

Supplementary Information

Nitrous Oxide Is a Potent Inhibitor of Bacterial Reductive Dechlorination

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This supplementary file contains: 6 tables and 3 figures on 7 pages.

Supplementary Tables S1-S6 (pages 2-4); Supplementary Figures S1-S3 (pages 5-7)

Table S1. Inhibition models used in whole cell suspension assays.

Michaelis-Menten Equation: $v_0 = \frac{V_{\max} [S]}{K_m + [S]} \quad (1)$	Noncompetitive inhibition: $v_0 = \frac{V_{\max} [S]}{\alpha(K_m + [S])} \quad (3)$
Competitive inhibition: $v_0 = \frac{V_{\max} [S]}{(\alpha K_m + [S])} \quad (2)$	Uncompetitive inhibition: $v_0 = \frac{V_{\max} [S]}{(K_m + \alpha[S])} \quad (4)$
For simplification, the inhibitor concentrations and inhibition constants in equations (2), (3) and (4) are expressed as α , whereby $\alpha = 1 + \frac{[I]}{K_i}$	

Table S2. Initial PCE-to-cDCE dechlorination rates versus PCE concentrations in Geo strain SZ cell suspension assays in the presence of 0, 10 and 60 μM N_2O .

PCE (μM)	V^a	PCE (μM)	V^a	PCE (μM)	V^a
No N_2O		10 μM N_2O		60 μM N_2O	
3.6	4.1	10.9	19.3	4.9	5.3
6.3	11.7	13.1	24.7	9.5	8.5
11.7	24.0	18.5	25.3	15.3	13.6
15.7	28.5	29.0	28.5	52.7	20.0
42.2	48.4	41.3	36.8	59.4	22.1
60.8	60.3	57.0	41.2	69.5	24.3
83.6	63.9	63.6	44.7	87.5	24.5
97.8	65.0	74.1	45.0	110.4	25.5
110.2	61.3	91.6	51.9	124.3	25.4
122.0	58.4	109.7	47.6	146.5	25.2
133.9	68.0				
153.9	56.7				

^a Initial dechlorination rate (nmol Cl^- released min^{-1} mg protein $^{-1}$).

Table S3. Initial cDCE-to-VC dechlorination rate versus cDCE concentrations in *Dhc* strain BAV1 cell suspension assays in the presence of 0, 10 and 60 μM N_2O .

cDCE (μM)	V^a	cDCE (μM)	V^a	cDCE (μM)	V^a
No N_2O		10 μM N_2O		60 μM N_2O	
2.7	15.4	1.9	10.3	1.7	2.4
4.4	27.6	5.0	16.9	5.4	4.9
7.5	48.6	8.4	23.6	10.1	7.2
17.5	53.2	17.5	36.4	17.5	14.2
31.7	79.5	31.7	38.5	29.5	18.5
64.5	93.1	63.4	54.6	63.4	24.7
174.9	110.3	161.8	75.4	208.8	39.6
250.3	106.5	211.0	80.5	225.2	42.6
377.1	111.4	488.6	86.4	424.6	42.1

^a Initial dechlorination rate ($\text{nmol Cl}^- \text{ released min}^{-1} \text{ mg protein}^{-1}$).

Table S4. Initial VC-to-ethene dechlorination rate versus VC concentrations in *Dhc* strain BAV1 cell suspension assays in the presence of 0, 15 and 50 μM N_2O .

VC (μM)	V^a	VC (μM)	V^a	VC (μM)	V^a
No N_2O		15 μM N_2O		50 μM N_2O	
4.4	18.5	3.2	7.9	4.1	3.0
13.9	53.3	11.4	19.2	11.3	7.0
23.5	72.4	22.0	25.2	21.7	10.9
56.3	89.9	51.4	34.4	47.3	14.2
98.5	104.3	87.9	39.2	72.6	16.7
116.0	107.6	118.0	42.6	113.3	17.1
149.0	107.6	126.0	43.5	140.8	20.9

^a Initial dechlorination rate ($\text{nmol Cl}^- \text{ released min}^{-1} \text{ mg protein}^{-1}$).

Table S5. Statistical parameters (R^2 , AICc and $Sy.x$ values) used for determining the best-fit inhibition model and inhibition constants in cell suspensions amended with N_2O as inhibitor.

Culture	Substrate	Inhibitor	Tested models	Statistical Parameters			K_i (μM)
				R^2	AICc	$Sy.x$	
Geo strain SZ	PCE	N_2O	Noncompetitive	0.971	81.338	3.350	40.8 ± 3.8
			Uncompetitive	0.966	86.477	3.639	29.1 ± 3.1
			Competitive	0.944	102.041	4.678	9.2 ± 1.9
Dhc strain BAV1	cDCE	N_2O	Noncompetitive	0.968	107.060	6.422	21.2 ± 3.5
			Competitive	0.952	117.928	7.853	2.3 ± 0.5
			Uncompetitive	0.938	124.696	8.902	25.9 ± 2.9
Dhc strain BAV1	VC	N_2O	Noncompetitive	0.996	43.791	2.386	9.6 ± 0.4
			Uncompetitive	0.986	69.426	4.393	7.0 ± 0.6
			Competitive	0.974	82.243	5.961	1.6 ± 0.3

R^2 , the Coefficient of Determination, gives information about the fit of the measured data to the different models tested, and the model with highest R^2 value provides the best data fit.

The AICc (i.e., corrected Akaike's Information Criterion) offers an estimate of the relative quality of tested models, and the model with the lowest AICc value represents the relative best fit among the tested models.

The $Sy.x$ represents the Standard Deviation of the Residuals, and the model with the lowest $Sy.x$ value provides best prediction of the data.

In all cell suspensions assays, the noncompetitive model (highlighted in bold) gave the highest R^2 and the lowest AIC and $Sy.x$ values.

Table S6. Initial versus final amounts of N_2O measured of Geo strain SZ and Dhc strain BAV1 cultures growing with chlorinated ethenes as electron acceptors in 160-mL vessels with 60 mL headspace.

Culture	e^- Acceptor	Inhibitor N_2O (μM)	Inhibitor N_2O (total $\mu mol/vessel$)	
			Initial	Final
Geo strain SZ	PCE	9.5	2.1 ± 0.1	2.1 ± 0.3
		19.1	4.1 ± 0.1	3.9 ± 0.5
		57.3	12.4 ± 1.0	12.6 ± 1.4
Dhc strain BAV1	cDCE	9.5	2.1 ± 0.2	2.1 ± 0.3
		29.0	6.3 ± 0.5	6.8 ± 0.8
		57.3	12.4 ± 0.5	12.2 ± 0.4
Dhc strain BAV1	VC	2.9	0.6 ± 0.1	0.6 ± 0.1
		5.7	1.2 ± 0.1	1.2 ± 0.2
		19.1	4.1 ± 0.6	4.1 ± 0.4

Error values represent the standard deviation based on measurements of triplicate cultures.

Supplementary Figures

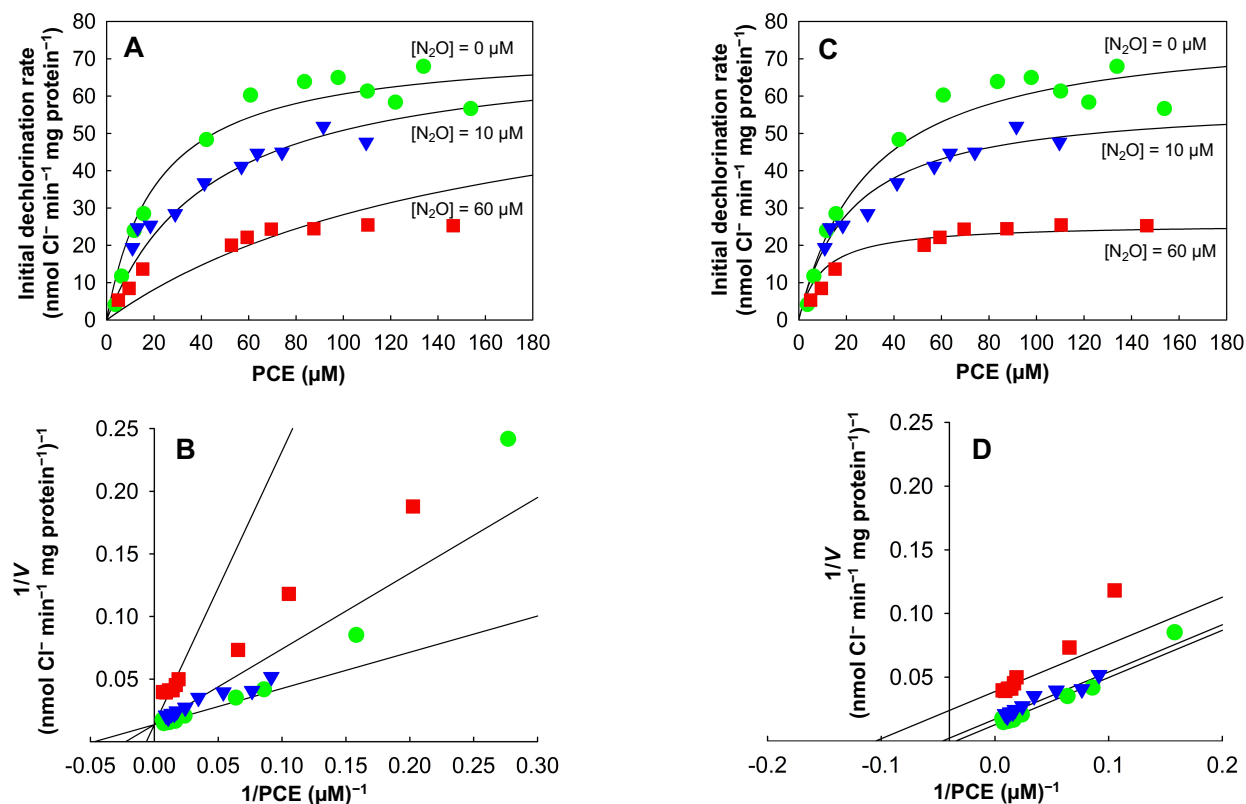


Figure S1. Competitive and uncompetitive N₂O inhibition kinetics of PCE-to-cDCE reductive dechlorination in *Geo* strain SZ cell suspensions. **Panels A and B** show Michaelis-Menten and Lineweaver-Burk plots, respectively, of competitive N₂O inhibition of PCE-to-cDCE reductive dechlorination in *Geo* strain SZ cell suspensions without and in the presence of 10 and 60 μM N₂O. **Panels C and D** depict Michaelis-Menten and Lineweaver-Burk plots, respectively, of uncompetitive N₂O inhibition of PCE-to-cDCE reductive dechlorination in *Geo* strain SZ cell suspensions without and in the presence of 10 and 60 μM N₂O. Solid lines represent the model simulation to each data set based on the nonlinear regression using the SigmaPlot 13 Enzyme Kinetic Module. Solid green circles represent rate data measured in the absence of N₂O; solid blue triangles show rate data measured in the presence of 10 μM N₂O; and solid red squares show rate data measured in the presence of 60 μM N₂O.

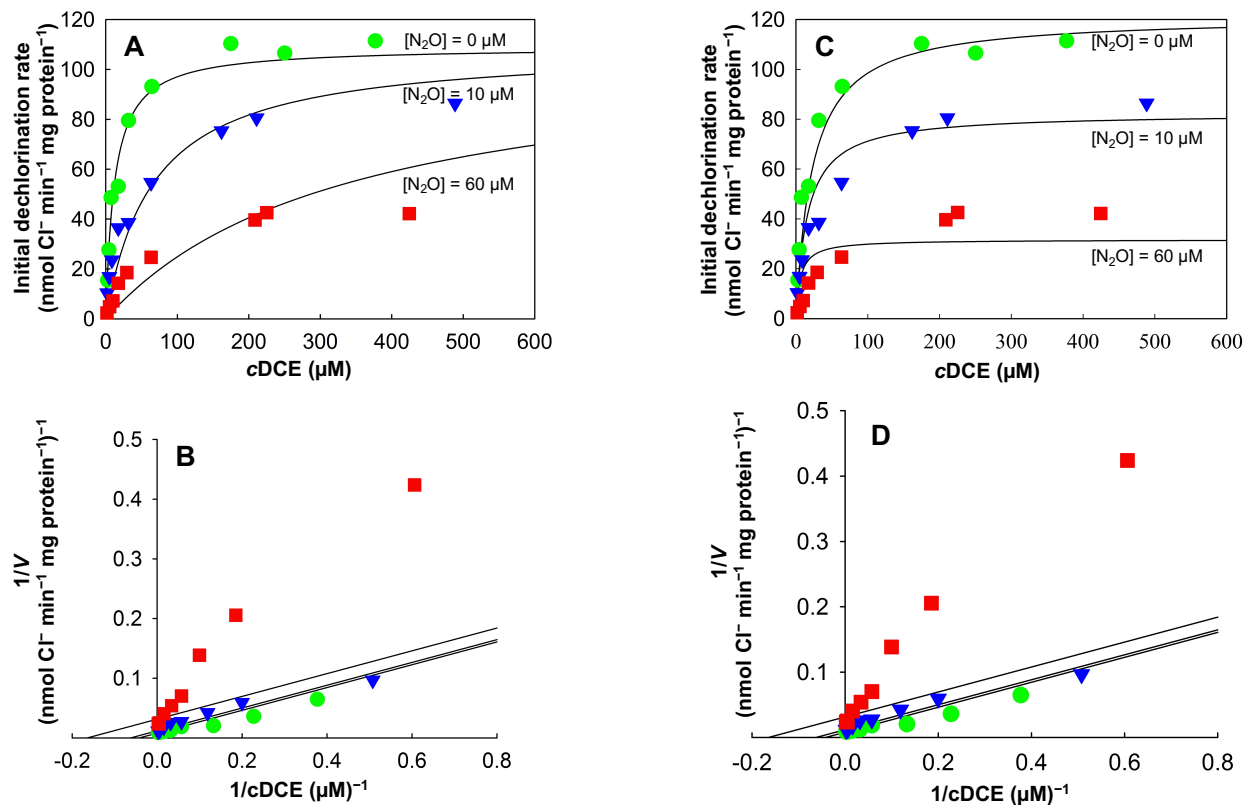


Figure S2. Competitive and uncompetitive N₂O inhibition kinetics of cDCE-to-VC reductive dechlorination in *Dhc* strain BAV1 cell suspensions. **Panels A and B** show Michaelis-Menten and Lineweaver-Burk plots, respectively, of competitive N₂O inhibition of cDCE-to-VC reductive dechlorination in *Dhc* strain BAV1 cell suspensions without and in the presence of 10 and 60 μM N₂O. **Panels C and D** depict Michaelis-Menten and Lineweaver-Burk plots, respectively, of uncompetitive N₂O inhibition of VC-to ethene reductive dechlorination in *Dhc* strain BAV1 cell suspensions without and in the presence of 10 and 60 μM N₂O. Solid lines represent the model simulation to each data set based on the nonlinear regression using the SigmaPlot 13 Enzyme Kinetic Module. Solid green circles represent rate data measured in the absence of N₂O; solid blue triangles show rate data measured in the presence of 10 μM N₂O; and solid red squares show rate data measured in the presence of 60 μM N₂O.

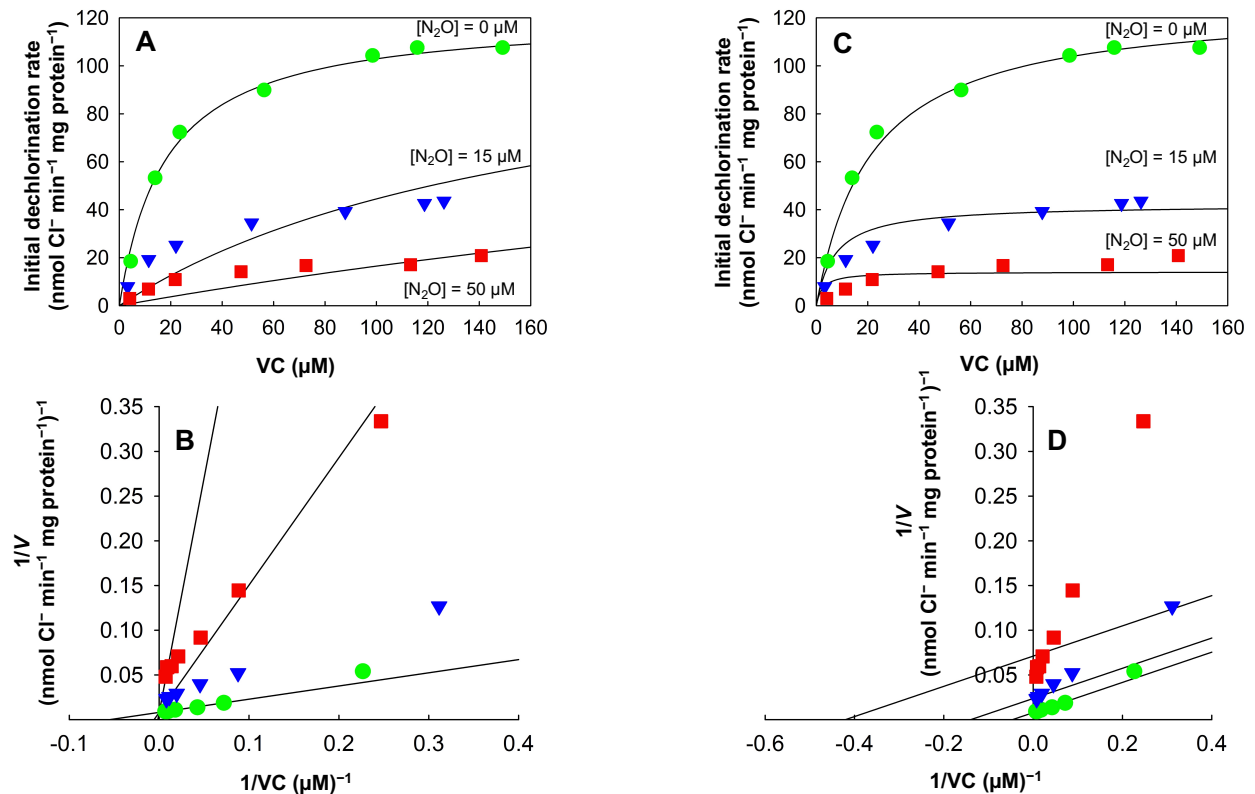


Figure S3. Competitive and uncompetitive N₂O inhibition kinetics of VC-to-ethene reductive dechlorination in *Dhc* strain BAV1 cell suspensions. **Panels A and B** show Michaelis-Menten and Lineweaver-Burk plots, respectively, of competitive N₂O inhibition of VC-to-ethene reductive dechlorination in cell suspensions of *Dhc* strain BAV1 without and in the presence of 15 and 50 μM N₂O. **Panels C and D** depict Michaelis-Menten and Lineweaver-Burk plots, respectively, of uncompetitive N₂O inhibition of VC-to-ethene reductive dechlorination in *Dhc* strain BAV1 cell suspensions without and in the presence of 15 and 50 μM N₂O. Solid lines represent the model simulation to each data set based on the nonlinear regression using the SigmaPlot 13 Enzyme Kinetic Module. Solid green circles represent rate data measured in the absence of N₂O; solid blue triangles show rate data measured in the presence of 15 μM N₂O; and solid red squares show rate data measured in the presence of 50 μM N₂O.