

# PKMine; AI for extracting input parameters from different sources to support PBK modelling

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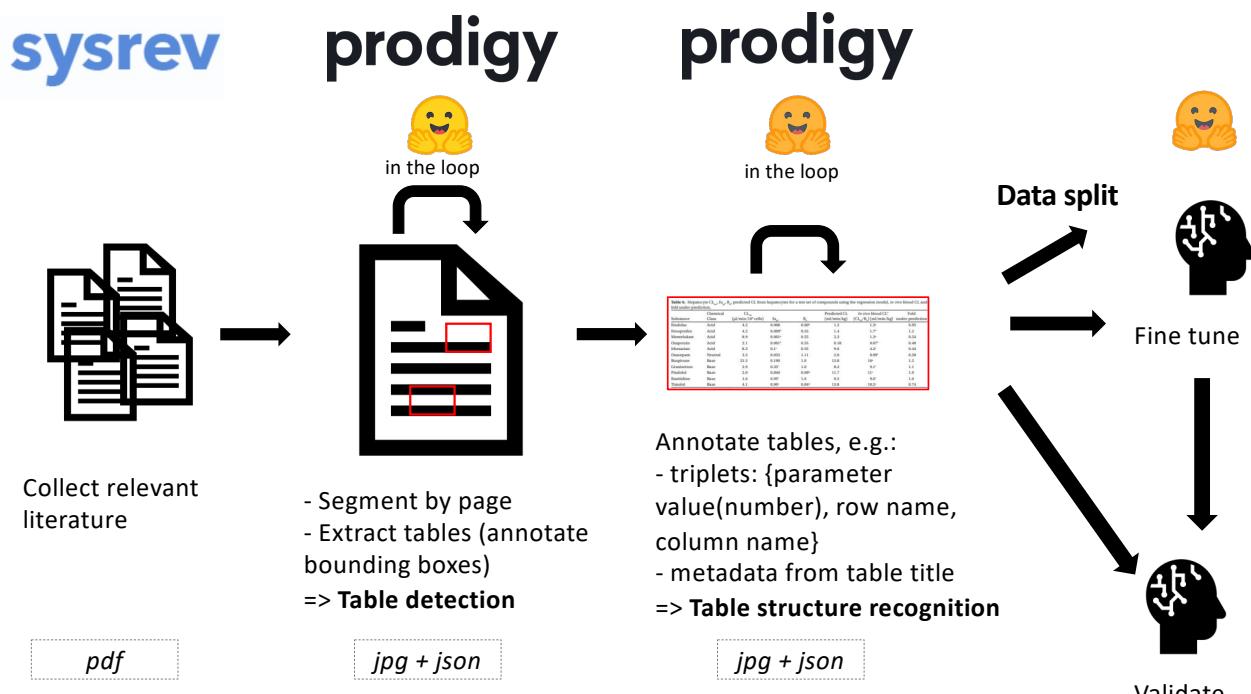
## Aim

Design a workflow to collect PBPK model parameters and validation data in a structured way. We propose to finetune AI models to extract these data from text, figures, tables and databases. Extracted data will include:

- Physicochemical properties
- *In vitro* ADME parameters
- *In vivo* ADME parameters

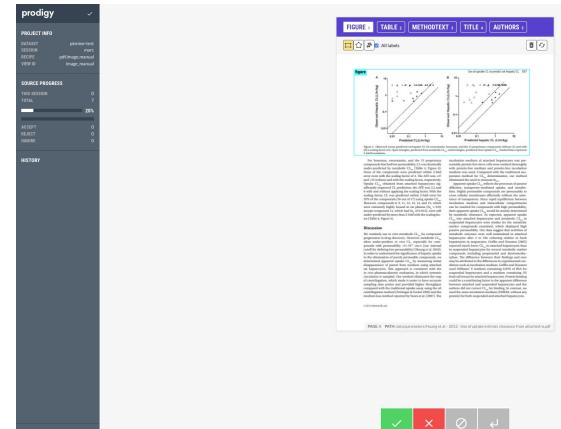
## Output

Parameters in a format that maximises interoperability with PBK tools and software (e.g. PKSim, R).



## Getting training data

Use annotation software **Prodigy** ([explosion.ai](https://explosion.ai)) on PDF files collected in the paper: Louisse, J. et al (2020). Towards harmonization of test methods for *in vitro* hepatic clearance studies. *Toxicology in vitro*; 63, 104722.



## Candidate models

### TableTransformerModel

[https://huggingface.co/docs/transformers/model\\_doc/table-transformer](https://huggingface.co/docs/transformers/model_doc/table-transformer)

class `transformers.TableTransformerModel`

### Abbreviations

ADME: Absorption, Distribution, Metabolism, Excretion

PBPK: Physiologically Based Pharmacokinetic modelling

TrOCR: Transformer Optical Character Recognition

### TrOCRProcessor

[https://huggingface.co/docs/\\_doc/trocr](https://huggingface.co/docs/_doc/trocr)

class `transformers.TrOCRProcessor`