

# Maximizing the Utility of Existing Data: Opportunities and Challenges for the Application of Systematic Review in Cross Species Extrapolation

Sara Vliet

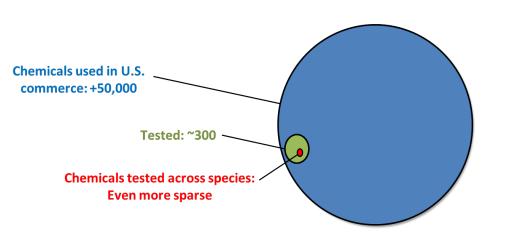
Biologist, Scientific Computing and Data Curation Division



The views expressed in this presentation are those of the author and do not necessarily reflect the views or policies of the US EPA

#### **New Approach Methodologies: Moving Away From Animal Testing**

 The mission of the EPA is to Protect Human Health and the Environment and the evaluation of ecological and human health risks associated with toxic chemicals represents one of the EPA's primary responsibilities



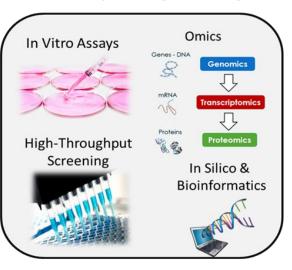
#### **Whole-Animal Models**

→ Observe Toxic Outcome



#### **New Approach Methods**

→ Identify Biological Target



#### NAMs come with their own set of challenges (just to name a few)...

- Uncertainties surrounding biological relevance (metabolism, kinetics, etc.)
- Uncertainties surrounding biological representation (cell-type, species, etc.)
- Uncertainties surrounding biological extrapolation (how well does the molecular/cellular data reflect organism responses?)

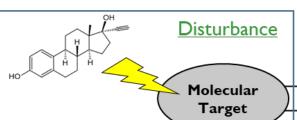


**EFFICIENCY** 

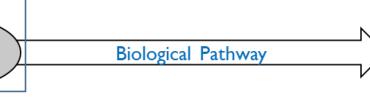


#### **Cross Species Extrapolation: A Challenge in New Approach Methodologies**

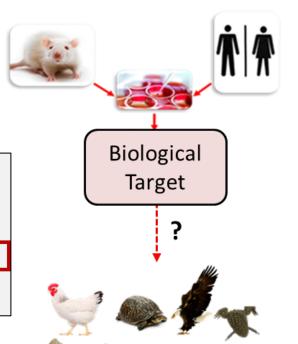
- Within animal testing, the sensitivity of species to a chemical is assumed to be a function of their relatedness, however, how well a model species represents the species of concern is often poorly understood
- This question still applies to many NAMs and the extrapolation of NAMs data from test species (e.g., mammalian cells) to other species of concern is essential



• For NAMs to play useful roles in decision-making, we need to understand how changes at the molecular level in cells and tissues are related to the apical adverse outcomes



Observed
Toxic Effect



#### In the lab

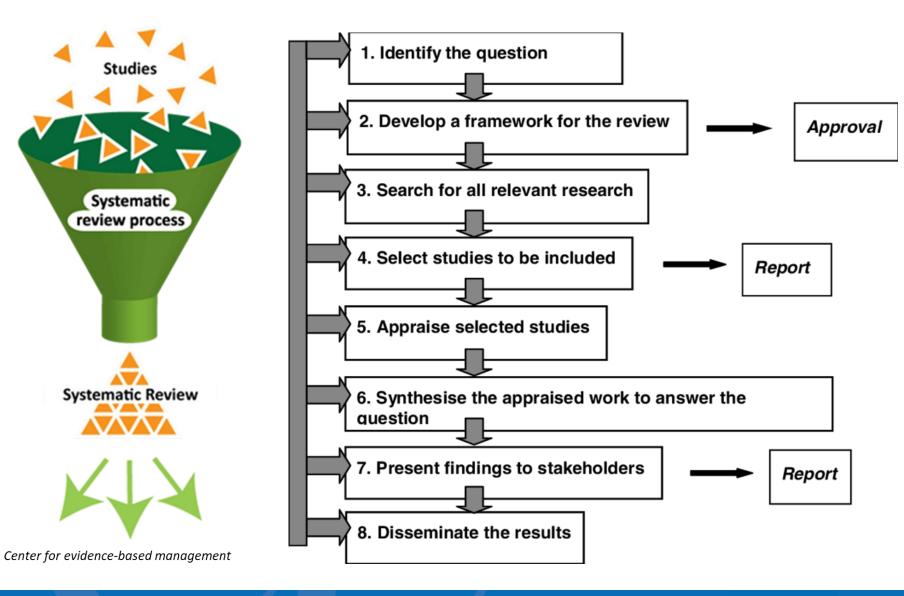
- Cross species In vitro studies
- Cross species In vivo studies
- Chemical proteomics
- Etc.

#### Out of the lab

- Homology modelling
- Molecular docking
- Review of existing evidence
- Etc.

Systematic Review: Literature review that uses systematic methods to collect secondary data, critically appraise research studies, and synthesize findings.

- First developed and used heavily within evidencebased medicine and recently applied more broadly to environmental health and cross-species applications
- Treats the searching, collection, and synthesis of existing evidence as a controlled experiment



#### **Systematic Review in Ecotoxicology**

## Opportunities & Challenges

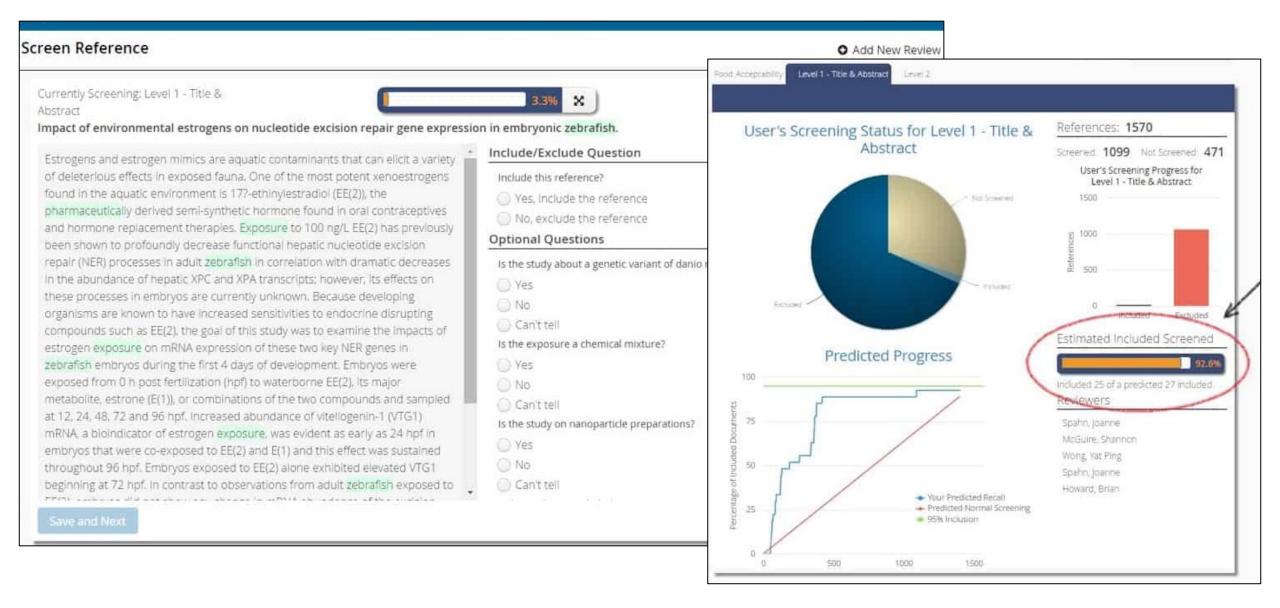


 As methods have evolved, a variety of tools have emerged to help manage the systematic review process and help address these challenges

- The use of existing data reduces the need for further animal testing
- The use of systematic methods promote increased transparency, objectivity, consistency, and reproducibility of information
- Strengthened confidence in risk assessments
- Reduces the influence of reviewer bias and error in the evaluation process
- Harmonized data collection
  - What data fields are being collected during the review?
  - What type of language is being used to describe this data?
- Standardized Techniques
  - What methods are being used to search the literature?
  - What type of QC/QA is being conducted?
- Need for flexibility
  - Ecotoxicology is diverse, approaches will need to be similarly flexible
  - SR may not always be necessary, what are you trying to achieve?
- Resource Requirements
  - Cost, time, expertise, training, etc.



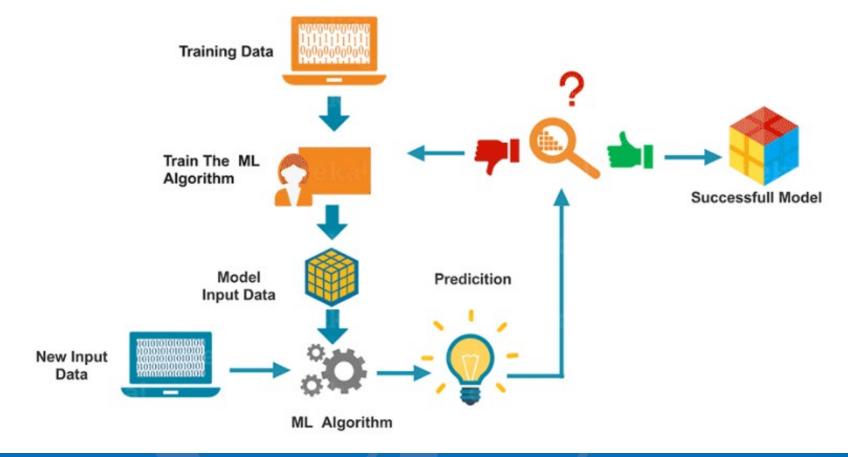
#### **Tools of the Trade: Systematic Review Software**



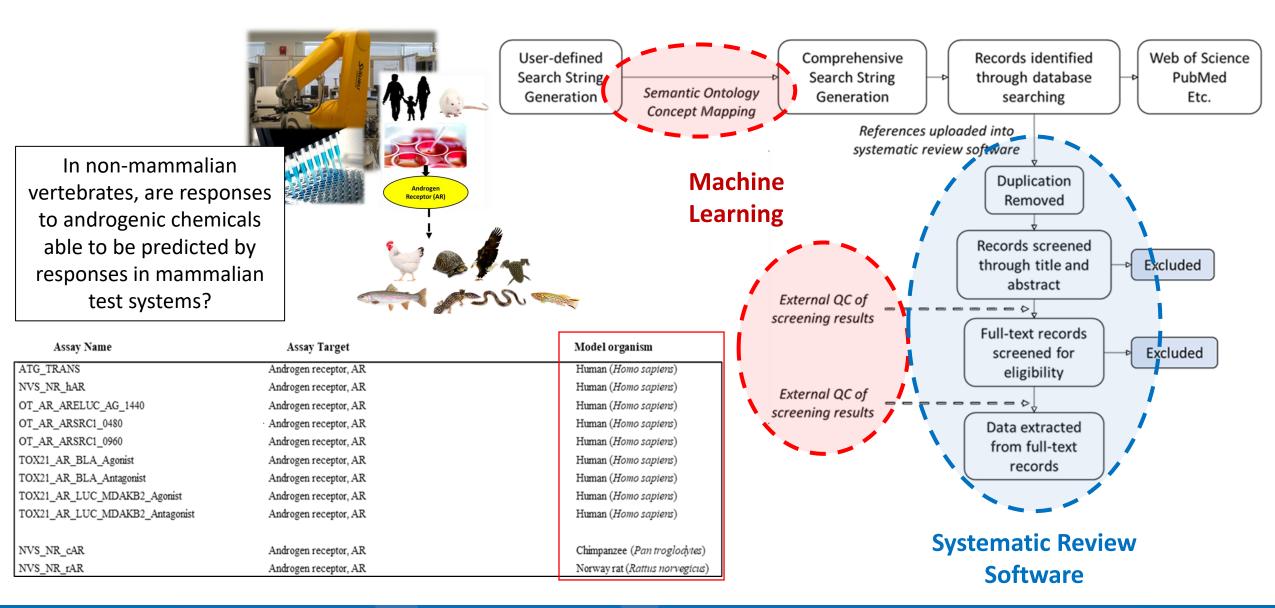
#### **Tools of the Trade: Machine Learning**

#### Machine learning can help reduce the human screening burden by predicting and prioritizing relevant references

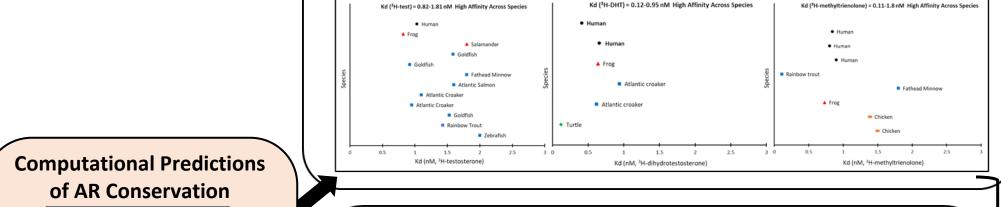
- Algorithms are also being explored as a potential "second reviewer" or to completely replace human screening
- Applications are being developed to extract information from scientific text, tables, and figures
- Different data/reporting formats and lack of standardization proves challenging
- In general, these methods can be time savers, but all have some limitations, and do not eliminate the need for expert input



#### Case Example: How can we apply Systematic Review Methods to Cross-Species Extrapolation?



#### Case Example: How can we apply Systematic Review Methods to Cross-Species Extrapolation?



**US EPA SeqAPASS Tool** 

**Systematic Evaluation of In Vivo Cross-Species Data** 

Systematic Evaluation of In Vitro Cross-Species Data

Observations by Vertebrate Class Chemical Bird Reptile · Female development of · Female development of Female development of male sex traits male sex traits Testosterone (endogenous androgen) · Male-biased populations Male-biased populations (Skewed sex-ratios) (Skewed sex-ratios) Female development of Female development of male sex traits Methyltestosterone Reduced gonadosomatic · Male-biased populations (synthetic androgen) Male-biased populations Reduced egg laying in index (GSI) (Skewed sex-ratios) (Skewed sex-ratios) Female development of Female development of · Female development of Female development of 17ß-trenbolone environmental androgen Reduced circulating E2 · Male-biased populations · Male-biased populations · Male-biased populations and vitellogenin levels (Skewed sex-ratios) (Skewed sex-ratios) (Skewed sex-ratios)

Weight of Evidence for AR Conservation Across Species

- Apply pathway to other targets of interest
- Repeat process to account for new information
- Inform future computational predictions



### Thank you!

Vliet.Sara@epa.gov

