

## **Modeling Chemical Fate and Metabolism for Computational Toxicology**

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#### **Environmental Issue**

The need to develop a scientifically credible approach for setting risk-based priorities for chemical testing requirements prior to having empirical data from which to estimate risks.

Computational Toxicology is the integration of modern computing and information technology with the technology of molecular biology and computational chemistry to improve EPA's prioritization of data requirements and risk assessments for toxic chemicals

#### **Objectives of ERD-Athens CompTox Team**

#### I. Modeling Chemical Fate in Ecosystems

- Perform laboratory and field experiments to fill data gaps Utilize input from genomic tools to:
- determine "threshold concentration" for toxic event - identify the toxic chemical species (in a mixture)
- Utilize computational tools for fate model parameterization and speciation (e.g., SPARC - SPARC Performs Automated Reasoning in Chemistry)

#### **II. Modeling Chemical Fate in Organisms (metabolism)**

- Evaluate existing metabolic simulators
- Improve simulator database (e.g., add new pathways)
- Apply advanced analytical tools for testing / improving metabolic simulator (e.g., NMR, Raman)
- Utilize SPARC outputs to inform metabolic simulator of species of concern (hydrolysis, speciation, etc.)

#### **III. Software Engineering**

- Design and implement a modeling framework (e.g., FRAMES)
- Perform uncertainty and sensitivity analysis

#### **The Problem**

Environmental fate and metabolic pathways are often complex even for relatively simple chemicals (e.g., bromobenzene). Pathways must be elucidated and prioritized for environmental fate and metabolic simulators to be developed.



in the process of developing and refining simulators for environmental transformation and metabolism.





# **Partnering to Protect Human Health and the Environment**

### **Scientific Approach**

Experimental systems and tools for populating databases for environmental fate and metabolic simulators

- Experimentally measured values. Systems include: -Well characterized sediments and soils -Intact liver -Liver microsomes
  - -S9 fraction
- Rate constants and molecular descriptors derived from mechanistic-based SPARC and/or OSAR models
- Advanced spectroscopic techniques (e.g., wide-bore NMR) for measuring metabolic rate constants and identifying metabolites in vivo and in vitro

