

Human In vitro Drug Metabolism Dataset

Transforming scientific data into clinical knowledge

The Drug Metabolism Dataset contains results from *in vitro* metabolism studies, where a drug is tested as an inhibitor/activator/inducer (precipitant) or a substrate (object) for a given human drug metabolizing enzyme (including variants).

- **Metabolism parameters** (IC_{50} , K_i , K_l , k_{inact} , % inhibition, % activation, % or fold increase, EC_{50} , $E_{max'}$, K_m , $V_{max'}$, and CL_{int}), along with detailed experimental conditions, are extracted from published articles (citations) and NDA/BLA reviews.
- Study results are organized according to the overall effect and mechanism of the interaction:
 - → Enzyme inhibition entry: drug as inhibitor or non-inhibitor
 - → Enzyme activation entry: drug as activator or non-activator
 - → Enzyme induction entry: drug as inducer, down-regulator, or non-inducer
 - → Enzyme substrate entry: drug as substrate
- Multiple queries allow users to retrieve an *in vitro* dataset by drug name, enzyme name, or mechanism of the interaction (drug as precipitant or as object).
- **Results** can be viewed, customized, and downloaded in multiple formats, allowing users to compile and organize the large body of information available.



FROM A CITATION OR NDA/BLA REVIEW

The latest, most relevant, peer-reviewed publications and regulatory documents are identified and fully analyzed. Study protocol and results are manually curated to update the knowledgebase on a daily basis.



Abstract

Many dietary supplements are promoted to patients with osteoarthritis (OA) including the three naturally derived compounds, glucosamine, chondroitin and diacerein. Despite their wide spread use, research on interaction of these antiarthritic compounds with human hepatic cytochrome P450 (CYP) enzymes is limited. This study aimed to examine the modulatory effects of these compounds on CYP2C9, a major CYP isoform, using in vitro biochemical assay and in silico models. Utilizing valsartan hydroxylase assay as probe, all forms of glucosamine and chondroitin exhibited IC50 values beyond 1000 µM, indicating very weak potential in inhibiting CYP2C9. In silico docking postulated no interaction with CYP2C9 for chondroitin and weak bonding for glucosamine. On the other hand, diacerein exhibited mixed-type inhibition with IC₅₀ value of 32.23 μM and K_i value of 30.80 μM, indicating moderately weak inhibition. Diacerein's main metabolite, rhein, demonstrated the same mode of inhibition as diacerein but stronger potency, with IC₅₀ of 6.08 μM and K_i of 1.16 µM. The docking of both compounds acquired lower CDOCKER interaction energy values, with interactions dominated by hydrogen and hydrophobic bondings. The ranking with respect to inhibition potency for the investigated compounds was generally the same in both in vitro enzyme assay and in silico modeling with order of potency being diacerein/rhein > various glucosamine/chondroitin forms. In vitro-in vivo extrapolation of inhibition kinetics (using 1 + [I]/K_i ratio) demonstrated negligible potential of diacerein to cause interaction in vivo, whereas rhein was predicted to cause in vivo interaction, suggesting potential interaction risk with the CYP2C9 drug substrates.

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TO A FULLY CURATED DATASET

Prior to integration, all data are carefully and critically evaluated. The richness of each citation, including relevant insights, is exploited, generating a highly detailed dataset.

Tree View of Citation Data 2 PubMed 29488228 & Comments: All the tested salt forms of glucosamine and chondroitin did not show inhibition on CYP2C9-mediated valsartan 4-hydrylation at concentrations up to 1000 µM in vitro. Object: valsartan Cardiovascular Drugs -- Angiotensin II Inhibitors (Angiotensin Receptor Blockers or ARBs) Mechanism: valsartan as a substrate of CYP2C9 Overall Effect: In Vitro Metabolism System: Microsomes (recombinant) entry Test system and Object Metabolite: 4-hydroxyvalsartan Enzymes: CYP2C9 test concentrations ■ Object Concentration: 10-1000 µM **Enzyme substrate** Study Results: 🔗 Study results: K_m, V_{max}, - K_m: 146.29 ± 53.62 μM and CL_{int} values - V_{max}: 43.04 ± 4.90 pmol/min/mg - CL_{int}: 0.2942 μL/min/mg Experimental conditions: Incubation: Volume = 200 μL Time = 30 min Other experimental details Protein concentration: 0.5 mg/mL Cofactors: NADPHgensys Precipitant: diacerein Treatments of Pain and Inflammation - Anti-inflammatory Drugs Mechanism: diacerein as an Object: valsartan Cardiovascular Drugs — Angiotensin II Inhibitors (Angiotensin Receptor Blockers or ARBs) inhibitor of CYP2C9-mediated Overall Effect: In Vitro Enzyme Inhibition valsartan 4-hydroxylation System: Microsomes (recombinant) Object Metabolite: 4-hydroxyvalsartan Enzymes: CYP2C9 Enzyme inhibition entry Object Concentration: 0.5-2Km Precipitant Concentration: 7.5-60 μΜ Study Results: 6 Study results: K, value (K_i: 30.80 μM) determination - Regression and mode of inhibition Inhibition type: Mixed Experimental conditions: - Incubation: Volume = 200 μL Time = 30 min Protein concentration: 0.5 mg/mL Cofactors: NADPHgensys Object Concentration: 146 µM Precipitant Concentration: 0-500 µM (estimated from Fig. 5) Study Results: 6 -□ IC₅₀: 32.23 μM Study results: IC₅₀ value * Experimental conditions: Diacerein inhibits CYP2C9 with an IC $_{so}$ value of 32.23 μM and a K $_{i}$ value of 30.80 $\mu M.$ What other drugs inhibit the CYP2C9 index substrate diclofenac?



POWERFUL TOOL FOR **DATA INTEGRATION**: FROM ONE CITATION TO METADATA ANALYSIS

The data are formatted for immediate use and can be filtered and re-arranged to allow metaanalysis of multiple results.

Query all CYP2C9 inhibitors

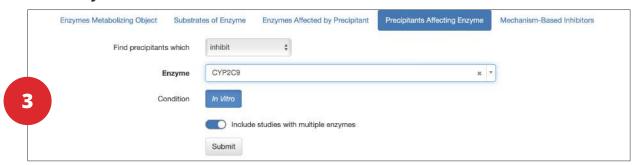


Table View of Query Results Filter Multiple formats for viewing and downloading Showing 201 to 300 of 1,078 entries (filtered from 3,910 total entries) 🔗 👌 Advanced Table Search Select columns Copy Excel CSV Print Precipitant Object Inhibition Accession # or Therapeutic Precipitant Class Object | Metabolite System Ki IC50 NDA/BLA# Published || Type diclo nifedipine Cardiovascular diclofenac Microsomes 0.57 µM Competitive PubMed 16963489 2006 Dec Drugs hydroxydiclofenac (recombinant) Calcium Channel Blockers triclabendazole sulfoxide Anti-Infective diclofenac Microsomes Ki,u = 0.565 Competitive NDA 208711 ± 0.03 µM; hydroxydiclofenac Antiparasitics Ki = 1.33 ± 0.063 µM honokial Natural Products diclofenac Microsomes 0.54 µM Competitive PubMed 24005963 2013 Sep 03 hydroxydiclofenac HL (pooled) Herbal Medications Cardiovascular PubMed 10064574 fluvastatin (acid) diclofenac 0.5 ± 0.1 Mixed Microsomes Drugs hydroxydiclofenac Mu HMG CoA Reductase Inhibitors (Statins) Natural Products galangin diclofenac Recombinant 0.50 µM Competitive PubMed 19074529 2009 Mar hydroxydiclofenac enzyme Herbal Medications Natural Products PubMed 26750984 diclofenac 0.50 µM 2016 Oct Microsomes Competitive glycyrol hydroxydiclofenac HL (pooled) Herbal Medications Anti-Infective diclofenac PubMed 11207028 2001 Feb Competitive Agents _ hydroxydiclofenac (recombinant) Antibiotics alpha-naphthoflavone Miscellaneous diclofenac Microsomes 0.41 uM Competitive PubMed 16963489 2006 Dec hydroxydiclofenac Agents -(recombinant) Obtain a complete list of in vitro inhibitors of CYP2C9 Copyright © 2023 Certara, USA All Rights Reserved.



IN VITRO METABOLISM DATASET IN NUMBERS

(as of October 16, 2023)

6,206

404

NDAs/BLAs

24,269 substrate

51,831

inhibition

735 activation

8,905

inductior entries

69,926

14,517

positive entries

negative entries

36 *in vitro* metabolism queries

with **117** possible searches

186 drug metabolizing isozymes

& **72** variants

2,764

5,306

compounds / compound as substrates / as inhibitor

265

1,975

compounds

compounds as inducers

705 food products & 1,960 herbal medications

APPLICATIONS OF THE IN VITRO METABOLISM DATASET



PROVIDES CONTEXT for RESULTS OBTAINED for candidate compouds



ALLOWS ASSESSMENT of MEASUREMENT VARIABILITY

(inter-lab, substrate- and system-dependency, etc.)



SUPPORTS STATIC
PREDICTIONS and PBPK
MODELING

with input parameters



HELPS OPTIMIZE IN VITRO STUDY DESIGN

(cell system, incubation conditions, test concentrations, choice of substrate/inhibitor, etc.)



ASSISTS with DOSE SELECTION for clinical trials



PROVIDES IN VITRO EVIDENCE to EXPLAIN CLINICAL RESULTS and improve understanding of drug interaction mechanisms



To learn more, visit www.druginteractionsolutions.org or email DIDBase@Certara.com



About Certara

Certara accelerates medicines using proprietary biosimulation software, technology and services to transform traditional drug discovery and development. Its clients include more than 2,000 biopharmaceutical companies, academic institutions, and regulatory agencies across 62 countries.