

Supporting Materials for
Reversible transformations of sulfamethoxazole and its submoieties by manganese-oxidizing bacteria and biogenic manganese oxides in the presence of humic substances

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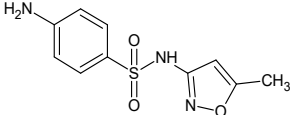
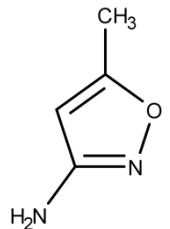
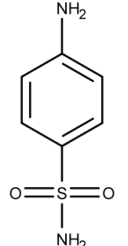
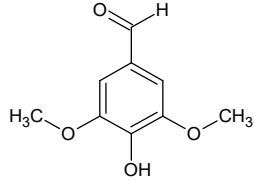
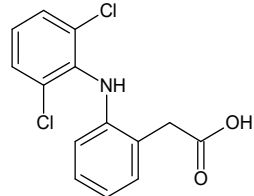
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ESI1 Substances used in the experiment as target contaminants and electron mediator (syringaldehyde) (Source: Kim et al., 2023; except electrophilicity index and ionisation potential calculated as described in Section 2.7 of the main text)

Property	Shortened name and abbreviation used in this study				
	Sulfamethoxazole (SMX)	Isoxazole (ISX)	Sulfanilamide (SNM)	Syringaldehyde (SAH)	Diclofenac (DCF)
IUPAC name	4-amino-N-(5-methyl-1,2-oxazol-3-yl)benzenesulfonamide	5-methyl-1,2-oxazol-3-amine	4-aminobenzenesulfonamide	4-hydroxy-3,5-dimethoxybenzaldehyde	2-[2-(2,6-dichloroanilino)phenyl]acetic acid
Molecular Formula	C ₁₀ H ₁₁ N ₃ O ₃ S	C ₄ H ₆ N ₂ O	C ₆ H ₈ N ₂ O ₂ S	C ₉ H ₁₀ O ₄	C ₁₄ H ₁₁ Cl ₂ NO ₂
Molecular Structure					
CAS no.	723-46-6	1072-67-9	63-74-1	134-96-3	15307-86-5
pKa	pKa1 = 1.6; pKa2 = 5.7	2.5	10.6	7.30	4.15
log K _{ow}	0.89	0.10	-0.62	1.07	4.51
Molecular weight	253.28	98.10	172.21	182.17	296.1
Electrophilicity index (eV)	pH4.5	2.813	2.195	2.565	4.361
	pH 7.5	2.238	2.184	2.561	3.857
Ionisation potential (eV)	pH4.5	6.172	6.711	6.184	6.394
	pH 7.5	5.673	6.700	6.178	5.878

ESI2 Substances used in the experiment as micropollutants or electron mediator (syringaldehyde). All substances were purchased from Sigma Aldrich (Czech Republic)

	Substance	Purity	Concentration, mg/L
Redox mediator	Syringaldehyde	98%	Added in a molar 2:1 or 20:1, syringaldehyde:target substance
Positive control	Diclofenac, DCF	Sodium salt, $\geq 98\%$	10
Sulfonamides	Isoxazole, ISX (3-Amino-5-methylisoxazole)	$\geq 97\%$	10
	Sulfadiazine, SDZ	99.0-101.0%	10
	Sulfadimethoxine, SDM	98.0-102.0%	2
	Sulfamerazine, SMR	$\geq 99\%$	10
	Sulfamethazine, SMZ	99.0-101.0%	10
	Sulfamethizole, SMI	$\geq 99\%$	10
	Sulfamethoxazole, SMX	$\geq 98\%$	10
	Sulfanilamide, SFN	$\geq 98\%$	10
	Sulfapyridine, SPD	$\geq 99\%$	10
	Sulfathiazole, STZ	99.0-101.0%	10

ESI3 Components of the growth medium for *Pseudomonas putida* MnB6

All reagents were used as received without further purification. Yeast extract was purchased from Oxoid; casamino acids from Difco™; HEPES and $\text{CoCl}_2 \cdot 6 \text{H}_2\text{O}$ from VWR Chemicals; D-glucose, CaCl_2 , MgSO_4 , FeCl_3 and $\text{CoCl}_2 \cdot 6 \text{H}_2\text{O}$ from Lach-Ner; $\text{CuSO}_4 \cdot 5 \text{H}_2\text{O}$ from Penta; and $\text{ZnSO}_4 \cdot 7 \text{H}_2\text{O}$ and MnCl_2 from Sigma-Aldrich.

Compound	Supplier	CAS Number	Dosage per 1 L
Yeast extract	Oxoid		0.5 g
Casamino acids	Difco™		0.5 g
HEPES	VWR Chemicals	7365-45-9	5mM
D-Glucose	Lach-Ner	50-99-7	10 mM
CaCl_2	Lach-Ner	10043-52-4	0,48 mM
MgSO_4	Lach-Ner	7487-88-9	0,83 mM
FeCl_3	Lach-Ner	7705-08-0	3.7 μM
$\text{CuSO}_4 \cdot 5 \text{H}_2\text{O}$	Penta	7758-99-8	10 mg
$\text{ZnSO}_4 \cdot 7 \text{H}_2\text{O}$	Sigma-Aldrich	7446-20-0	44 mg
$\text{CoCl}_2 \cdot 6 \text{H}_2\text{O}$	Lach-Ner	7791-13-1	20 mg
$\text{Na}_2\text{MoO}_4 \cdot 2 \text{H}_2\text{O}$	VWR Chemicals	10102-40-6	13 mg
MnCl_2	Sigma-Aldrich	7773-01-5	29 mg

ESI4 Additional information on the analysis of BioMnOx

The BioMnOx obtained after 24-h cultivation was pretreated by 0.17% sodium hypochlorite (12% Cl, stabilized, technical, Carl Roth, Germany) in several iterative steps to eliminate the adsorbed organic fraction originating from the cultivation media. The cleaning procedure was based on the protocol of Villalobos et al. (2003). Scanning electron microscopy (SEM) of the samples was performed using a MIRA 3 electron microscope (Tescan Orsay Holding, Brno, Czech Republic) with a secondary electron detector operated at 12 kV acceleration voltage. The energy-dispersive X-ray spectroscopy (EDX) of the samples was conducted using an energy-dispersive spectroscopy system (Bruker XFlash X-ray detector, Karlsruhe, Germany, and ESPRIT 2 software). The photoelectron spectrum was measured by X-ray photoelectron spectroscopy (XPS) method (Kratos ESCA 3400, X-ray source MgK α 120 W). A wide scan was performed using 0-1100 eV, 0.2 eV/step, 5x acquisitions, and then the identified elements were measured in high resolution (0.05 eV/step, 15x-30x acquisitions). Additionally, the Mn 3s region was separately measured to determine the oxidation state of manganese in the supplied sample (as received) and after Ar⁺ ion sputtering. The C 1s 284.8 eV lines were used as calibration for the evaluation. The specific surface area was determined from the adsorption isotherm of N₂ at T = 77 K using the method of BET surface analysis (Coutler SA 3100 automatic analyzer, Beckman Coutler, Inc., Brea, CA, USA).

The X-ray diffraction (XRD) analysis was performed using a desktop diffractometer Bruker D2 Phaser with an LYNXEYE XE detector (CuK α radiation, 30 kV, 10 mA, and measuring increment step of 0.022° 2 θ , time step 2.5 s, in the range from 5° to 80° 2 θ). The identification of all phases was performed using Diffrac. Suite EVA software (version 4.3) and the ICDD PDF-2 database (2018).

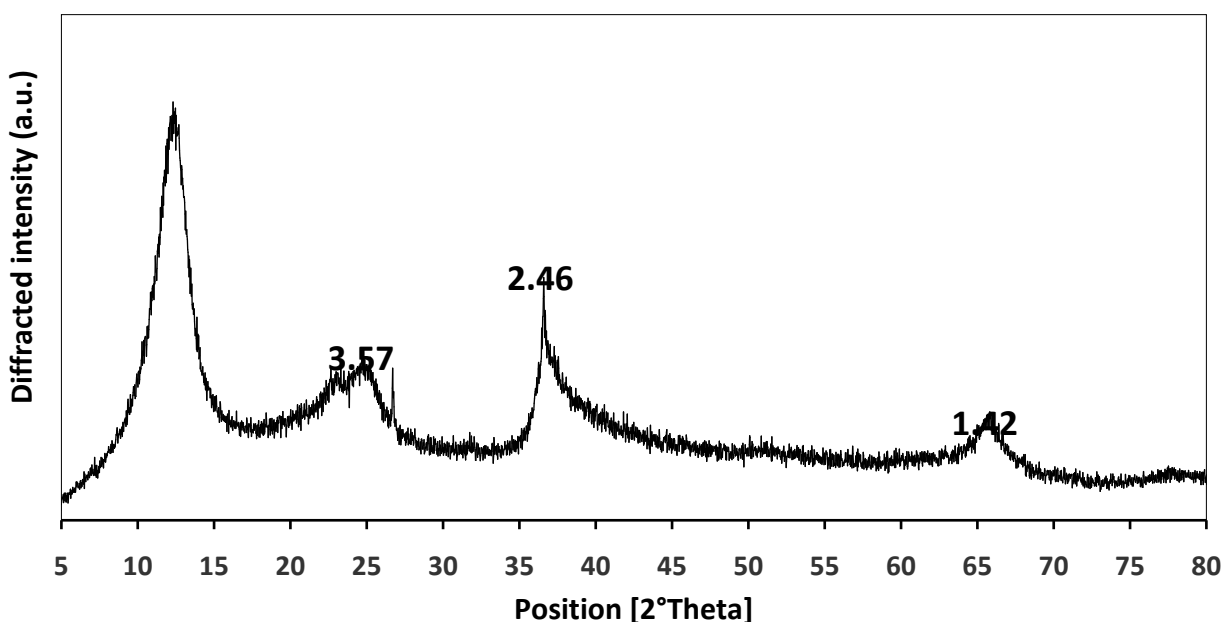
ESI5 HPLC analysis details: all the analyses were performed with a mobile phase: A – acetonitrile, B- formic acid/0.01M ammonium formate, pH 3.3.

Substance	Program (isocratic or gradient) ACN/buffer	Retention time [min]	Wavelength [nm]	LOD/LOQ [mg/L]	R ²
Diclofenac	Iso., 70/30	5.55	283	0.06/0.17	0.999
Isoxazole	Iso., 30/70	3.95	220	0.4/0.12	0.999
Sulfanilamide	Iso., 30/70	3.92	260	0.02/0.05	0.999
Sulfamethoxazole	Iso., 40/60	5.64	269	0.03/0.08	0.997
Syringaldehyde and Sulfamethoxazole	Iso., 30/70	5.71 8.06	307 269	0.22/0.66 0.03/0.08	0.998 0.998
Isoxazole and Sulfanilamide and Syringaldehyde	Gradient mode, 15/85 0-7 min 30/70 7-14 min 80/20 14-22 min 15/85 22-30 min	5.06 4.66 12.11	260 220 307	0.02/0.07 0.07/0.11 0.20/0.66	0.995 0.998 0.993
Sulfadiazine	Iso., 30/70	5.92	269	0.04/0.13	0.995
Sulfadimethoxine	Iso., 50/50	4.79	271	0.03/0.09	0.999
Sulfamerazine	Iso., 30/70	4.92	269	0.02/0.07	0.999
Sulfapyridine	Iso., 30/70	4.66	266	0.02/0.07	0.999
Sulfathiazole	Iso., 30/70	4.39	287	0.02/0.07	0.999
Sulfamethizole	Iso., 30/70	5.28	282	0.02/0.07	0.999
Sulfamethazine	Iso., 30/70	5.48	268	0.03/0.08	0.996

ESI6 Identification of transformation products

The information-dependent acquisition (IDA) mode, combining EMS with EPI, was used to maximize the information obtained in one scan. Data recorded by the EMS-IDA-EPI method were collected in positive electrospray ionization (ESI+) and negative (ESI-) ionization modes, while the p-MRM method data were collected only in positive ionization mode. The EMS and EPI mass ranges were from m/z 50 to m/z 700, and the scan rates were 1000 Da/s. The IDA criteria were as follows: the trigger for EPI was the 1–2 most intense ions that exceeded 100 cps; the mass tolerance was 250 mDa; former target ions were excluded for 30 s after two occurrences; the maximum rolling collision energy allowed was 80 eV in ESI+ and -80 eV in ESI-; and the dynamic background subtraction was turned on. The presence of TPs in the samples identified by the p-MRM mode was confirmed by analyzing the mass spectra recorded in the EMS-IDA-EPI mode. Non-targeted analysis was performed using a retrospective approach to mass spectral analysis.

ESI7 X-ray diffraction spectrum for MnOx used in the experiment



ESI8 SEM-EDS analysis

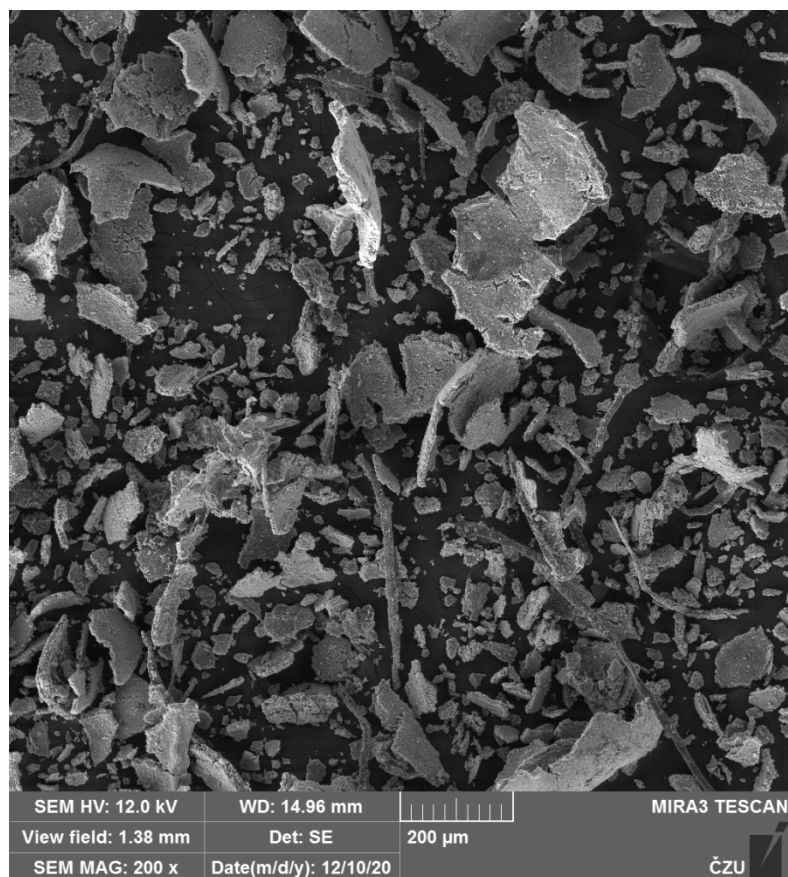


Figure SEM morphology

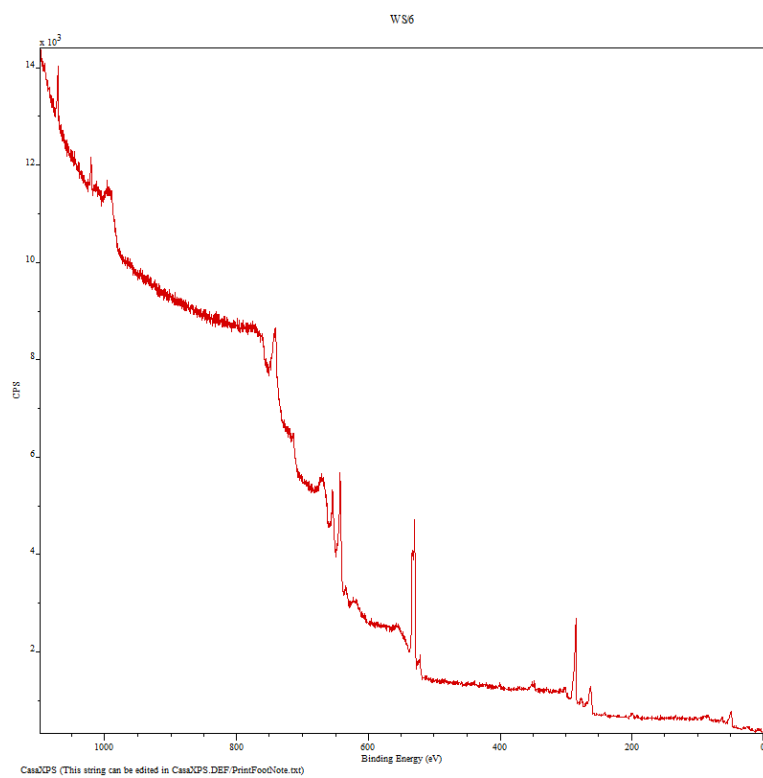
Table SEM-EDS analysis of the elemental composition of the filtration materials and amendments (wt%), n=10

	C	O	Na	Al	Si	P	Cl	Ca	Mn
Average	12.28%	26.02%	3.20%	0.94%	0.84%	0.46%	2.27%	2.36%	51.62%
SD	2.83%	3.61%	0.30%	0.01%	0.01%	0.01%	0.38%	0.20%	4.52%

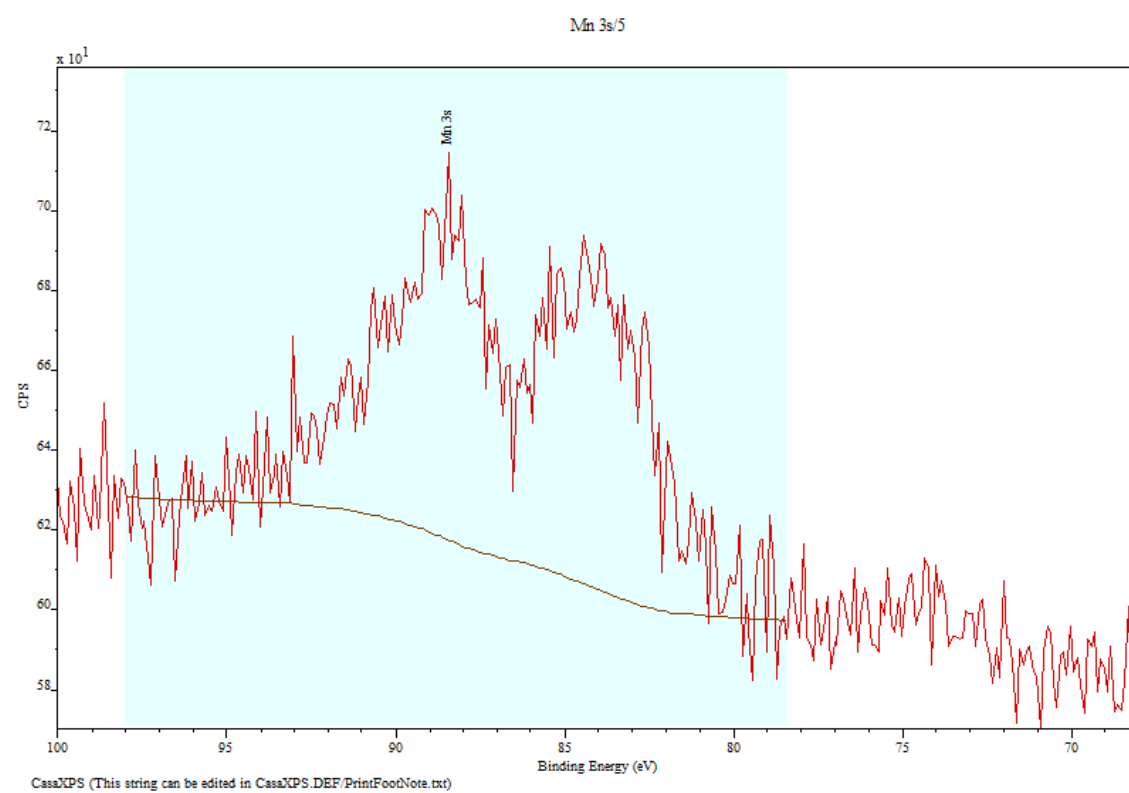
ESI9 Details on the XPS analysis of BioMnOx produced by *Pseudomonas putida* MnB6

	atom. %
C 1s	48.27
Mn 2p	7.38
O 1s	38.28
Cl 2p	1.34
Zn 2p	0.39
P 2p	0.70
Na 1s	2.63
Ca 2p	1.00

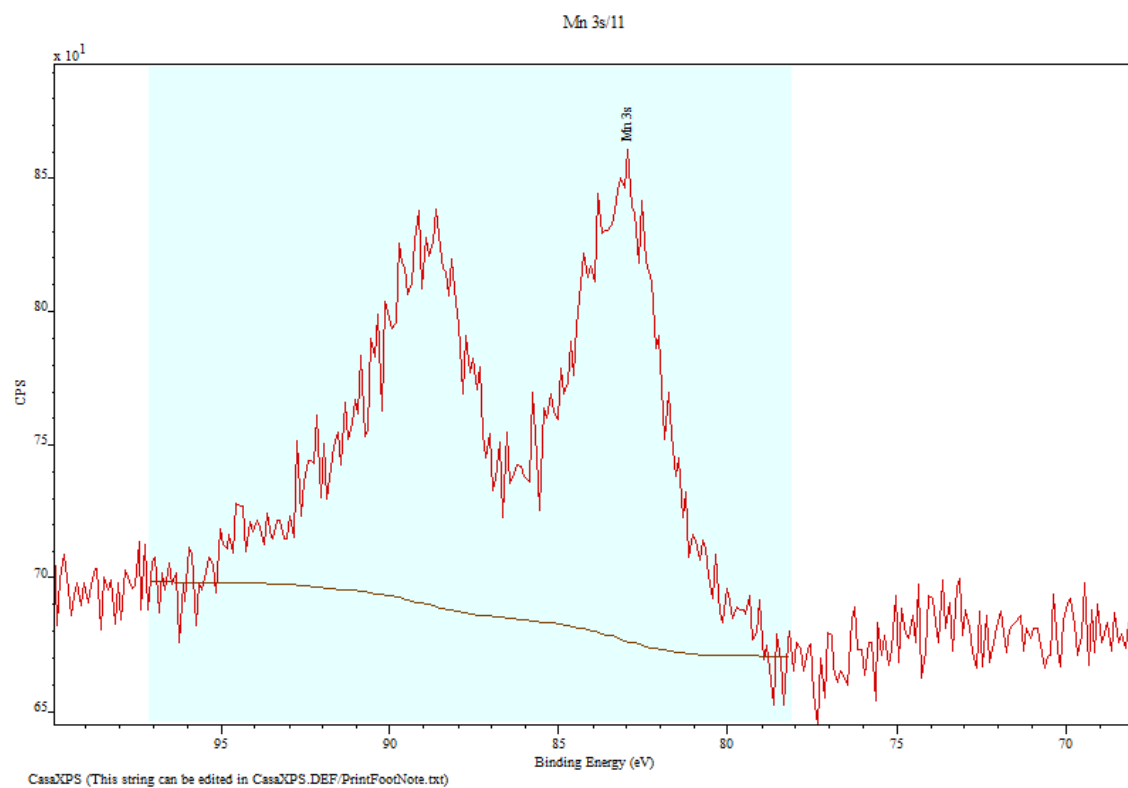
Full scan reveals that biogenic material contains mostly Mn, O and C.



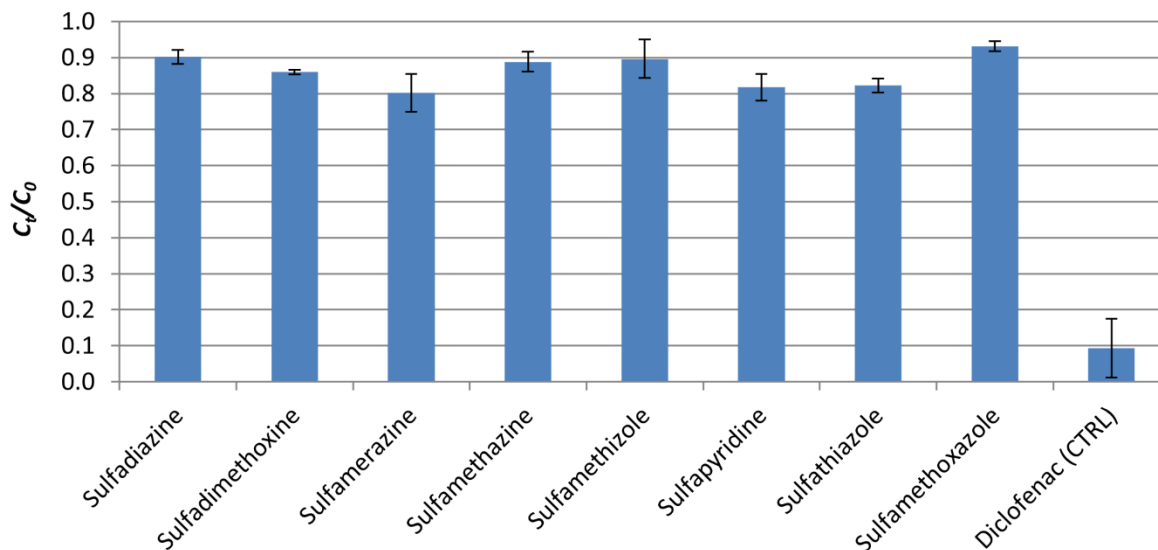
Mn 3s/5 before sputtering



Mn 3s/5 after sputtering



ESI10 Removal (C_t/C_0) of selected sulfonamides and diclofenac (positive control) in active cultures of *P. putida* MnB6 with peroxidized BioMnOx (Bio-Mn experiment) after time (t) 48 h.



ESI11 Improvements of the degradation of SMX, ISX and SNM in the presence of SAH

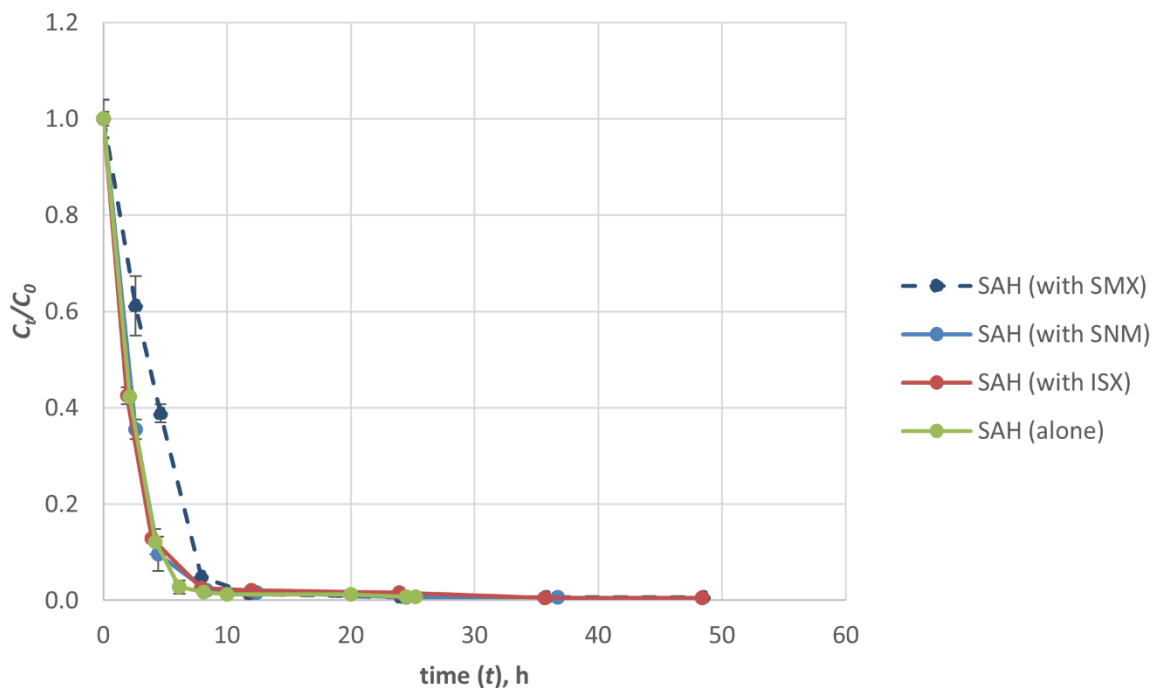
	SMX	SNM	ISX
Bio-Mn	+26%	+58%	+27%
Bio-noMn	+6%	+5%	+13%
Abio-Mn	+14%	+34%	+6%

ESI12 First-order kinetics parameters in the Bio-Mn and Bio-Mn-SAH tests (for the SAH experiments without the decoupling period)

Substance	k , 1/h	SE	$t_{50\%}$, h	$t_{99\%}$, d	R
SMX, w/o SAH	0.001	0.0002	693	192	0.842
SMX, w. SAH	0.029	0.0042	23.9	6.62	0.893
ISX, w/o SAH	0.001	0.0001	693	192	0.983
ISX, w. SAH	0.065	0.0128	10.6	2.95	0.906
SNM, w/o SAH	no fit				
SNM, w. SAH	0.174	0.0351	3.98	1.10	0.907
SAH (alone)	0.452	0.0194	1.53	0.42	0.998

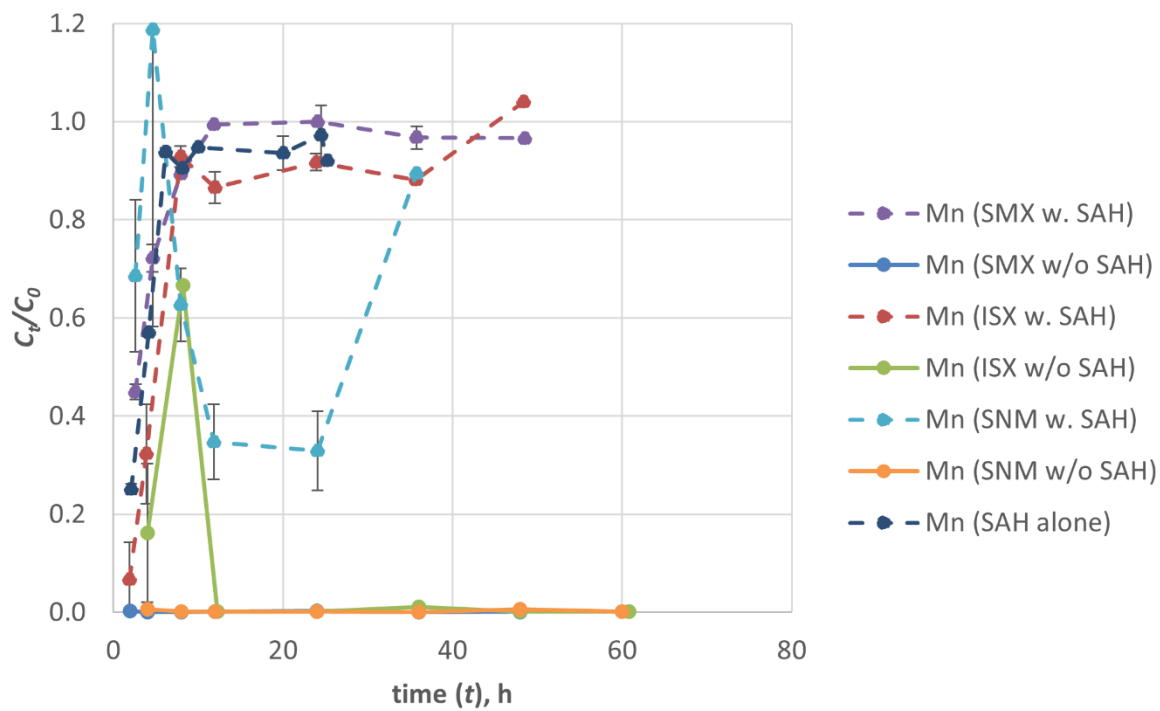
The parameters were determined using least squares non-linear regression (Levenberg-Marquardt estimation method in Software Statistica 14 (from Tibco Software Inc.).

ESI13 The removal (C_t / C_0) time profile of SAH in the Bio-Mn experiments with specific compounds SMX, SNM or ISX.

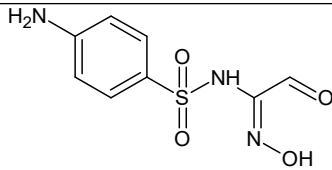
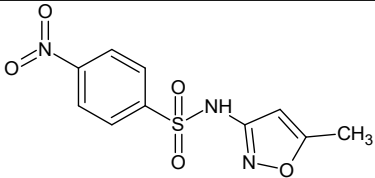
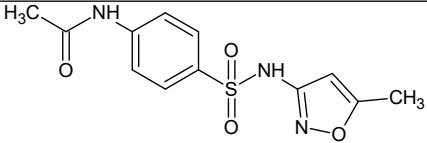
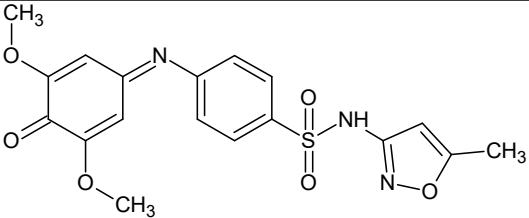


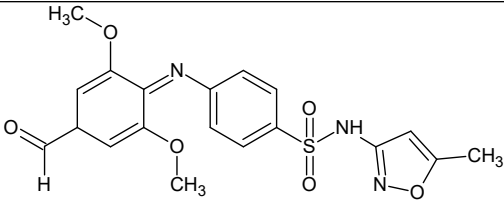
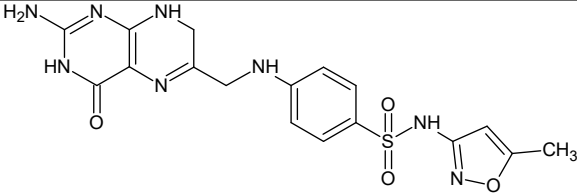
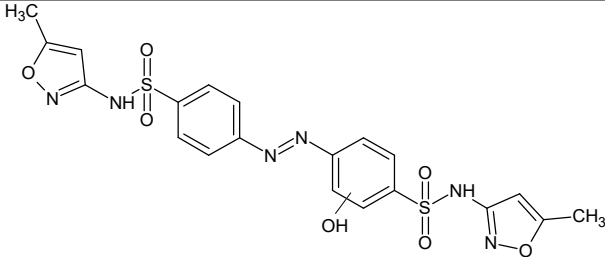
The removal (C_t/C_0) of SAH in the Abio-Mn experiment was 0.75 ± 0.04 , and in the Bio-noMn was 0.009 ± 0.001 .

ESI14 The removal (C_t/C_0) time profile of dissolved Mn in the experiments with specific compounds SMX, SNM or ISX (in the presence or absence of SAH) and SAH alone

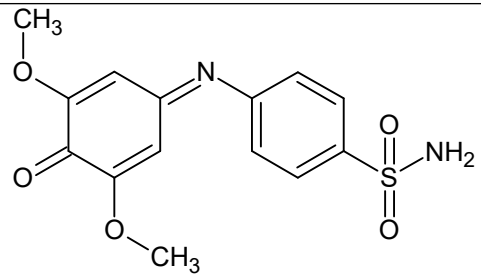
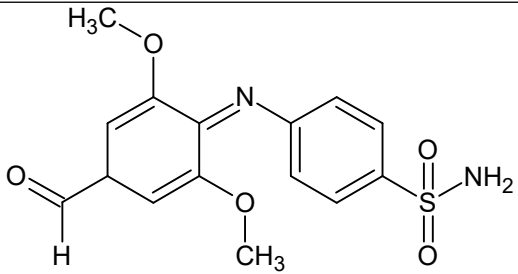


ESI15 Transformation products of sulfamethoxazole (SMX)

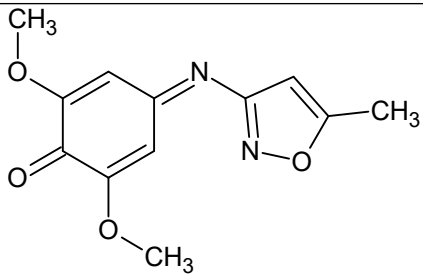
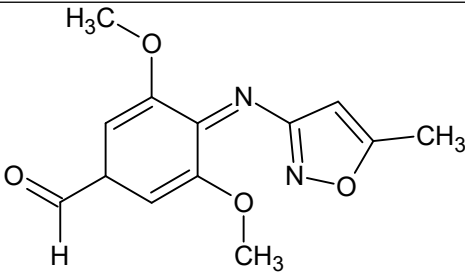
Symbol	Name	Transition ions/ product mass, m/z	Ionisation mode	Proposed chemical structure	Reference
TP244	4-amino-N-(1-hydroxyimino-2-oxaethyl)benzenesulfonamide	245.0 217.0	ESI+		Sochacki et al., 2021
TP283	4-nitro-SMX	284.2 267.0	ESI+		Sun et al., 2019
TP295	N4-acetyl-SMX	295.9 215.0; 249.9; 277.9	ESI+		Sochacki et al., 2021
TP403	SMX-DMBQ	403.9 243.0; 375.9	ESI+		Song et al., 2019

TP417	SMX-SAH	417.9 362.0; 389.9	ESI+	 <p>The structure shows a 5-methoxy-2-(4-methoxyphenyl)-6-(3-methylisoxazol-5-ylsulfamoyl)benzaldehyde derivative. It features a central benzene ring with a methoxy group at position 5, a 4-methoxyphenyl group at position 2, and a 3-methylisoxazol-5-ylsulfamoyl group at position 6.</p>	Margot et al., 2015
TP430	7,8-Dihydropterin-SMX	431.1 311.1	ESI+	 <p>The structure shows a 7,8-dihydropterin derivative linked to a 3-methylisoxazol-5-ylsulfamoyl group via a methylene bridge. The pterin ring has an amino group at position 6 and a carbonyl group at position 4.</p>	Xiong et al., 2020
TP518	Hydroxyazosulfamethoxazole	519.1 419.0; 373.1	ESI+	 <p>The structure shows a hydroxyazosulfamethoxazole derivative linked to a 3-methylisoxazol-5-ylsulfamoyl group via an azo bridge. The hydroxyazosulfamethoxazole part consists of a 5-methoxyisoxazole ring connected to a sulfamoyl group, which is further connected to a 4-hydroxyphenyl ring via an azo bridge.</p>	Yang et al., 2017

ESI16 Transformation products of sulfanilamide (SNM)

Symbol	Name	Transition ions/ product mass, m/z	Ionisation mode	Proposed chemical structure	Reference
TP322(SNM)	SNM-DMBQ	322.9 263.0; 295.0	ESI+		This work
TP337(SNM)	SNM-SAH	338.2 303.2; 321.1	ESI+		This work

ESI17 Transformation products of isoxazole (ISX)

Symbol	Name	Transition ions/ product mass, m/z	Ionisation mode	Proposed chemical structure	Reference
TP248(ISX)	ISX-DMBQ	248.9 178.1; 221.1	ESI+		This work
TP262(ISX)	ISX-SAH	263.0 207.0; 234.9	ESI+		This work

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