepatocytes
 - Active transport, passive diffusion, intracellular/ extracellular binding, metabolism
- Extended incubation times to obtain steady-state
- Both drug and metabolites are measured
- Data analysis is more complex but has less error
- Provides a mechanistic description of in vitro process
- Can account for canalicular efflux using sandwich-cultured hepatocytes
 - Single low substrate concentration, multiple time points
- Applicable for compounds that are metabolized by enzymes with no selective inhibitors (e.g., glucuronidation)
- Determination of uptake K_m and V_{max} is preferred to CL for $\ensuremath{\mathsf{IVIVE}}$

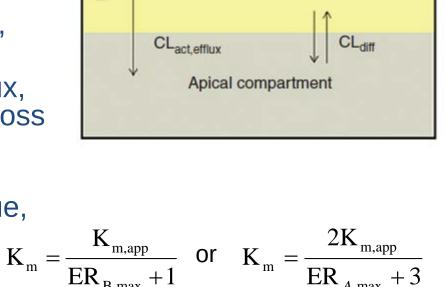
$$\frac{dC_{cell}}{dt} = \frac{\frac{V_{max} \times C_{med,u}}{K_{m,u} + C_{med,u}} + CL_{diff} \times C_{med,u} - C_{cell} \times f_{u,cell} \times (CL_{diff} + CL_{met,u})}{V_{cell}}$$

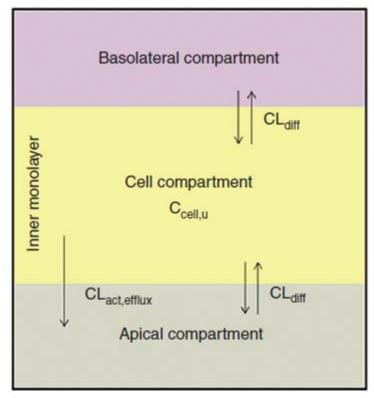


- Vesicles are best for low permeable compounds
- Commonly used for polar substrates (MRPs, MATES, BCRP)
- Inhibition of bile acid transport (BSEP)
- Not the test system of choice for P-gp and BCRP since these substrates are lipophilic and have higher permeability

Efflux transporters: Confluent cell monolayers (Model I)

- 3 compartment model (apical, basal, cellular)
- Basolateral ER_B : transcellular passive permeability/ $P_{app,A \rightarrow B}$
- Asymmetry $ER_{\alpha}: P_{app, B \rightarrow A}/P_{app, A \rightarrow B}$
- Cellular ER_c :cellular concentration in absence of efflux/cellular concentration in presence of efflux
- ER values used to calculate K_m , V_{max} and IC_{50}
- Assumes minimal lag time in flux, similar passive permeability across the apical and basolateral membranes
- If these assumptions are not true, this model is not valid



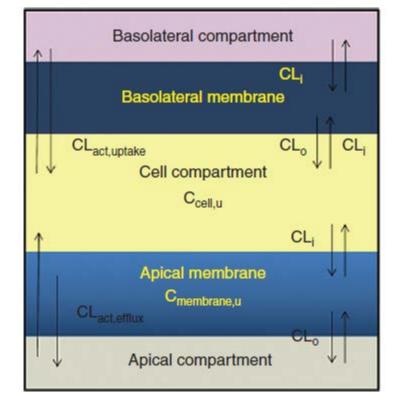


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Efflux transporters: Confluent cell monolayers (Model II)

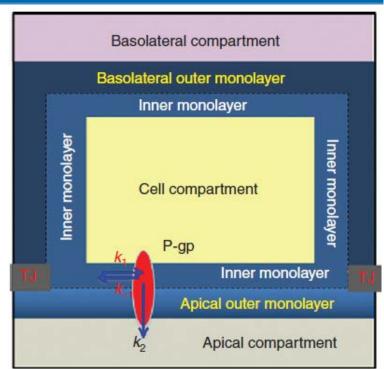
- 5 compartment model (includes apical and basolateral membrane compartments)
- Membrane-to-water partition coefficient (K_p) estimated in human liver microsomes
- K_m and IC₅₀ estimated in terms of the unbound drug concentration in membrane and cytosol, respectively
- 3- and 5-compartment models predict similar intracellular concentrations following apical addition of substrate, but differ in prediction of intracellular concentrations for flux in the $B \rightarrow A$ direction
- Appropriate when appreciable lag flux time and for prediction of tissues with basolateral exposure to the drug (e.g., liver or kidney).
- Metabolism can also be incorporated into the five-compartment model



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Efflux transporters: Confluent cell monolayers (Model III)

- Structural model: considers binding of substrate from within the inner leaflet of the apical membrane, with on $[k_{on} (M^{-1}s^{-1})]$ and off $[k_{off} (s^{-1})]$ rate constants and an efflux rate constant from P-gp into the apical chamber $[k_{out} (M)]$
- Estimates P-gp efflux active surface density, *T(0)*
- These parameters are used to calculate K_{m} and V_{max}
- Requires use of multiple initial drug concentrations and time points until steady state is reached between P-gpmediated efflux into the apical chamber and passive permeability from the apical chamber back into the cytosol
- K_i can be calculated from the IC₅₀
- Accounts for basolateral transport



$$\mathbf{K}_{\mathrm{m}} = \frac{\mathbf{k}_2 + \mathbf{k}_{-1}}{\mathbf{k}_1}$$

$$\mathbf{V}_{\max} = \mathbf{k}_2 \times T(0)$$



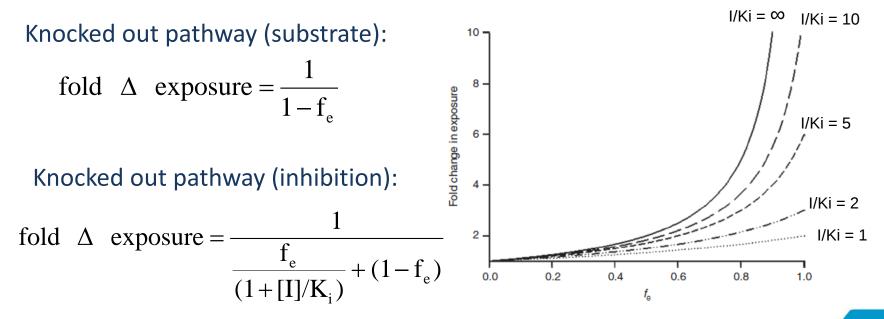


- Probe concentration $<< K_m$ for $IC_{50} = K_i$ (uptake transporters and efflux in vesicles)
- Method for K_i in polarized cell lines have not been agreed upon, IC_{50} is preferred in this test system
- Substrates requiring both basolateral uptake and apical efflux (e.g., digoxin) should use net value for inhibition to determine IC₅₀
- Evidence of substrate-dependent inhibition suggests transporters have multiple binding sites. Inhibition determined with relevant comedications and prototypical substrates.
- Preincubation may increase inhibition. With and without preincubation should be evaluated.
- Incubations should be conducted at relevant physiological pH

Knockout animal models



- With exception of OAT and MATE, all major transporters can be evaluated in knockout animals
- P-gp and BCRP data from knockout models: predict rate and extent of adsorption.
 - If complete adsorption, minimal in vivo efflux, may not need to conduct a clinical DDI study of intestinal efflux
 - May be useful for BBB predictions
- Fraction excreted (fe) method is preferred:





- Confirm compound exhibits similar pharmacokinetics between humans and knockout species
 - Species differences should be assessed in vitro to support translation
- Potential up- or down-regulation of transporters and drug-metabolizing enzymes may complicate interpretation

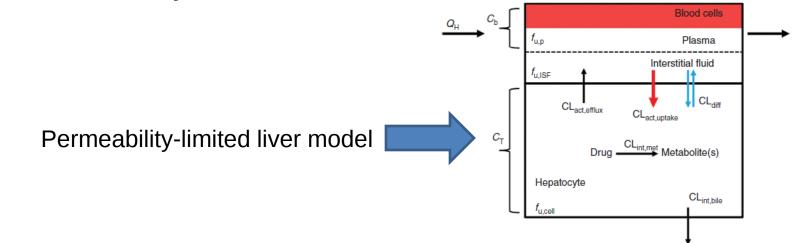


- Based on clearance concepts
- Assume a single transporter dominates victim drug adsorption, distribution and clearance
- Assume constant concentration of inhibitor or substrate during dosing
- May eliminate false negatives by assessing "worse-case scenario", but can result in false positives

PBPK for prediction of transporter-mediated PK



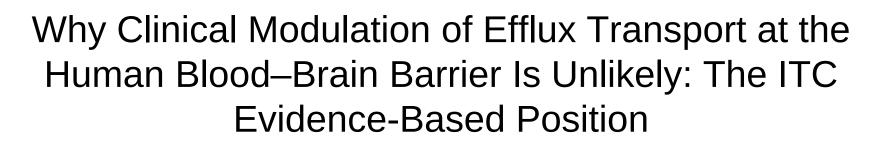
- Incorporate permeability-limited tissue compartments to account for diffusional barriers
- Can be optimized with existing clinical data for improved predictability ("top-down" approach)
- Clinical data can bridge the gap in transporter IVIVE by generating empirical scaling factors (ESFs)
- Caveats: Interindividual variability, cellular system differences, difference in transporter expression, differences in test system vs tissue activity, allelic variants



Summary



- IVIVE require information from numerous processes and organs
- Each has its advantages and limitations in translating data to the clinic
- Compartmental modeling is the best approach
- Full kinetic characterization of transporters is required
- Knockout animals provide valuable information and can improve the understanding of human PK (with caution)
- Expression and localization of transporters is needed
- More work is required to determine scaling factors



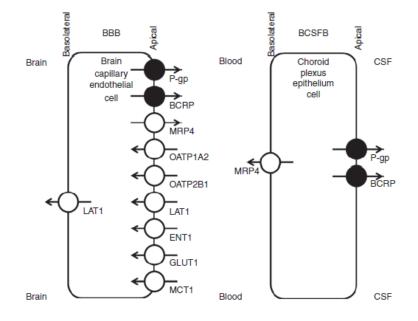
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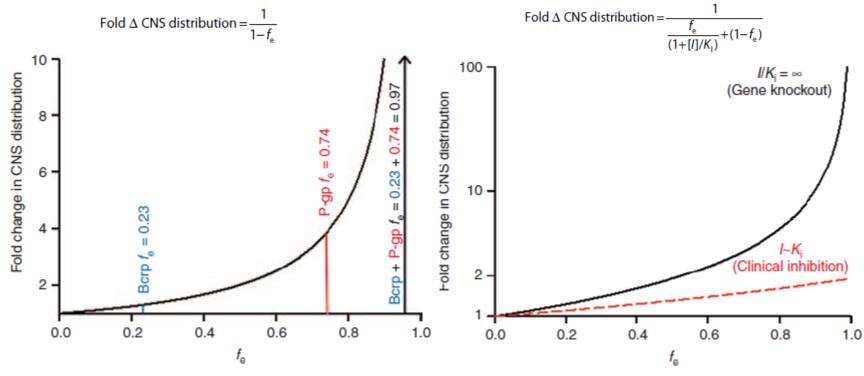


- BBB is tightest endothelium in human body
- ↑ lipophilicity ↑ passive permeability
- Efflux transporters limit CNS penetration
- Transporter function at blood-cerebrospinal fluid (CSF) barrier can function differently than BBB
- Example, P-gp and BCRP secrete
 substrates into CSF,
 but at BBB efflux into
 blood



P-gp/BCRP synergy (lapatinib)





- Knockout mice
 - Bcrp-knockout: $f_e = 0.23$ (1.3-fold)
 - P-gp-knockout: $f_e = 0.74$ (3.8-fold)
 - Both: $f_e = 0.97$ (33-fold)
 - Actual: 26-fold

- Mdr1a/1b/Bcrp^{-/-} mice
 - Genetic ablation
- Clinical inhibition
 - Only weak inhibition is obtained (I = Ki)
 - Cyclosporin Cmax/Ki (P-gp) = 1.6
 - Efavirenz Cmax/Ki (BCRP) = 0.06
 - Largest DDI with lapatinib = 1.9-fold



- Multidrug-resistance proteins (MRPs)
 - Only MRP4 in human and rodent BBB
 - Limits penetration of adefovir and topotecan
 - Mrp4 knockout mice: < 2-fold increase in exposure
 - Risk of DDI is low
- Uptake transporters (LAT1, ENT1, MCT1, Glucose transporter 1, OATPs)
 - Competitive inhibition not likely due to Km >> Cmax
 - No clinical interactions



- CNS exposure is defined by Kp (Brain to plasma ratio)
- ↑ systemic drug conc, ↑ brain exposure
- Systemic DDI ≠ BBB DDI
 brain distribution is not altered
- Models
 - Brain capillary endothelial cells
 - Noncerebral cell lines: MDCK and LLCPK1
 - Animal models (including knockout)
 - PET imagining
 - CSF concentration
 - CNS pharmacodynamic effects



- DDI predicted based on KO animal studies with complete loss of transporter function
- Complete inhibition of transporters at the BBB is not likely
 - Unbound Cmax \leq Ki, P-gp and BCRP inhibition \leq 50% (2-fold increase in exposure)
- No clinical induction of BBB P-gp reported
- No clinical correlation between P-gp function and CNS disorders





Upcoming Events: September 2013

XenoTech will be attending the following meetings:

• 9/17 – IQPC: Clinically Relevant Drug Transporters (London, UK)

Tom Zaleski, Ph.D. tzaleski@xenotechllc.com

 9/30 – 10th International ISSX Meeting (Toronto, Ontario, Canada) Booth # 301





Thank You!

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