

This is the accepted manuscript version of the contribution published as:

Woods, A., Kuntze, K., Gelman, F., Halicz, L., Nijenhuis, I. (2018):
Variable dual carbon-bromine stable isotope fractionation during enzyme-catalyzed reductive dehalogenation of brominated ethenes
Chemosphere **190** , 211 – 217

The publisher's version is available at:

<http://dx.doi.org/10.1016/j.chemosphere.2017.09.128>

1 **Variable dual carbon-bromine stable isotope fractionation during enzyme-catalyzed**
2 **reductive dehalogenation of brominated ethenes**

3

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13 **Manuscript for Chemosphere**

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19 Abstract

20 The potential of compound-specific stable isotope analysis (CSIA) to characterize
21 biotransformation of brominated organic compounds (BOCs) was assessed and compared to
22 chlorinated analogues. *Sulfurospirillum multivorans* and *Desulfitobacterium hafniense* PCE-S
23 catalyzed the dehalogenation of tribromoethene (TBE) to either vinyl bromide (VB) or ethene,
24 respectively. Significantly lower isotope fractionation was observed for TBE dehalogenation by
25 *S. multivorans* ($\epsilon_C = -1.3 \pm 0.2\text{‰}$) compared to *D. hafniense* ($\epsilon_C = -7.7 \pm 1.5\text{‰}$). However, higher
26 fractionation was observed for dibromoethene (DBE) dehalogenation by *S. multivorans* ($\epsilon_C = -$
27 $16.8 \pm 1.8\text{‰}$ and $-21.2 \pm 1.6\text{‰}$ for *trans*- and *cis*-1,2- (DBE) respectively) compared to *D.*
28 *hafniense* PCE-S ($\epsilon_C = -9.5 \pm 1.2\text{‰}$ and $-14.5 \pm 0.7\text{‰}$ for *trans*-1,2-DBE and *cis*-1,2-DBE,
29 respectively). Significant, but similar, bromine fractionation was observed for for *S. multivorans*
30 ($\epsilon_{Br} = -0.53 \pm 0.15\text{‰}$, $-1.03 \pm 0.26\text{‰}$, and $-1.18 \pm 0.13\text{‰}$ for *trans*-1,2-DBE, *cis*-1,2-DBE and TBE,
31 respectively) and *D. hafniense* PCE-S ($\epsilon_{Br} = -0.97 \pm 0.28\text{‰}$, $-1.16 \pm 0.36\text{‰}$, and $-1.34 \pm 0.32\text{‰}$ for
32 *cis*-1,2-DBE, TBE and *trans*-1,2-DBE, respectively). Variable C-Br dual-element slopes were
33 estimated at $\Lambda (\epsilon_C / \epsilon_{Br}) = 1.03 \pm 0.2$, 17.9 ± 5.8 , and 29.9 ± 11.0 for *S. multivorans* debrominating
34 TBE, *cis*-1,2-DBE and *trans*-1,2-DBE, respectively, and at 7.14 ± 1.6 , 8.27 ± 3.7 , and 8.92 ± 2.4 for
35 *D. hafniense* PCE-S debrominating *trans*-1,2-DBE, TBE and *cis*-1,2-DBE, respectively. A high
36 variability in isotope fractionation, which was substrate property related, was observed for *S.*
37 *multivorans* but not *D. hafniense*, similar as observed for chlorinated ethenes, and may be due to
38 rate-limiting steps preceding the bond-cleavage or differences in the reaction mechanism.
39 Overall, significant isotope fractionation was observed and, therefore, CSIA can be applied to
40 monitor the fate of brominated ethenes in the environment. Isotope effects differences, however,
41 are not systematically comparable to chlorinated ethenes.

42

43 **Keywords:** CSIA, reductive dehalogenation, brominated organic compounds

44

45 **1. Introduction**

46 Several chlorinated as well as brominated organic compounds (BOCs) are pervasive
47 environmental contaminants (de Wit, 2002; Alaei, 2003); however, while biotransformation
48 processes have been extensively investigated for chlorinated compounds (Bradley and Chapelle,
49 2010), very little is known regarding the environmental fate and transport of BOCs, such as the
50 flame-retardants like polybrominated diphenyl ethers (PBDEs), and fumigants like ethylene
51 dibromide (EDB) and methyl bromide (Waaijers and Parsons, 2016).

52 Microbial reductive dehalogenation is widely regarded as a key process in organohalide removal
53 in environmental systems, and reductive dehalogenation of chloroethenes, for example, has been
54 subject to considerable study (Smidt, 2004; Nijenhuis and Kuntze, 2016). Furthermore, validated
55 concepts and approaches have been developed to address the fate of these chlorinated substances
56 *in situ* (Bombach et al., 2010), but have yet to be developed and validated for their brominated
57 analogues.

58 Compound-specific stable isotope analysis (CSIA) may be a suitable approach to monitor and
59 characterize biodegradation of BOCs, as it is currently well developed for many chlorinated
60 groundwater contaminants (for a review see (Hunkeler et al., 2008; Nijenhuis and Kuntze,
61 2016)). CSIA utilizes the relatively faster chemical reaction of molecules with a lighter isotope at
62 the reactive position, resulting in an enrichment of heavy isotopes in the residual contaminant

63 pool (i.e. fractionation) and allowing for the calculation of reaction-specific enrichment factors
64 (ϵ), identifiers for detecting and monitoring *in situ* contaminant transformation (Hunkeler et al.,
65 2008).

66 Carbon isotope analysis has been extensively reported during microbial reductive dechlorination
67 of chloroethenes (see e.g. (Lee, 2007; Cichocka, 2008)), and dual-element carbon-chlorine
68 isotope analysis has also emerged for the same transformation process (Abe, 2009; Wiegert,
69 2013; Badin, 2014). For example, dual-element isotope analysis was described for reductive
70 dechlorination of chloroethenes by *Sulfurospirillum multivorans* and *Desulfitobacterium*
71 *hafniense* PCE-S (Cretnik, 2013; Renpenning, 2014). These bacterial strains are known also to
72 reductively debrominate bromoethenes (Ye, 2010), reported soil and groundwater contaminants
73 (Patterson et al., 2007). In this study, therefore, *S. multivorans* and *D. hafniense* PCE-S were
74 selected as model strains for a systematic comparison of stable isotope fractionation during
75 enzymatic debromination versus dechlorination.

76 Principally, the magnitude of isotope fractionation is determined by the rate-determining step of
77 a reaction. The carbon isotope effects during the expected rate-determining step for reductive
78 dehalogenation, the carbon-halogen bond cleavage, are expected to be similar based on the
79 comparison of theoretical maximum kinetic isotope effects (KIEs) with $KIE_C = 1.057$ for C-Cl
80 (Elsner and 2005) and 1.043 for C-Br bond cleavage (Zakon, 2013). Bromine stable isotope
81 effects are, however, expected to be considerably smaller compared to chlorine stable isotope
82 effects considering $KIE_{Br} = 1.002$ (Zakon, 2013) compared to $KIE_{Cl} = 1.013$ (Elsner, 2010).
83 Previous reports for the reductive dehalogenation of chloroethenes, however, have already
84 shown that fractionation patterns are difficult to predict based on theoretical KIE alone but may
85 be masked by uptake and binding of the substrate to the enzyme influenced by the substrate

86 hydrophobicity and cell composition (Nijenhuis, 2005; Cichocka, 2007; Renpenning, 2015). The
87 reaction mechanism of carbon-chlorine bond cleavage was reflected in dual-element, carbon-
88 chlorine, fractionation patterns for trichloroethene (TCE) however, not for tetrachloroethene
89 (PCE) during microbial dehalogenation (Cretnik, 2013; Cretnik, 2014; Renpenning, 2014). This
90 variability was proposed to be associated with the relatively fast intrinsic reaction rate for PCE
91 compared to its relatively slow overall transport rate. Therefore, stable isotope patterns are
92 assumed to reflect reactions steps, such as enzyme binding, prior to bond cleavage (Renpenning,
93 2014). Similar effects were observed comparing the abiotic vs. biotic dihaloelimination of
94 ethylene dibromide (Kuntze et al., 2016). Recently, in a computational modeling study, Ji et al.
95 (2017) inferred that highly chlorinated ethenes (e.g. TCE and PCE) primarily react via an inner
96 sphere nucleophilic substitution mechanism, whereas the less chlorinated ethenes (e.g. cis- and
97 trans-DCE) mainly react through an inner sphere nucleophilic addition pathway. Although the
98 exact mechanism of dehalogenation by cobamide-based dehalogenases is still under debate,
99 several pathways can be considered: Co-C bond formation after direct Co(I) attack on the carbon
100 backbone of the organohalide (Schrauzer et al., 1969); Co-X bond formation after direct Co(I)
101 attack on the halogen atom (Payne et al., 2015); long-range electron transfer leading to substrate
102 radical formation (Kunze et al., 2017).

103 Thus far, it is not clear if there are similarities or differences in the stable isotope fractionation
104 for brominated and chlorinated analogues and if the fractionation patterns for brominated
105 compounds can be predicted from their chlorinated analogues. In this study, therefore, the
106 carbon and bromine stable isotope fractionation was investigated during reductive
107 dehalogenation of bromoethenes by crude extracts of *S. multivorans* and *D. hafniense* PCE-S, for
108 a direct comparison to previous results for chlorinated ethenes. Enrichment factors, ϵ_C and ϵ_{Br} ,

109 dual-element plots $\Lambda (\epsilon_C / \epsilon_{Br})$ and apparent kinetic isotope effects (AKIE) were determined for
110 tribromoethene (TBE) and *cis/trans*-1,2-dibromoethene (DBE) and evaluated against those
111 reported for their chlorinated analogs.

112

113

114 2. MATERIALS AND METHODS

115 ***2.1 Chemicals.***

116 All chemicals were purchased from Sigma-Aldrich Chemie (Seelze, Germany) or Merck
117 (Darmstadt, Germany) and were of analytical grade. Additionally, ethene was purchased from
118 Airproducts (Hattingen, Germany), tribromoethene (TBE) and 1,2-dibromoethene (1,2-DBE; *cis*-
119 and *trans*- mixture) were purchased from ABCR (Karlsruhe, Germany), and vinyl bromide (VB)
120 was purchased from Aldrich.

121 ***2.2 Cultivation of bacteria and preparation of crude extracts.***

122 *S. multivorans* (Miller et al., 1998) and *D. hafniense* strain PCE-S (Miller et al., 1997) were
123 cultivated as previously described. During cultivation, PCE was provided as terminal electron
124 acceptor and pyruvate as electron donor. Crude extracts were prepared in triplicate as previously
125 described (Nijenhuis, 2005) under anoxic conditions (N_2/H_2 atmosphere) within an anoxic
126 glovebox (Coy Laboratory Products Inc., USA), or under a steady stream of nitrogen gas.

127 ***2.3 Reductive debromination assays.***

128 For determination of carbon and bromine stable isotope fractionation, reductive debromination
129 assays were developed as previously described for reductive dechlorination (Nijenhuis, 2005).
130 Provided as sole electron acceptor, either TBE or 1,2-DBE dissolved in ethanol were transferred
131 at a final concentration of 1 mM to 4 ml reduced 1.6 mM methyl viologen buffer. Degradation
132 was then facilitated by the addition of crude extracts of active enzyme and the extent of
133 degradation was controlled by the addition of different concentrations of artificial electron
134 acceptor titanium (III) citrate. Experiments were prepared in triplicate from three independent
135 crude extracts. Abiotic controls were prepared for each assay to control for chemical reduction
136 of the substrate by titanium(III) citrate and methyl viologen. All reactions were stopped by the
137 addition of 1 mL saturated Na₂SO₄ (pH 1). Immediately following termination of reactions, 0.5
138 mL headspace was analyzed for concentrations of parent compound and products. Assays were
139 kept at 4° C until stable isotope analysis.

140 ***2.4 Analytical methods.***

141 A gas chromatograph (Chrompack CP-3800; Varian, Middelburg, The Netherlands) equipped
142 with a flame ionization detector (GC-FID) and a GS-Q fused-silica capillary column (30 m x
143 0.53 mm; J&W Scientific, Waldbronn, Germany) was used for concentration analysis of
144 samples. Analysis was performed as previously reported (Nijenhuis, 2005), and is described in
145 detail in the Supplementary Information (SI).

146 Gas chromatography combustion isotope ratio mass spectroscopy (GC-C-IRMS) was used for
147 carbon stable isotope analysis of samples. Analysis was performed as previously reported
148 (Nijenhuis, 2005), and is described in detail in the SI.

149 Gas chromatography with multi-collector inductively coupled plasma mass spectrometry (GC-
150 MC-ICPMS) was used for bromine stable isotope analysis of samples. Analysis was performed
151 as previously reported (Gelman, 2010), and is described in detail in the SI.

152 *2.5 Calculations and evaluation of data.*

153 Carbon and bromine isotope enrichment factors ($\epsilon_{\text{C or Br}}$) were determined for TBE and *cis*- and
154 *trans*-1,2-DBE according to the Rayleigh equation (Elsner and 2005):

$$155 \quad \epsilon \times \ln\left(\frac{C_t}{C_0}\right) = \ln\left(\frac{R_t}{R_0}\right) \quad [1]$$

156 where C_t and C_0 are the concentrations of the parent compound at time t and time 0 respectively,
157 and $R = 1 + \delta$. $\epsilon_{\text{C or Br}}$ is reported in parts per thousand (‰). As assays were performed in
158 triplicate, data was combined in linear regressions (not forced through zero) used to generate
159 final ϵ values within a 95% confidence interval (95% CI) using Excel Analysis Toolpak
160 (Microsoft, USA).

161 The apparent kinetic isotope effect (AKIE) for carbon and bromine for each debromination
162 reaction was estimated using the following equation (Elsner, 2010):

$$163 \quad AKIE = \frac{1}{(1 + z \times \epsilon_{rp})} \quad [2]$$

164 where z is the number of atoms in intramolecular competition, and ϵ_{rp} is the reactive position-
165 specific ϵ which was calculated as follows:

$$166 \quad \epsilon_{rp} = \epsilon \times \frac{n}{x}$$

167 [3]

168 where $\epsilon_{C \text{ or } Br}$ is the bulk enrichment, n is the number of atoms of the element considered in the
169 molecule, and x is the number of atoms of the element considered at the reactive site.

170 Assuming reactions are stepwise and secondary isotope effects negligible, $x = z$, and after
171 elimination upon combining equations 2 and 3, the term can be simplified as:

$$172 \quad AKIE = \frac{1}{1+n \times \epsilon} \quad [4]$$

173 C-Br dual-element fractionation correlation was expressed as Λ and determined by linear
174 regression (not forced through zero) of $\delta^{81}Br$ data as a function of $\delta^{13}C$ data. As before, linear
175 regressions were performed within a 95% CI using Excel Analysis Toolpak.

176

177 3. RESULTS

178 ***3.1 Dehalogenation and isotope fractionation of brominated ethenes***

179 Reductive dehalogenation assays were prepared with crude extracts from *S. multivorans* and *D.*
180 *hafniense* PCE-S. Figure 1. illustrates the pathway reported by Ye *et al.* (2010) and subsequently
181 observed in this study. Dehalogenation occurred in all assays with TBE sequentially converted to
182 VB via branched intermediates *trans*-, *cis*-, and 1,1-DBE, with *D. hafniense* PCE-S additionally
183 producing small amounts of ethene.

184 ***3.2 Carbon stable isotope fractionation***

185 TBE and 1,2-DBE carbon stable isotope fractionation was evaluated for crude extracts of both *S.*
186 *multivorans* and *D. hafniense* strain PCE-S (SI Figure S1 & S3; Table 1). For *D. hafniense* PCE-
187 S, ϵ_c was calculated at $-7.7 \pm 1.5\%$, $-9.5 \pm 1.2\%$ and $-14.5 \pm 0.7\%$ for TBE, *trans*-1,2-DBE and

188 *cis*-1,2-DBE respectively. Fractionation of TBE was comparable to that reported for
189 dechlorination of TCE by *D. hafniense* PCE-S with $\epsilon_C = -10.9\text{‰}$ (Cichocka, 2007), and is within
190 range of that reported for PCE at -5.2 to -8.9 (Nijenhuis, 2005; Cichocka, 2007).

191 For *S. multivorans*, however, a weak carbon isotope enrichment at $\epsilon_C = -1.2 \pm 0.2\text{‰}$ was
192 observed for TBE, while a one order of magnitude larger enrichment was measured at $\epsilon_C = -16.8$
193 $\pm 1.8\text{‰}$ and $-21.2 \pm 1.6\text{‰}$ for *trans*- and *cis*-1,2-DBE, respectively. Observed carbon isotope
194 effects for TBE were much smaller than those reported for reductive dechlorination of TCE by *S.*
195 *multivorans*, where ϵ_C ranged from -13.2‰ to -26.0‰ , however, were within range of that
196 reported for PCE dechlorination ($\epsilon_C = -0.4\text{‰}$ to -2.2‰) (Nijenhuis, 2005; Lee, 2007; Cichocka,
197 2008; Renpenning, 2014; Renpenning, 2015).

198 AKIE_C values were calculated and compared to theoretical KIE (Table 1). All values, ranging
199 from 1.035 ± 0.0036 to 1.044 ± 0.0032 for *S. multivorans* debrominating *trans*- and *cis*-1,2-DBE
200 respectively, and 1.016 ± 0.0030 , 1.019 ± 0.0024 , and 1.030 ± 0.0007 for *D. hafniense* PCE-S
201 debrominating TBE, *trans*-1,2,DBE and *cis*-1,2-DBE respectively, were similar (within the same
202 order of magnitude) to theoretical KIE calculated at 1.043 for a simple C-Br bond cleavage
203 (Zakon *et al.*, 2013). One notable exception was TBE debromination by *S. multivorans* which
204 was one order of magnitude smaller with an AKIE value of 1.002 ± 0.0004 , a further indication
205 that in this particular reaction, either rate-limitation preceding C-Br bond cleavage contributes to
206 considerable masking effects or a different reaction mechanism is employed.

207 The occurrence of significantly smaller fractionation for *S. multivorans* debrominating TBE fits
208 with the proposition by Renpenning *et al* (2015) that substrate hydrophobicity may cause rate-
209 limitation at the active-site of the enzyme, resulting in masking of intrinsic isotope effects. It is

210 expected that this is an enzyme-specific effect, related to physico-chemical properties of the
211 PceA reductive dehalogenase of *S. multivorans*. With the relative higher hydrophobicity of TBE
212 (log K_{ow} = 3.20, (Canton and Wegman, 1983)), similar to that of PCE (log K_{ow} = 3.40, (Hansch
213 et al., 1995)), compared to DBE (log K_{ow} = 1.76, (Hansch et al., 1995)) and TCE (log K_{ow} =
214 2.42, (Hansch et al., 1995)), this explanation is consistent with our results.

215 **3.3 Bromine stable isotope fractionation**

216 Similarly, bromine stable isotope fractionation was assessed (Table 1; SI Figure S2 & S4). ϵ_{Br}
217 was estimated at $-0.53 \pm 0.15\text{‰}$, $-1.03 \pm 0.26\text{‰}$, and $-1.18 \pm 0.13\text{‰}$ for *S. multivorans*
218 debrominating *trans*-1,2-DBE, *cis*-1,2-DBE and TBE, respectively, and -0.97 ± 0.28 , $-1.16 \pm$
219 0.36 , and -1.34 ± 0.32 for *D. hafniense* PCE-S debrominating *cis*-1,2-DBE, TBE and *trans*-1,2-
220 DBE, respectively. Notably, no or little variability between microorganisms or within the
221 debromination pathway was seen.

222 Generally, variability in bromine stable isotope fractionation between all assays was negligible.
223 After correcting for non-reacting bromines within the molecule, all values were similar to the
224 estimated Streitweiser limit at 1.002 (Table 1) with $AKIE_{Br} = 1.001 \pm 0.0003$, 1.002 ± 0.0005 ,
225 and 1.004 ± 0.0004 for *S. multivorans* debrominating *trans*-1,2-DBE, *cis*-1,2-DBE and TBE
226 respectively, and 1.002 ± 0.0006 , 1.003 ± 0.0006 , and 1.003 ± 0.0011 for *D. hafniense* PCE-S
227 debrominating *cis*-1,2-DBE, *trans*-1,2-DBE and TBE respectively, nor was there any indication
228 of a strong masking effect for bromine during debromination of TBE by *S. multivorans* as there
229 was for carbon in the same assay. $AKIE_{Br}$ for this reaction did in fact exceed the estimated
230 Streitweiser limit, as did $AKIE_{Br}$ for two other assays, but as all are approximations and are
231 within the same order of magnitude, the difference is negligible.

232 **3.4 Dual-element C and Br isotope analysis**

233 Carbon stable isotope fractionation was plotted as a function of bromine stable isotope
234 fractionation (Figure 1, Table 1), where linear regression slopes (expressed as $\Lambda \approx \epsilon_C/\epsilon_{Br}$)
235 represent the combination of kinetic isotope effects for each assay. Dual-element slopes were
236 estimated at 7.1 ± 1.6 , 8.3 ± 3.7 , and 8.9 ± 2.4 for *D. hafniense* PCE-S during debromination of
237 *trans*-1,2-DBE, TBE and *cis*-1,2-DBE respectively (Table 1). Distinctly different and more
238 variable values were estimated at 1.03 ± 0.2 , 17.9 ± 5.8 , and 29.9 ± 10.95 for *S. multivorans*
239 during debromination of TBE, *cis*-1,2-DBE and *trans*-1,2-DBE respectively (Table 1). Again,
240 this difference observed for TBE was similar for that for PCE, while DBE, behaved similar to
241 TCE. This variability in isotope fractionation pattern for TBE may be due to slight differences in
242 reaction mechanism, but, may also be due to different kinetics of the overall reaction, i.e. not C-
243 Br bond cleavage but another, preceding, step in the reaction determining the observed isotope
244 effect (Renpenning, 2014). In *S. multivorans*, the dehalogenation reaction mediated by PceA was
245 recently suggested to function via long-range electron transport, leading to carbon-halogen bond
246 cleavage, however, in case of TBE may be via an undirected electron transfer mechanism (Kunze
247 et al, 2017; Ye et al 2010).

248 **Discussion**

249 **4.1 Application of CSIA for assessment of in situ degradation of brominated organic** 250 **compounds**

251 Single-element, carbon stable isotope fractionation was previously reported for ethylene
252 dibromide (EDB) (Henderson et al., 2008) and bromine stable isotope fractionation for
253 brominated phenols (Bernstein, 2013), both for microbial dehalogenation. Dual-element carbon-
254 bromine stable isotope fractionation was reported for several abiotic and biotic transformation

255 reactions of EDB (Kuntze et al., 2016) and abiotic degradation of tribromoneopentyl alcohol
256 (TBNPA) (Kozell, 2015). Similarly, in our case significant carbon and bromine fractionation was
257 observed for all tested bromoethenes, supporting the utility of CSIA for evaluating
258 biotransformation of BOCs. Due to the observed variability in enrichment factors, a
259 complementary approach including the assessment of the degrading microorganisms *in situ* may
260 be necessary to allow a quantification of biodegradation applying the Rayleigh concept
261 (Meckenstock et al., 2004; Nijenhuis and Kuntze, 2016).

262 ***4.2 Evaluation of the debromination reaction using CSIA***

263 The ϵ_C values determined in this study, for what is assumed to be a stepwise, sequential
264 hydrogenolysis reaction via single-electron transfer ranged from $-7.5 \pm 1.4\text{‰}$ to $-16.5 \pm 2.8\text{‰}$,
265 with larger fractionation (when excluding the small *S. multivorans* TBE outlier) than that
266 reported for the debromination of EDB (Henderson et al., 2008; Kuntze et al. 2016), in which
267 reported ϵ_C were at $-5.6 \pm 1\text{‰}$ and $-5.3 \pm 0.5\text{‰}$ for ethylene dibromide (EDB) for biodegradation
268 in anaerobic microcosms (Henderson et al., 2008) and by *S. multivorans* (Kuntze et al., 2016),
269 respectively. Similarly, assuming a step-wise reaction for EDB, AKIE values for *cis*- and *trans*-
270 DBE were also higher compared to EDB dehalogenation by *S. multivorans* with $AKIE_C = 1.044$,
271 1.035 and 1.0107 (Kuntze et al., 2016), respectively. Similar differences were observed in
272 bromine isotope fractionation with $AKIE_{Br} = 1.002$, 1.001 and 1.0046 for *cis*- and *trans*-DBE
273 and EDB (Kuntze et al., 2016), respectively, supporting a difference in the reaction mechanism
274 between hydrogenolysis and dihaloelimination. The ϵ_{Br} values determined ranged from $-0.53 \pm$
275 0.15‰ to $-1.34 \pm 0.32\text{‰}$ and generally exceeded the values reported for microbial reductive
276 debromination of brominated phenols at $-0.20 \pm 0.06\text{‰}$ to $-0.76 \pm 0.08\text{‰}$ (Bernstein, 2013),
277 likely due to the difference in involved microbial community and enzymes as well as substrate

278 properties. Thus far, the described microorganisms and their reductive dehalogenases were
279 reported to be highly substrate specific, each dehalogenating a restricted set of organohalides
280 (Richardson, 2013). Structural differences in the enzyme structure may, therefore, affect the
281 reaction mechanism to a certain extent (Kunze, 2017).

282 **4.3 Rate-limitation and masking of isotope effects**

283 The extent of fractionation for debromination observed in the present study was within the order
284 of magnitude of that theoretically predicted by the semi-classical Streitweiser limit equation for
285 C-Br bond cleavage for all reactions with the exception of debromination of TBE by *S.*
286 *multivorans*. Here, carbon isotope fractionation was significantly lower at $\epsilon_C = -1.3 \pm 0.2\text{‰}$ than
287 for all other assays, including debromination of TBE by *D. hafniense* PCE-S at $\epsilon_C = -7.5 \pm 1.4\text{‰}$,
288 and was remarkably similar to the previously reported small ϵ_C for *S. multivorans*-mediated
289 reductive dehalogenation of PCE at -0.4‰ to -2.2‰ (Nijenhuis, 2005; Cichocka, 2007;
290 Renpenning, 2015). This extent of masking was not seen for bromine during debromination of
291 TBE by *S. multivorans*, with ϵ_{Br} evaluated at $-1.18 \pm 0.13\text{‰}$, an average value within all assays
292 which ranged from -0.53 ± 0.15 for *S. multivorans* debrominating *trans*-1,2-DBE to $-1.34 \pm$
293 0.32‰ for *D. hafniense* PCE-S debrominating *trans*-1,2-DBE. As fractionation patterns were
294 different for carbon than for bromine, the correlation between carbon and bromine fractionation
295 was variable, with Δ_{C-Br} ranging from $7.1 \pm 1.6\text{‰}$ to $8.9 \pm 2.4\text{‰}$ for *D. hafniense* PCE-S, and
296 from $1.03 \pm 0.2\text{‰}$ to $29.9 \pm 10.95\text{‰}$ for *S. multivorans*. This variability between strains and,
297 most pronouncedly for *S. multivorans* within the dehalogenation pathway, suggests differences
298 in the reaction between these substrates at a single enzyme, similarly to the previous
299 observations for the chlorinated ethenes (Cretnik, 2013; Wiegert, 2013; Renpenning, 2014).

300 **5. Implications for the application of CSIA for assessment of *in situ* biodegradation of BOC**

301 BOCs have gained the interest of policy makers and environmental researchers alike due to their
302 toxicity and suspected carcinogenic, endocrine disruptive, and neurodegenerative properties (de
303 Wit, 2002; Alaei, 2003), and have been restricted or even banned by regulatory agencies such
304 the U.S. Environmental Protection Agency (EPA) (Betts, 2008). However, due to their persistent
305 nature, as evidenced by detection in house dust, soil, and water samples, as well as in human
306 breast milk (de Wit, 2002), BOCs continue to pose a threat to human health and ecosystems, and
307 their environmental fate and transport should be elucidated. The use of stable isotope analysis to
308 detect and characterize transformation of BOCs *in situ* may be a valid option as fractionation has
309 been reported for both biotic and abiotic processes. And while much work is still needed, this
310 study now adds to the still quite small database of measurable and statistically significant carbon
311 and bromine isotope enrichment factors determined for microbial reductive debromination.

312

313 **Acknowledgements**

314 We thank Hans Richnow for discussion and Falk Bratfisch, Ursula Günther, and Matthias Gehre
315 for technical support during isotope analysis. This research has been financially supported by the
316 European Union under the 7th Framework Programme (project acronym CSI: ENVIRONMENT,
317 contract number PITN-GA-2010-264329) and by the Bundesministerium für Bildung und
318 Forschung (02WU1221; INTIME), Germany, and Ministry of Science and Technology, Israel
319 (BMBF-MOST).

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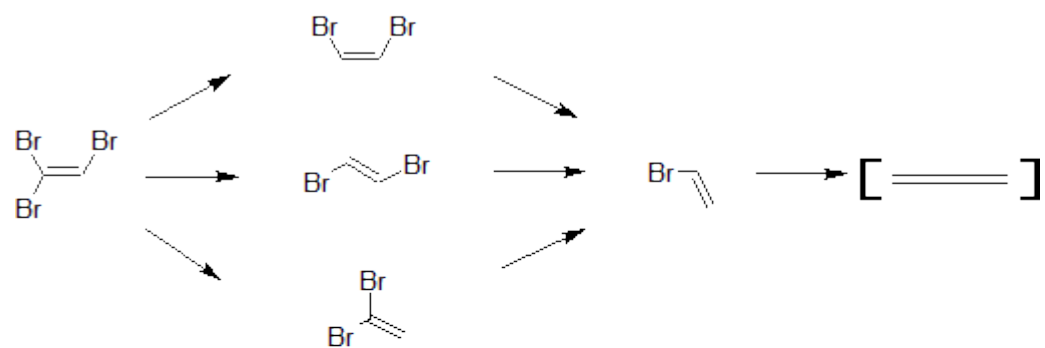
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Tables and Figures

Table 1. Carbon and bromine isotope fractionation during enzymatic reductive debromination of TBE, and *cis*- and *trans*-1,2-DBE expressed as enrichment factors (ϵ_C and ϵ_{Br}) and AKIE values assuming reactions are stepwise and secondary isotope effects negligible. Sample size is given as N. 2-D analysis was also performed and linear regression slopes are denoted Λ .

	<i>S. multivorans</i>				<i>D. hafniense</i> PCE-S					
	ϵ_C [‰]		R^2	N	AKIE _C	ϵ_C [‰]		R^2	N	AKIE _C
TBE	-1.2	±0.2	0.829	39	1.002 ±0.0004	-7.7	±1.5	0.799	30	1.016 ±0.0030
<i>cis</i> -DBE	-21.2	±1.6	0.937	48	1.044 ±0.0032	-14.5	±0.7	0.988	23	1.030 ±0.0007
<i>trans</i> -DBE	-16.8	±1.8	0.939	25	1.035 ±0.0036	-9.5	±1.2	0.870	42	1.019 ±0.0024
	ϵ_{Br} [‰]				AKIE _{Br}	ϵ_{Br} [‰]				AKIE _{Br}
TBE	-1.18	±0.13	0.937	27	1.004 ±0.0004	-1.16	±0.36	0.828	13	1.003 ±0.0011
<i>cis</i> -DBE	-1.03	±0.26	0.912	10	1.002 ±0.0005	-0.97	±0.28	0.777	18	1.002 ±0.0006
<i>trans</i> -DBE	-0.53	±0.15	0.911	9	1.001 ±0.0003	-1.34	±0.32	0.876	14	1.003 ±0.0006
	Λ					Λ				
TBE	1.0	±0.2	0.814	25		8.3	±3.7	0.711	12	
<i>cis</i> -DBE	17.9	±5.8	0.862	10		8.9	±2.4	0.797	17	
<i>trans</i> -DBE	29.9	±11.0	0.856	9		7.1	±1.6	0.885	14	



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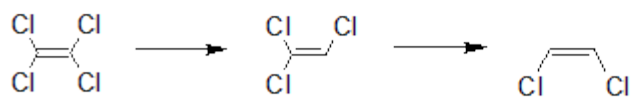


Figure 1. Enzymatic reductive debromination versus reductive dechlorination catalyzed by crude extracts of *Sulfurospirillum multivorans* and *Desulfitobacterium hafniense* PCE-S. Tribromoethene is sequentially debrominated to vinyl bromide via branched intermediates 1,2-*cis*- and *trans*-, and 1,1-dibromoethene, compared to the exclusive formation of *cis*-1,2-DCE as final product of dechlorination. Additionally, small amounts of ethene are produced when the debromination reaction is catalyzed by crude extracts of *D. hafniense* PCE-S.

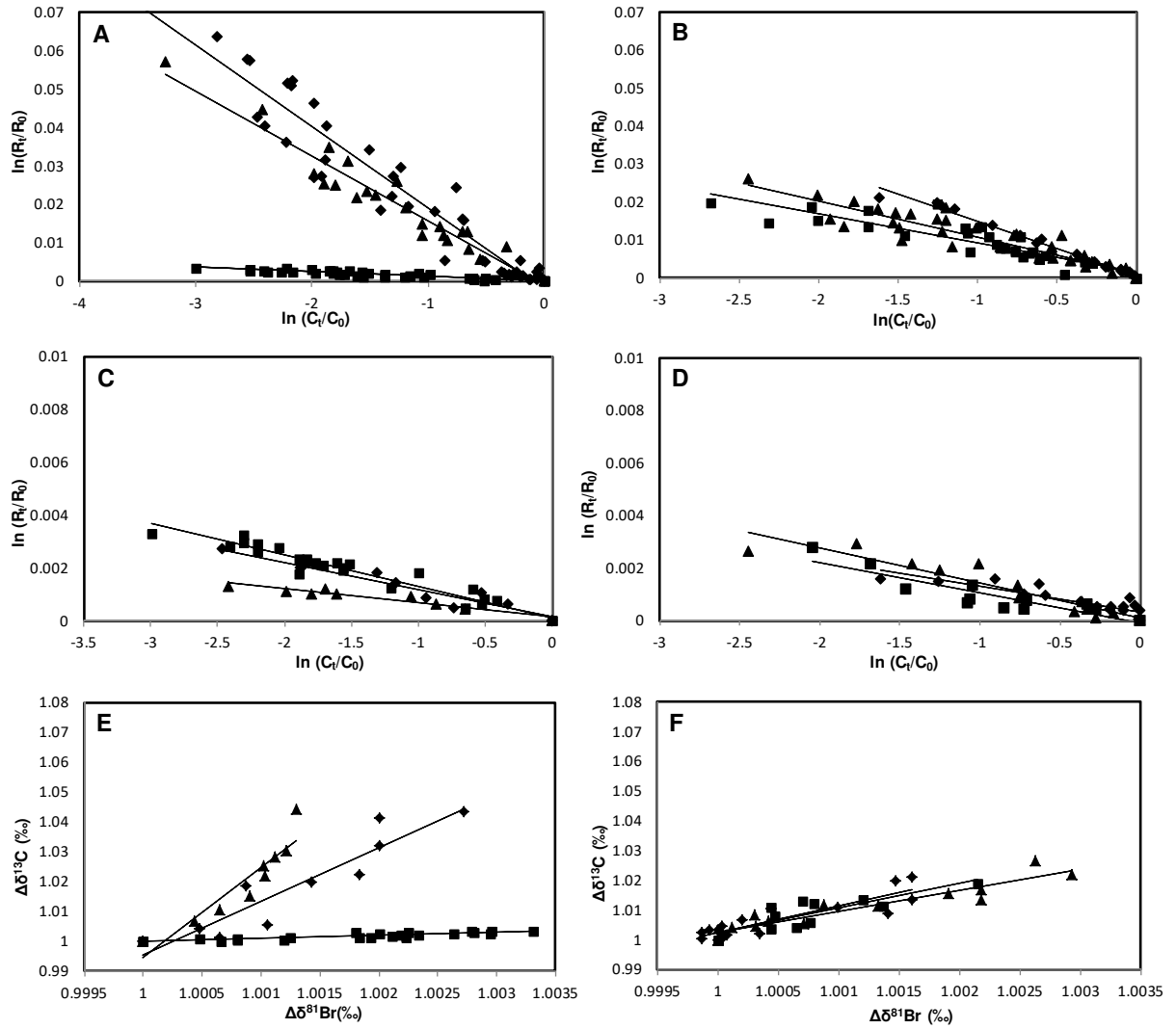


Figure 2: Rayleigh plots of carbon (A,B) and bromine (C,D) stable isotope fractionation and dual-element plots (E,F) for TBE (square), *cis*-DBE (diamond) and *trans*-DBE (triangle) dehalogenation by *S. multivorans* (A,C,E) and *D. hafniense* strain PCE-S (B,D,F).