

# A quantitative structure-activity relationship (QSAR) model to estimate serum half-lives of per- and polyfluoroalkyl substances (PFAS) in multiple species

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# Trying to characterize the toxicokinetics of PFAS

#### What are per- and polyfluoroalkyl substances (PFAS)?

- Highly diverse group of anthropogenic molecules that substitute fluorine for hydrogen along carbon backbones
- 1223 PFAS are currently in the TSCA inventory
- 602 PFAS are known to be commercially active
- Widely distributed in environmental substrates
- Generally non-biodegradable
- Often have relatively long half-lives in body tissues

#### What do we want to do?

- Describe and quantify toxicokinetics (TK) of PFAS
- Extrapolate TK parameters from data-rich to data-limited substances
- Prioritize chemicals for in-depth assessment based on predicted half-life and predicted exposure

#### What's the problem?

- TK parameters vary widely and non-allometrically across species
- Some TK parameters vary by sex within species
- Read-across methods and cross-species extrapolations are unreliable

#### A potential solution: Machine Learning

- Use data from in vivo testing of PFAS across multiple species and chemicals
- Integrate multiple chemical and physiological characteristics
- Produce fit-for-purpose predictions of TK parameters
- Contribute to prioritization of chemicals for more in-depth evaluation

# Putting it together: datasets and model assembly

#### In vivo serum half-life (SHL) data

- Curated dataset of values published in peer-reviewed literature 1-20
- 66 data points
  - 11 PFAS chemicals
  - 4 species (Human, Monkey, Rat, Mouse), sex-specific values when available

## Model predictors: capturing PFAS chemical/structural diversity

- Considered 118 predictors in 6 general categories
- **❖** Physio-chemical characteristics (OPERA)<sup>21,22</sup> **❖** Protein binding affinities<sup>27,28</sup>
- **❖** Species-specific kidney characteristics<sup>23,24,25</sup> **❖** Critical micelle concentrations<sup>29</sup>
- **❖** Similarity to endogenous compounds<sup>26</sup>
- Inclusion of ether bond

#### Predictors pruned in an iterative process

- Identified pairs of predictors with Spearman's  $\rho > 0.9$
- Retained member of pair with better correlation with SHL
- 29 predictors remaining for model construction

#### Machine learning method: Random Forest (RF)

- Investigated two approaches with *caret* 30,31 package of R
  - RF Classification model with SHL binned into 3 bins
    - Fast (<1 day), Moderate (1 day 1 year), and Slow (>1 year)
  - RF Regression Model
- Both models fit with 10-fold cross validation, replicated 10 times

# **Evaluation of machine learning models**

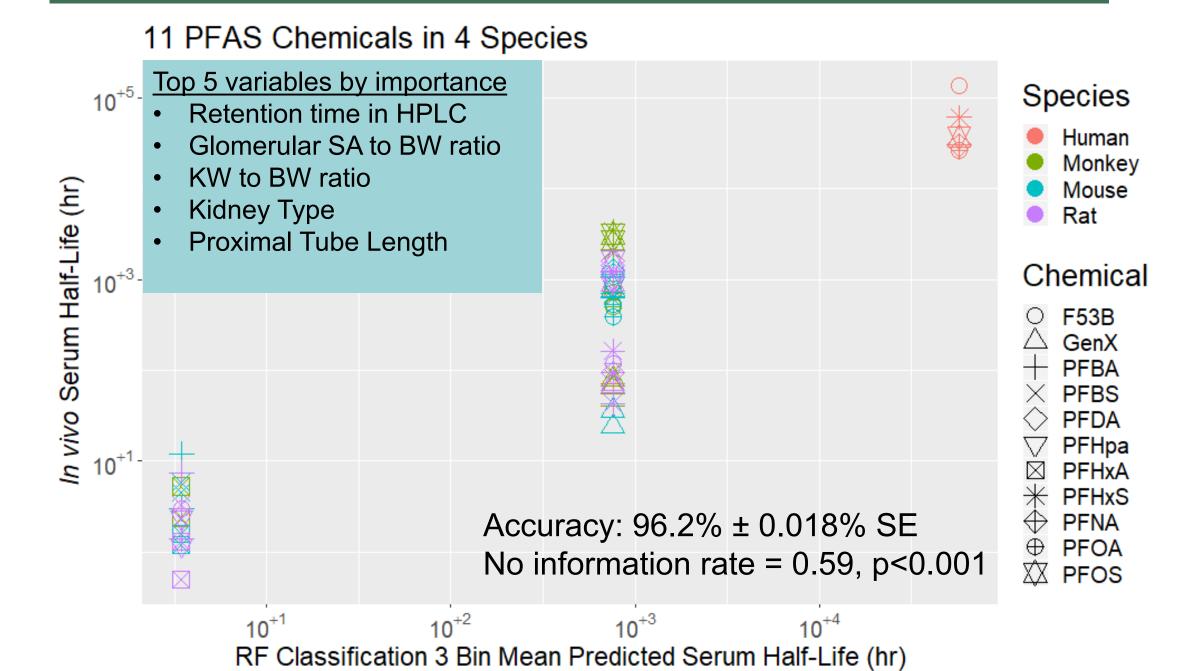


Figure 1. RF classification model results. Predicted (fast, moderate, or slow) vs in vivo SHL, shown by chemical and species.

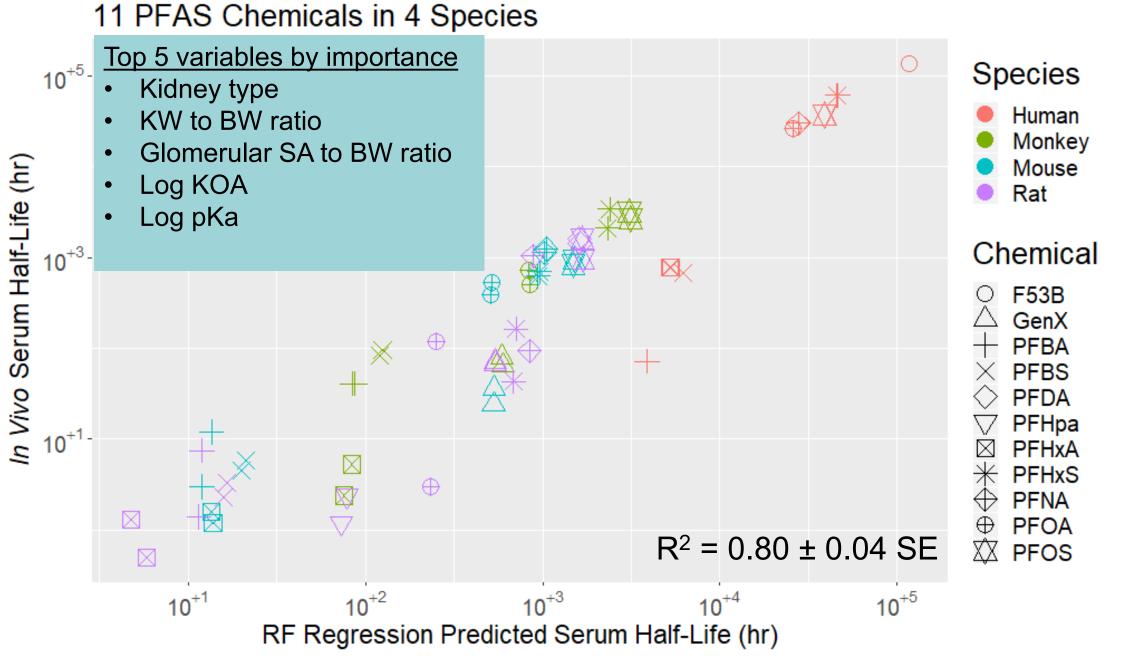


Figure 2. RF regression model results. Predicted vs in vivo SHL, shown by chemical and species.

# **Application to the DSSTox PFAS dataset**

Classification model used to predict SHL for 6648 PFAS and PFAS-like chemicals listed in the USEPA Distributed Structure-Searchable Toxicity (DSSTox) database

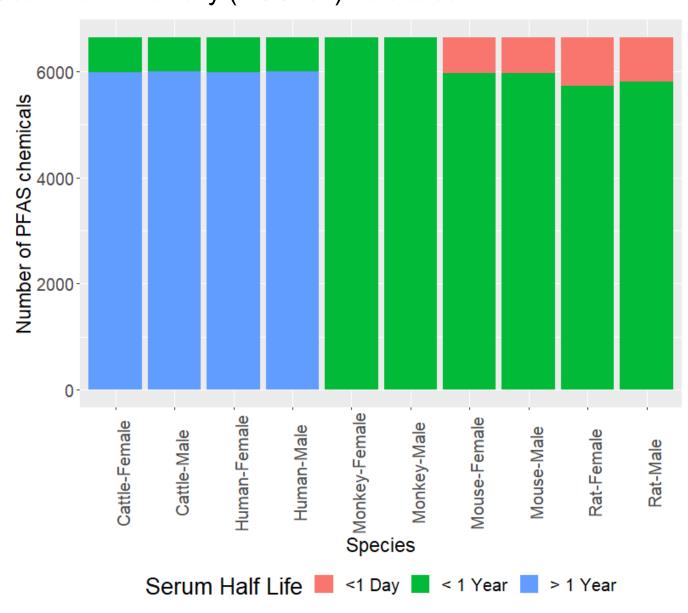


Figure 3. Classification of 6648 DSSTox chemicals by species (5) and sex into fast, moderate, or slow SHL categories. Based on kidney characteristics<sup>23</sup>, model can potentially be extended to other species.

## **Conclusions and future directions**

- Both classification (Fig 1) and regression (Fig 2) models are promising initial efforts to estimate an important TK parameter of PFAS chemicals using machine learning
- Though less precise, the classification model is highly accurate
- Application of classification model to DSSTox database suggests most PFAS may have long SHLs (> 1 year) in humans (Fig 3)
- Limitations
- Models are likely overfit (29 parameters with 66 obs)
- Regression model tends to overestimate length of SHL for quickly clearing chemicals (Fig 2)
  - Additional training data for such chemicals needed
- Lack of independent test set due to data paucity

#### Future work

- Expansion of models with additional SHL data
- Chemicals
- Species
- Development of evaluation sets of in vitro and in vivo data
- Incorporation of predictions into TK modeling effort