

Discovery Research

In-Vitro ADME Capabilities



Discovery Research
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| | |
|-------------------------------|---|
| Absorption | Solubility (Kinetic), Lipophilicity (LogD7.4), Permeability (PAMPA and Caco-2) and Efflux assays (Caco-2) |
| Distribution | Protein binding (plasma, brain homogenate, and microsomes), Blood to plasma partitioning (Species: rat, dog, and human) |
| Metabolism | Metabolic stability and intrinsic Clearance (S9, microsomes, rP450, hepatocytes), Reaction phenotyping (CYP and non-CYP, Enzyme kinetics, Chemical inhibition method and RAF), Metabolite identification and GSH trapping |
| Drug-Drug Interactions | Direct inhibitor (IC_{50} and K_i), Time dependent inhibition (Single point, IC_{50} shift, K_I and K_{inact} , Dialysis) CYP Induction (mRNA expression and enzyme activity) Uptake transporters: substrate and inhibition assays (OAT1, OAT3, OCT1, OCT2, OATP1B1, OATP1B3, MATE-1 and MATE-2K) |



Solubility and Distribution coefficient

Kinetic solubility can help interpret the complication arising from the compound precipitation during biochemical, functional, and cell based assays
Identify poor soluble compounds that reduce productivity in drug discovery and development

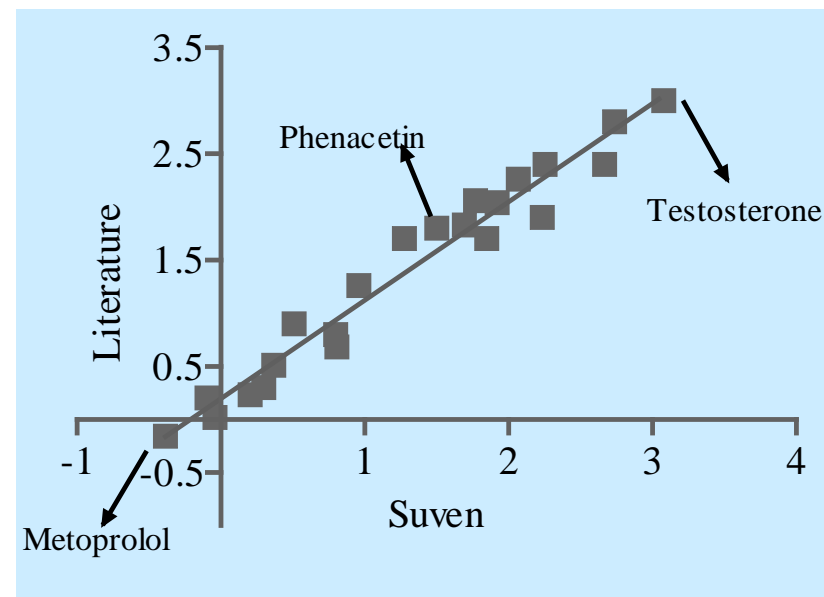
Kinetic solubility

Phosphate buffer 7.4
Simulated gastric fluid
Simulated intestinal fluid

Lipophilicity is a key factor in determining the permeation of physiological membrane, protein binding, and target affinity

Log D 7.4

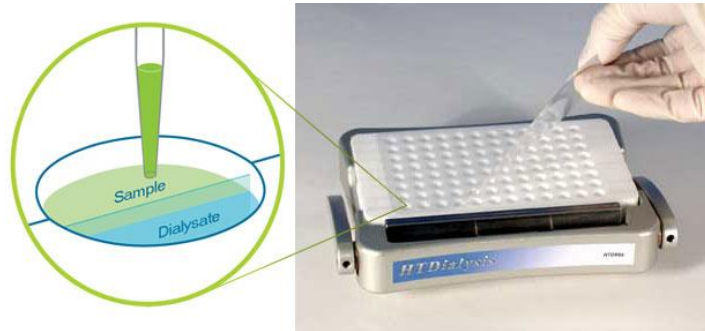
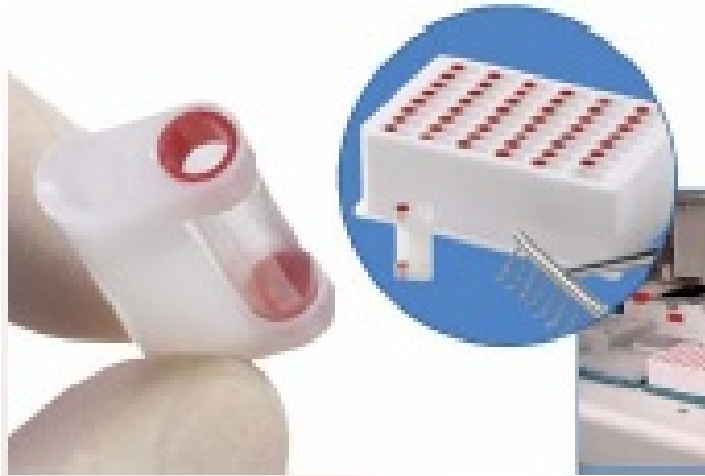
Miniaturized shake-flask method
n-Octanol / Phosphate buffer 7.4
Cyclohexane / Phosphate buffer 7.4



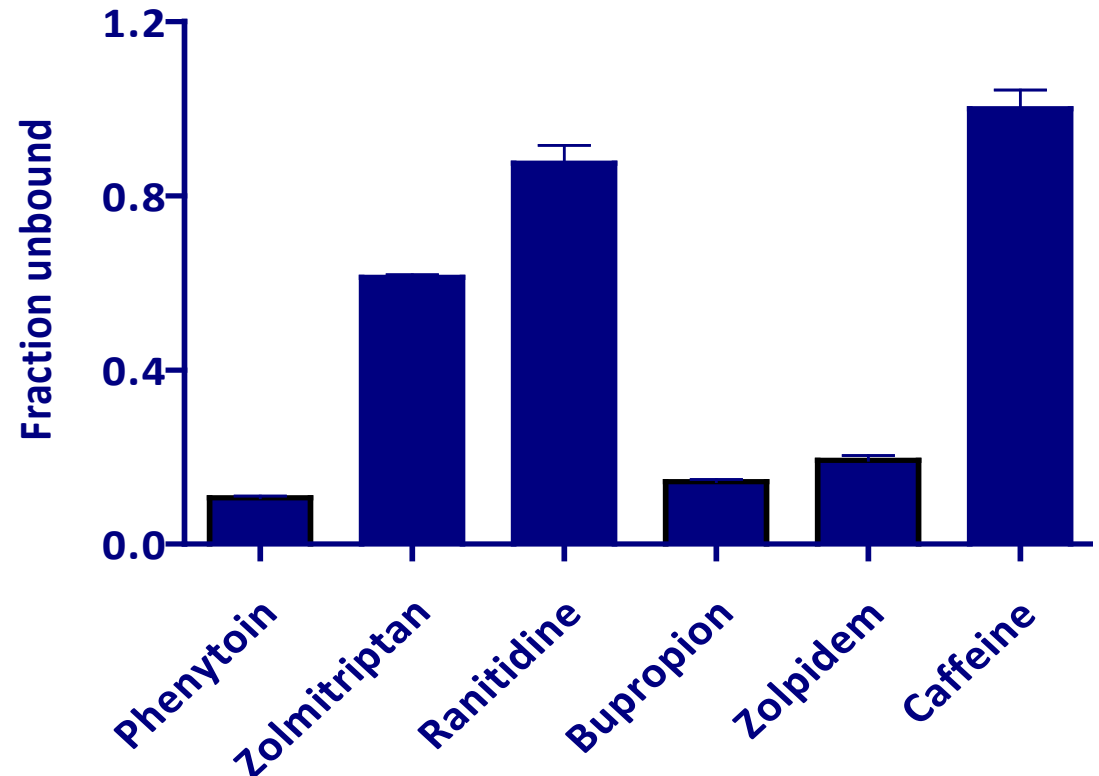


Unbound fraction (plasma/brain homogenate/microsomes)

Knowledge of Unbound fraction is essential to correct the total plasma concentration and correlate with efficacy



Brain homogenate



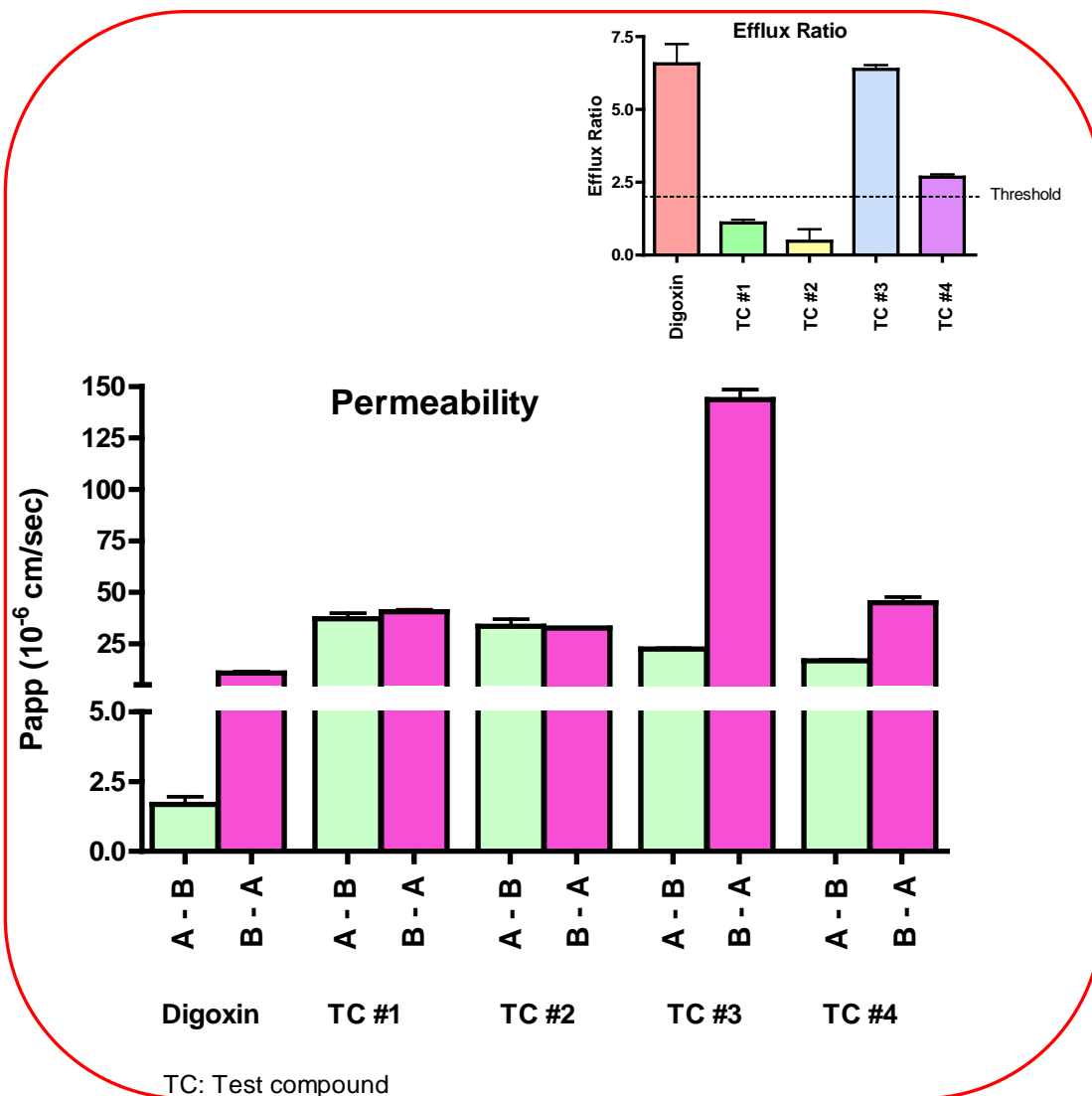
Rapid Equilibrium Dialysis (RED) and High Throughput (HT) Dialysis



P-glycoprotein (P-gp) Substrate assessment

A net flux ratio (or efflux ratio (ER)) of ≥ 2 in cells that express P-gp (e.g., Caco-2 cells) suggest that New chemical entity is a P-gp substrate

| | |
|------------------|-----------------------|
| Test System: | Caco-2 |
| Test Conc.: | 10 μ M |
| Incubation: | 60 min, 37 °C |
| Positive control | Digoxin |
| Membrane | TEER & Lucifer Yellow |
| Analysis | LC-MS/MS |

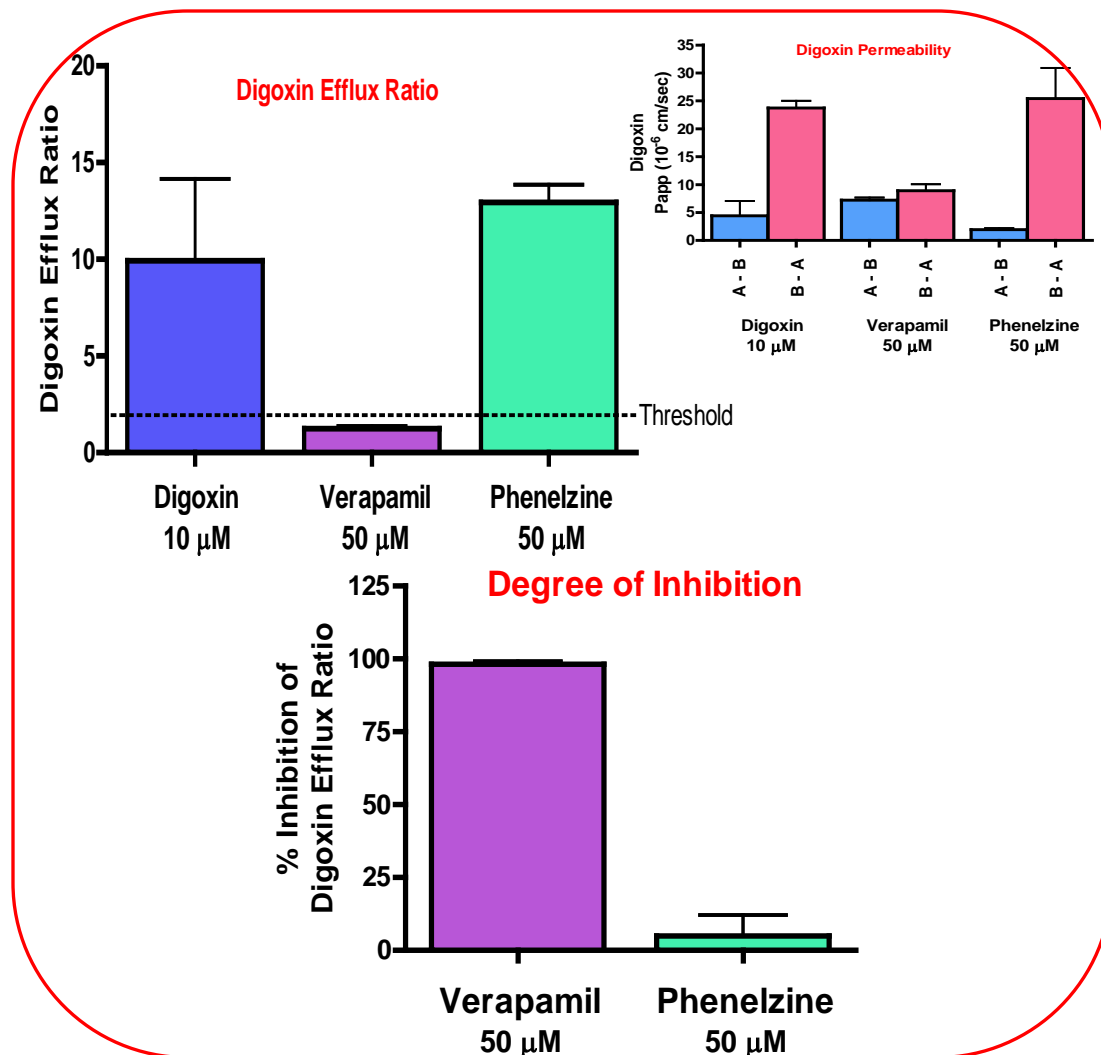


Suven Discovery



P-glycoprotein (P-gp) Inhibitor assessment

| | |
|--------------------|-----------------------|
| Test System | Caco-2 |
| Test Concentration | 50 μ M |
| Positive control | Verapamil |
| Negative Control | Phenelzine |
| P-gp Substrate | Digoxin |
| Incubation | 1 hour |
| Membrane Integrity | TEER & Lucifer Yellow |
| Analysis | LC-MS/MS |





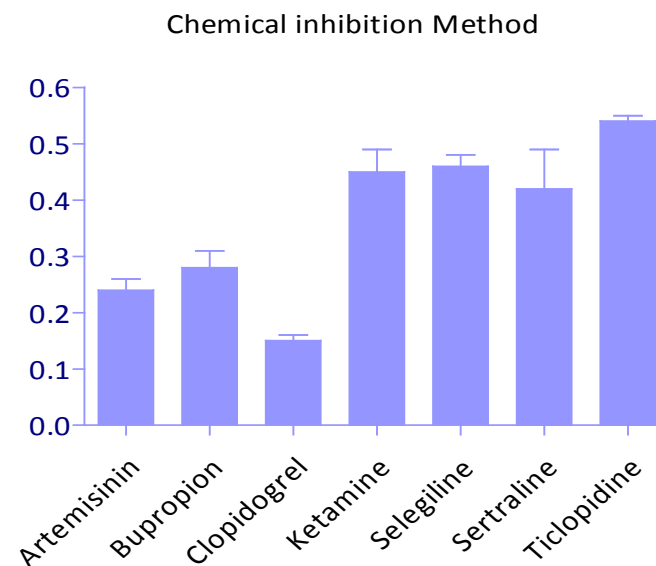
Phenotyping (Chemical inhibition method and RAF)

Contribution $\geq 25\%$ by an enzyme is considered significant based on in vitro phenotyping studies and Human Pharmacokinetic study

Evaluate the role of CYP1A2, CYP2B6, 121 CYP2C8, CYP2C9, CYP2C19, CYP2D6, and CYP3A4/5 and additional enzymes including MAO, FMO, and UGT

In vitro Phenotyping studies include chemical inhibition and Metabolism in recombinant enzymes (RAF approach)

| P450 | HLM CL_{int} ($\mu\text{L}\cdot\text{min}^{-1}\cdot\text{mg}^{-1}$) | rCYP CL_{int} ($\mu\text{L}\cdot\text{min}^{-1}\cdot\text{pmol CYP}^{-1}$) | CL-RAF ($\text{pmol}\cdot\text{mg}^{-1}$) | Mean CYP abundance ($\text{pmol}\cdot\text{mg}^{-1}$) | CL-ISEF |
|------|---|--|--|---|------------------|
| 1A2 | 7 ± 0.3 | 1.2 ± 0.05^a | 5.6 ± 0.5 | 39 | 0.14 ± 0.01 |
| 2A6 | 1298 ± 58 | 23 ± 0.5 | 56.9 ± 3 | 27 | 2.11 ± 0.1 |
| 2B6 | 3 ± 0.2 | 0.13 ± 0.01 | 19.8 ± 1 | 16 | 1.24 ± 0.07 |
| 2C8 | 777 ± 38 | 13 ± 2 | 58.8 ± 6 | 22.4 | 2.62 ± 0.3 |
| 2C9 | 96 ± 6 | 4.3 ± 0.1 | 21.2 ± 1 | 61 | 0.35 ± 0.01 |
| 2C19 | 0.4 ± 0.1 | 0.15 ± 0.01 | 2.8 ± 0.4 | 11 | 0.25 ± 0.04 |
| 2D6 | 33 ± 0.3 | 32 ± 1 | 1.0 ± 0.03 | 12.6 | 0.08 ± 0.002 |
| 2 E1 | 8 ± 0.3 | 0.13 ± 0.001 | 62.0 ± 2 | 64.5 | 0.96 ± 0.03 |
| 3A4 | 399 ± 33 | 18 ± 1.3 | 22 ± 1 | 93 | 0.24 ± 0.01 |





Enzyme Kinetics and CYP Inhibition

CYP Inhibition (IC_{50} and K_i)
 CYP1A2, CYP2B6, 121 CYP2C8, CYP2C9, CYP2C19, CYP2D6, and CYP3A4/5

Basic Model (I/K_i) or Mechanistic Static Model to predict drug interaction risk

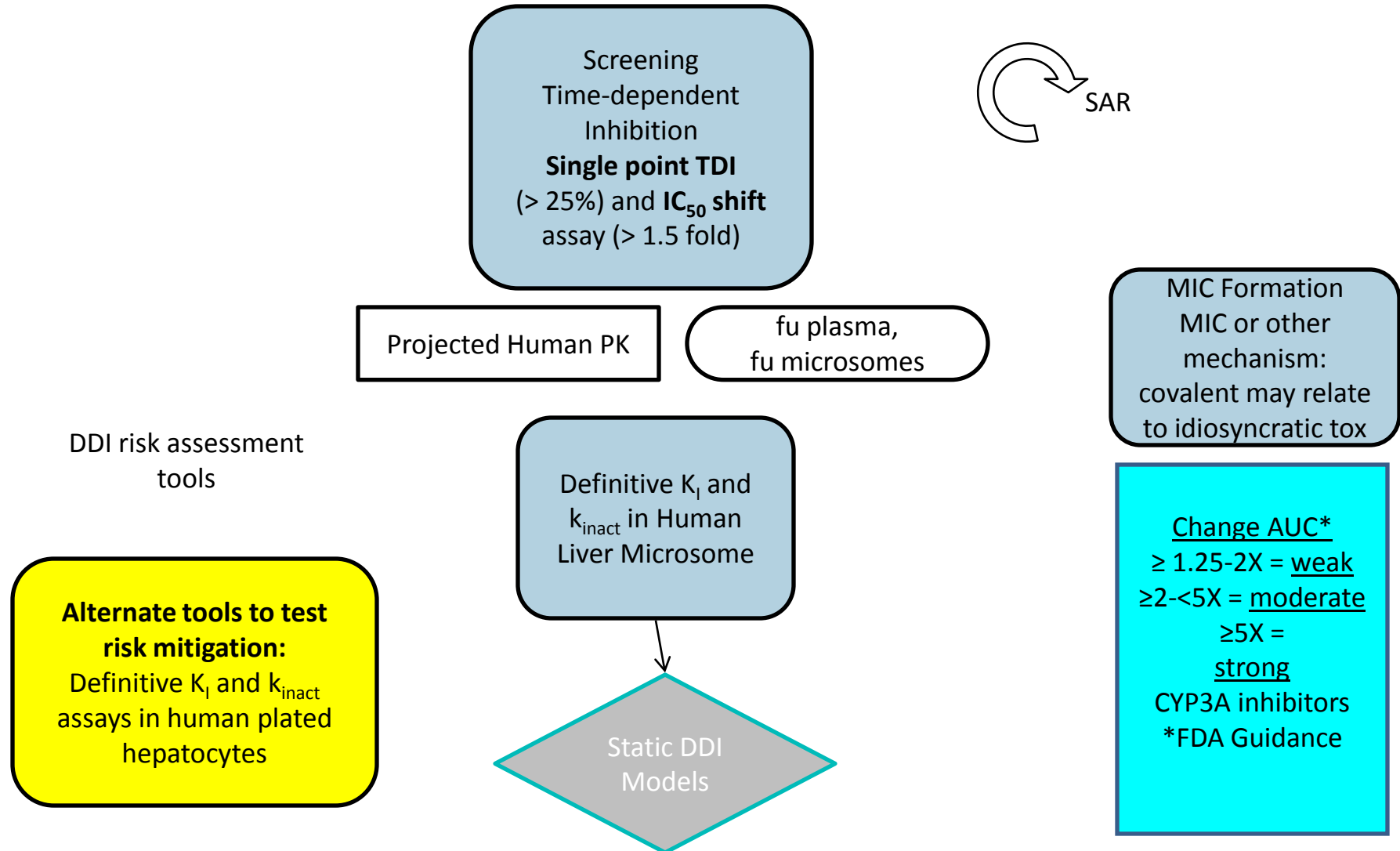
Validated marker activities for major CYP isoforms in HLM and rCYP

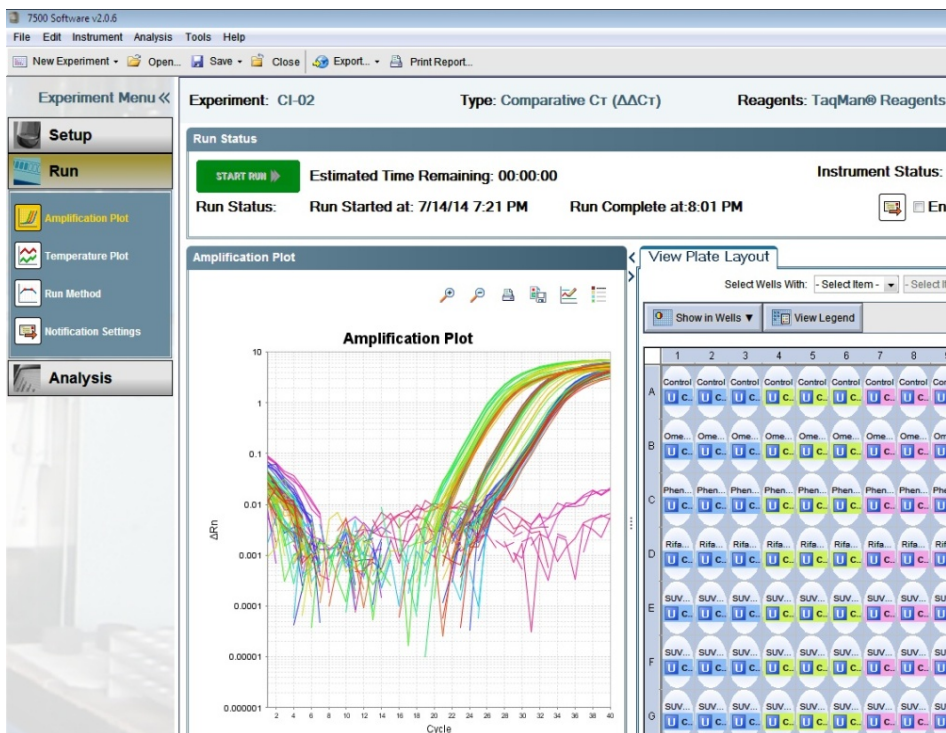
| P450 | HLM | | | rCYP | | |
|------|----------------------|--|----------------------|------------------|--|----------------------|
| | K_m (μM) | V_{max} ($pmol \cdot min^{-1} \cdot mg^{-1}$) | Kinetic Mechanism | K_m μM | V_{max} ($pmol \cdot min^{-1} \cdot pmol CYP^{-1}$) | Kinetic Mechanism |
| 1A2 | 59 ± 3 | 395 ± 10 | MM | 82 ± 5.9^a | 31 ± 2.7 | NC |
| 2A6 | 0.5 ± 0.07 | 649 ± 10 | MM | 0.5 ± 0.004 | 11.4 ± 0.3 | MM |
| 2B6 | 70 ± 3 | 187 ± 7 | SI | 88.4 ± 10.6 | 11.9 ± 0.5 | SI |
| 2C8 | 2.7 ± 0.2 | 2097 ± 51 | MM | 0.5 ± 0.1 | 6.6 ± 0.1 | MM |
| 2C9 | 9.7 ± 0.5 | 933 ± 17 | MM | 1.1 ± 0.04 | 5.0 ± 0.1 | MM |
| 2C19 | 92 ± 8 | 38 ± 2 | MM | 13.4 ± 0.7 | 2.0 ± 0.01 | MM |
| 2D6 | 4.6 ± 0.1 | 152 ± 4 | MM | 0.2 ± 0.1 | 6.3 ± 0.1 | MM |
| 2E1 | 177 ± 9 | 1384 ± 61 | MM | 101.5 ± 2.2 | 12.8 ± 0.1 | MM |
| 3A4 | 3.0 ± 0.2 | 1197 ± 63 | MM | 3.1 ± 0.2 | 56.1 ± 0.9 | SI |

^a- S_{50} instead of K_m ; ^b- Negative co-operativity, clearance read from plot of $v/[S]$ vs $[S]$;

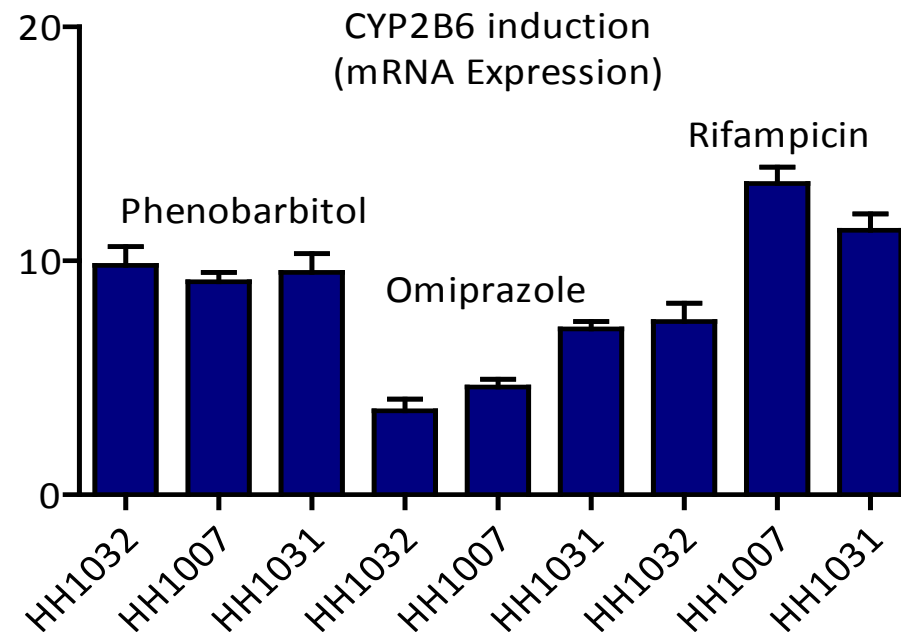


Time Dependent Inhibition





Fold change relative to control
(Mean ± SEM)

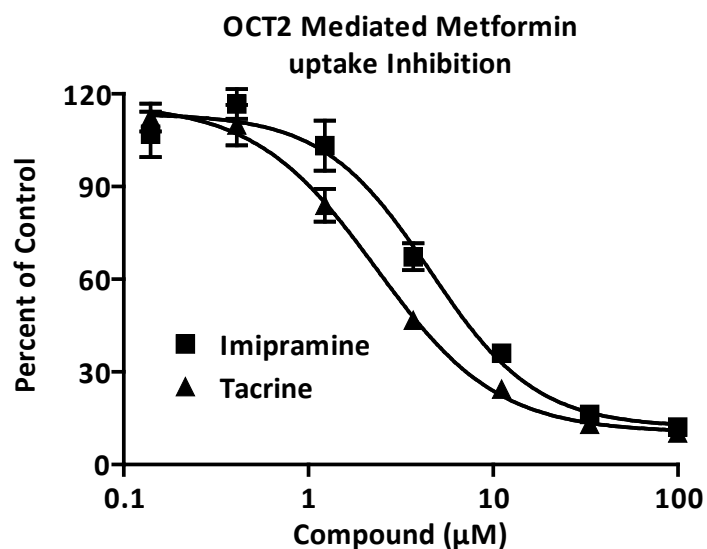


CYP Induction evaluated by “Gold standard method” using cryopreserved plateable hepatocytes from three donors at three concentrations of new chemical entity with both mRNA expression and enzyme activities monitored using RT-PCR and LC-MS/MS respectively. Vehicle control, Positive control, and Negative controls are included in the assay



Uptake Transporter (IC₅₀ assay)

| Transporter | Substrate | Positive control | Absolute IC ₅₀ (uM) |
|-------------|--------------------------------|------------------|--------------------------------|
| OAT1 | Para-amino hippuric acid (PAH) | Flufenamic acid | 0.5 |
| OAT3 | Estrone 3-sulfate (E3S) | Indomethacin | 0.7 |
| OCT1 | Tetraethyl ammonium (TEA) | Verapamil | 5.0 |
| OCT2 | Metformin | Imipramine | 6.4 |
| OATP1B1 | Estradiol β-D Glucuronide | Sulfasalazine | 3.4 |
| OATP1B3 | Estradiol β-D Glucuronide | Rifampicin | 2.3 |



| Compound ID | OCT2 Abs IC ₅₀ (µM) | Literature Reported |
|-------------|--------------------------------|---------------------|
| Imipramine | 6.4 | 3.3 |
| Tacrine | 3.4 | 3.1 |



- Independent Quality Assurance team
- Quality System Procedures (QSP's) for Quality System Management and Standard Operating Procedures (SOP's) for Operation, Calibration, Maintenance of Equipment's
- Document and Data Control, Conducting Internal Audits, Study Specific Audits
- Dedicated Archive facility for the retention of the records
- Facility audited and approved by many global pharmaceutical companies and majority of Indian Pharma Companies

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