

Eawag Swiss Federal Institute of Aquatic Science and Technology



Uploading, Sharing, and Extracting Biotransformation Data in enviPath

May 11th 2025





Synthesizing data on environmental contaminant biotransformations requires community-driven efforts







There are two ways you can upload your data to enviPath











The enviPath website – good for uploading small datasets







Getting started with the enviPath website



ENVIPORTAL THE ENVIRONMENTAL CONTAMINANT BIOTRANSFORMATION PATHWAY RESOURCE

enviPath is a database and prediction system for the microbial biotransformation of organic environmental contaminants. The database provides the possibility to store and view experimentally observed biotransformation pathways. The pathway prediction system provides different relative reasoning models to predict likely biotransformation pathways and products. You can try it out below.

Learn more >>

2





Add a pathway starting with the parent compound



Test Package 1	F Actions -
Test package for SETAC Workshop	13
Pathways	
Rules	
Compounds	
Reactions	
telative Reasoning	
Scenarios	





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Contact

enviPath UG (haftungsbeschränkt





Add transformation products (TPs) to the pathway







Connect reactants (educts) with products







Add experimental information to the pathway using scenarios



Create a new empty scenario



Add experimental information

Text from paper

"Fresh **aerobic** activated sludge (**volatile suspended solids** 3.8 g/L) was obtained from the aeration basin at the Palo Alto WWTP in **Palo Alto, California**"





Search Info -

Add experimental information to the pathway using scenarios

Test Package 1						
Test package for SETAC Workshop						
Pathways						
Rules						
Compounds						
Reactions						
Relative Reasoning						
	Relative Reasoning Scenarios	Relative Reasoning Scenarios	Relative Reasoning Scenarios	Relative Reasoning Scenarios	Relative Reasoning Scenarios	Relative Reasoning Scenarios





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Impressum/Imprint

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Actions -





Add experimental information to the pathway using scenarios



Add more experimental information

Text from paper

"The observed first-order **rate constant** for the disappearance of N-EtFOSE calculated from the linear regression in Figure 2 (R^2 = 0.967), is k_{obs} = 0.99 ± 0.08 d⁻¹, corresponding to a **half-life** of 0.71 ± 0.06 d (Table 2)."





Search

📕 Actions 🗸

Remove

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Info 🚽

Add experimental information to the pathway using scenarios

2)	Package Pathway Rule Compound Reaction Relative Reasoning Scenario Home / Package / Test Package 1 / Scenario / Rhoads et al 2008 N-EtFOSE - (00000)	Setting l
Scenario entries	Rhoads et al 2008 N-EtFOSE - (00000)	
Rate constant Lower = 0.91 d^{-1} Upper = 1.07 d^{-1}	N-EtFOSE pathway in activated sludge Type: Sludge	
fe r = 0.65 d	Property Location	Value Palo A
967	Redox condition	aerobio
	Volatile suspended solids concentration (VSS)	Start: End:





SCHOOL OF COMPUTER SCIENCE SCHOOL OF COMPUTER SCIENCE MACHINE LEARNING

Unit

g/L

Delete All:

Impressum/Imprint

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3





Add experimental information to the pathway using scenarios

Attach the scenario to a compound in the pathway

Package	Pathway	Rule	Compound	Reaction	Relative Reasoning	Scenario	Setting	User	Group	Predict -		S	earch	Info 🚽
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	N-EtFOSE	E Path	way											
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The Final Pathway





3		Μ	lake	package public		
Pathway	Rule	Compound	Reaction	Predict -	Search	Info
					_	

Home / Package / Test Package 1

Test Package 1	✓ <u>Actions</u> -
Test package for SETAC Workshop	Update Q Ask for review
Pathways	License
Rules	·III Publish
Compounds	 Import scenarios
Reactions	Download
Relative Reasoning	Export as JSON
Scenarios	 Engineer Package Get Missing Rules Info
	C Update References





Adding data for large datasets can get tedious... Add your data to the empty Create an empty Make package public package package Compounds Pathways Reactions "We built our random-forest model using the Scenarios enviPath-PFAS package in addition to our own experimental data" ^{⊲' shall} keep∎ hy data "Our dataset is publicly available on the enviPath platform"





We developed a BiotransformAtion Reporting Tool (BART) to help upload pathways

Find it on GitHub

https://github.com/FennerLabs/BART





- Spreadsheet-based
- Compatible with enviPath Python
- Easy to edit and reuse



Why do we need this tool?

There is a large amount of experimental data available on chemical contaminant biotransformations in the environment, but most of this data is stored in an inaccessible, non machine-readable format (e.g. behind paywalls, in PDFs). We aim to fix this by encouraging researchers to share their data in an open source, standardized format that can be easily used to upload the data into open-source, freely available online software (e.g. <u>enviPath</u>).

About the template

This repository contains an empty version of the BiotransformAtion Reporting Tool (BART), and a filled-out version with data from the publication (<u>10.1021/acs.est.3c05506</u>) as an example.

Empty template: https://github.com/FennerLabs/BART/blob/main/BART_AUTHOR_YEAR_PathwayName_Template.xlsx

Filled-out Example: https://github.com/FennerLabs/BART/blob/main/BART_FANG_2024_6-2FTNO_Example.xlsx





You can download and empty version and a filled-out example at the GitHub repository

ピ main 👻 ピ 1 Branch 📀 2 Tags	Q Go to file
🚳 slr257 Update README.md 📼 🛛 Emp	ty Template
BART_AUTHOR_YEAR_PathwayName_Template	Add files via upload
BART_FANG_2024_6-2FTNO_Example.xlsx	Add files via upload
	Initial commit
	Opuate README.md

There is a FAQ page in case you get stuck

\mathbf{O}	FennerLab	s / BART								Q Type // to se	arch
Code	 Issues 	11 Pull requests	₽ Discussions	Actions	Projects	🕮 Wiki	Security	🗠 Insights	鐐 Sett	ings	
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BART also simplifies uploading large, complicated pathways



Figure 4. Proposed transformation pathways of 6:2 FTNO and 6:2 FTSA in aerobic sludge.





Filling out BART – Start with the Description Tab







Add SMILES for compounds in Compounds tab

	А	В	(C D E
	1 Type	Name	SMILES	
	2 Parent	6:2 FTNO	C[N+](C)(CCCNS(=O)(=O)CCC(C(C(C(C(C(F)(F)F)(F)F)(F)F)(F)F)(F	
	3 TP	6:2 FTAA	FC(F)(C(C(C(C(CCS(=O)(=O)NCCCN(C)C)(F)F)(F)F)(F)F)(F)F)(F)F)F)F	
		6:2 FTUAA	FC(F)(C(C(C(C(C(CCS(=O)(=O)NCC/C=[N+](\C)/C)(F)F)(F)F)(F)F)(F)F)F)F)F	
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Farent compo	Junu	6:2 FTPrAm	FC(F)(C(C(C(C(CCS(=O)(=O)NCCCN)(F)F)(F)F)(F)F)(F)F)(F)F)F	
is clearly lab	blod	6:2 FTSAm-PrOH	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NCCCO)(=O)=O)F	
is clearly labe	eleu	6:2 FTSAm-PrAL	FC(F)(C(C(C(C(CCS(=O)(=O)NCCC=O)(F)F)(F)F)(F)F)(F)F)(F)F)	
		6:2 FTSAm-PrA	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NCCC(O)=O)(=O)=O)F	
	10 TP	N-Me-6:2 FTSAm-PrA	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(N(C)CCC(O)=O)(=O)=O)F	
	11 TP	N-Et-6:2 FTSAm	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NCC)(=O)=O)F	
	12 TP	6:2 FTSAm-EtOH	FC(C(F)(C(F)(C(F)(C(F)(CCS(NCCO)(=O)=O)F)F)F)F)F)(F)F	
	13 TP	6:2 FTSAm-AcAL	FC(C(F)(C(F)(C(F)(C(F)(CCS(NCC=O)(=O)=O)F)F)F)F)F)(F)F	
	14 TP	6:2 FTSAm-AcA	FC(C(F)(C(F)(C(F)(C(F)(CCS(NCC(O)=O)(=O)=O)F)F)F)F)F)F)F)F	
	15 TP	6:2 FTSAm-ForAL	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NC=O)(=O)=O)F	
	16 TP	6:2 FTSAm	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(N)(=O)=O)F	SIVILES for all compounds in a
	17 TP	6:2 FTSAm-Conjunct-1	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NC(C(N)C(C)CC)=O)(=O)=O)F	
	18 TP	6:2 FTSAm-Conjunct-2	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NC(C(N)CCC)=O)(=O)=O)F	pathway should be provided in one
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Connect Reactants and Products in the Connectivity Tab

	Α	В	С	D	E	F	G 🔺
	1 Reactant	Reactant_SMILES	Product	Product_SMILES	Multistep	Comment	1
	2 6:2 FTNO	C[N+](C)(CCCNS(=O)(=O)CCC(C(C(C(C(C(F)(F)F)(F)F)(F)F)(F)F)(F	6:2 FTAA	FC(F)(C(C(C(C(CCS(=O)(=O)NCCCN(C)C)(F)F)(F)F)(F)F)(F)F)(F)F)			
	3 6:2 FTNO	C[N+](C)(CCCNS(=O)(=O)CCC(C(C(C(C(C(F)(F)F)(F)F)(F)F)(F)F)(F	6:2 FTUAA	FC(F)(C(C(C(C(CCS(=O)(=O)NCC/C=[N+](\C)/C)(F)F)(F)F)(F)F)(F)F)(F)F)F			
	4 6:2 FTNO	C[N+](C)(CCCNS(=O)(=O)CCC(C(C(C(C(C(F)(F)F)(F)F)(F)F)(F)F)(F	6:2 FTSAm-PrOH	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NCCCO)(=O)=O)F			
	5 6.2 FTUAA	<u>EC(F)(C(C(C(C(C(C(C(C(C(C(C(C(C(C(C(C(C(C</u>	6:2 FTSeAm	FC(F)(C(C(C(C(CCS(=O)(=O)NCCCNC)(F)F)(F)F)(F)F)(F)F)(F)F)F			
• • •		CCS(=O)(=O)NCCCN(C)C)(F)F)(F)F)(F)F)(F)F)(F)F)F	6:2 FTSeAm	FC(F)(C(C(C(C(CCS(=O)(=O)NCCCNC)(F)F)(F)F)(F)F)(F)F)(F)F)F			
Columns II	n blue are		6:2 FTPrAm	FC(F)(C(C(C(C(CCS(=O)(=O)NCCCN)(F)F)(F)F)(F)F)(F)F)(F)F)(F)F)			
dow	un hoxoc	CCS(=0)(=0)NCCCN)(F)F)(F)F)(F)F)(F)F)(F)F)F	6:2 FTSAm-PrOH	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NCCCO)(=O)=O)F			
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	11 6:2 FTSAm-PrA	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NCCC(O)=O)(=O)=O)F	6:2 FTSAm	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(N)(=O)=O)F			
	12 6:2 FTSAm-PrA	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NCCC(O)=O)(=O)=O)F	N-Et-6:2 FTSAm	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NCC)(=O)=O)F	Reac	tions car	be flagged
	13 N-Et-6:2 FTSAm	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NCC)(=0)=0)F	6:2 FTSAm-EtOH	FC(C(F)(C(F)(C(F)(C(F)(CCS(NCCO)(=O)=O)F)F)F)F)F)F)F)F)	noud		, so naggoa
	14 6:2 FISAm-EtOH	F(C(F)(C(F)(C(F)(C(F)(C(F)(C(F)(C(F)(C(6:2 FISAm-ACAL	F(C(F)(C(F)(C(F)(C(F)(C(F)(C(F)(C(S(NCC(O)-O)(-O)-O)(-F)))))))	2	as multiste	ep here
	15 6:2 FTSAM-ACAL	F(C(F)(C(F)(C(F)(C(F)(C(F)(C(F)(C(G)(G(C(G))-G)(F)(F)(F)(F)(F)(F))))))	6:2 FTSAM-ForAl	F(C(r)(C(r)(C(r)(C(r)(C(r)(C(r)(C(0)-0)-0)-0)-0)-0)-0)=0			
	17 6:2 FTSAm-ForAl	FC(F)(C(F)(F)(F)C(F)(F)(F)C(F)(F)(F)C(F)(F)(F)(F)C(F)(F)(F)(F)(F)(F)(F)(F)(F)(F)(F)(F)(F)(6:2 FTSAm	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(S(N)(=0)=0)F			-
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	21 6:2 FTSAm	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(N)(=O)=O)F	6:2 FTSAm-ForAL	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NC=O)(=O)=O)F			
	22 6:2 FTSAm-Conjunct-1	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(NC(C(N)C(C)CC)=0)(=0)=0)F	6:2 FTSAm	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(N)(=O)=O)F			
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Fill-out Scenario information in a tabular format







Add kinetics, confidence levels, and proposed intermediates last

Image: Control of contro	A	B		E	F G	H	1	Confidence Level DEAC	K Confidence Level Color	L Duran e d'ante mar e d'ante 2	M
Bet R. 201	Scenario	Compound	SMILES Model	Parameter	Value Value Unit	Corrected	R2 0.0076	Confidence Level_PFAS	Confidence Level_Schy	Proposed Intermediate?	Comment
0 0.00000 0.00000	Fang, B. et al. 2024	6:2 FINO	C[N+](C)(C)(C)(S)(=0)(=0)(C)(C)(C)(C)(C)(F)(F)(F)(F)(F)(F)(F)(F)(F)(F)(F)(F)(F)	Hall-life	1.2 d		0.9970	1		1	
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	Fang, B. et al, 2024	N-Me-6:2 FTSAm-PrA	FC(F)(C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)CCS(N(C)CCC(O)=O)(=O)=O)F					38	1	3	
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What should you do with the finished BART?

- 1. Include BART in the Supporting Information of your manuscript at submission
- 2. Send your BART files to the enviPath team (<u>stephanie.rich@uzh.ch</u>) and we can generate the package for you in 1-2 days
- 3. Upload your pathways using our command-line executable (this will be available at the GitHub repo soon)







Extracting data from enviPath currently requires knowledge of python, and we are working to make this more user-friendly

Please visit the tutorial website to try out enviPath Python







Acknowledgements









CNVi**Path**





stephanie.rich@uzh.ch





Questions?





Hands-On Example





Either follow along with our example, or try it on a pathway of your choice

Steps:

- 1. Find a paper with a biotransformation pathway
- 2. Go to BART and download and empty template