1 Supporting information for:

2 Natural chlordecone degradation revealed by numerous transformation
3 products characterized in key French West Indies environmental
4 compartments

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41 S1 Supporting Methods

42 Chemicals and reagents

43 Chlordecone was obtained from Cluzeau Info Labo (C.I.L.) (Dr. Ehrenstorfer, reference C11220000, batch number 10725, and purity 96.7%). ¹³C-enriched CLD was obtained from 44 Azur Isotopes with a ${}^{13}C/{}^{12}C$ ratio estimated to 80%. Vitamin B₁₂ (\geq 96%), palladium on 45 carbon (12%), zinc powder, K_2CO_3 and Na_2S ($\geq 98\%$) were purchased from Sigma Aldrich. 46 Titanium (III) citrate was prepared from titanium (III) chloride (> 12% in HCl) (Sigma-47 Aldrich) and sodium citrate and neutralized with Na₂CO_{3[1]}. Micrometric Fe⁰ was obtained 48 from Dousselin (iron powder NC 100.24; Dousselin & Geoffray Jacquet Réunis, 69270 49 Couzon au Mont d'Or, France). Dichloromethane (DCM), heptane and pentane (HPLC grade) 50 51 were obtained from Fisher Chemical; cyclohexane (99.8%), ethyl acetate (EtOAc) (99.6%), chloroform (99.9%) and NMR solvents from Acros Organics; and acetonitrile (MeCN), 52 ethanol (EtOH) and methanol (MeOH) (LC-MS grade) from VWR Chemicals. H₂SO₄ (95-53 97%) was obtained from Fluka. Chemical products used for microbiological media were 54 55 obtained from Sigma Aldrich.

56 Analytical methods

57 GC-FID analyses

1 ml samples were collected, acidified to pH 3 with HCl (1 M) and extracted three times with dichloromethane (1:1 sample/DCM). After slow evaporation of the combined organic layers overnight, the crude residue was solubilized in an isooctane/acetone 85:15 mixture (350 μ l) and 3 μ l were injected in duplicate using split mode (1:3; 350°C) on a Thermo Scientific Trace 1300 connected to a flame-ionization detector (320°C). The instrument was equipped with a non-polar 20 m × 0.15 mm × 0.15 μ m TG-5Sil column (Thermo Fisher Scientific). The temperature program started at 150°C for 0.05 min, increased at a linear rate of 23°C/min
until 290°C and remained 0.6 min at 290°C.

66 GC-MS analyses

67 GC-MS 1

500 μL samples were collected, acidified to pH 1 with HCl (6 M) and extracted with dichloromethane (1:1 sample/DCM). The organic layer was then analyzed using a Thermo Fisher Focus GC coupled to a single-quadrupole mass spectrometer (Thermo Fisher DSQ II). The instrument was equipped with a non-polar 30 m × 0.25 mm × 0.25 μm DB-5MS column (Agilent J&W) and a split/splitless injector. Ionization conditions and GC program were described elsewhere[2, 3].

74 *GC-MS 2*

GC-MS analyses were performed using the instrument described in Chaussonnerie et al, 2016[2]. GC program started at 125°C (hold time 0.5 min), continued with 50°C min⁻¹ to 180°C, followed by 12.5°C min⁻¹ to 275°C (hold time 0.5 min). The study was done using the SIM (single ion monitoring) mode. Two ions were selected for CLD (m/z = 272; 237), B1 (m/z = 288; 253) and one for B2-3-4 (m/z = 217). Time ranges were also defined for these targeted ions.

81 LC-HRMS analyses

82 *LC-MS 1*

83 20 μ l samples were collected and mixed with 480 μ l of NH₄OAc buffer (10 mM; pH 7 84 adjusted with NH₄OH)/MeCN (38:10; V/V). The resulted mixture was filtered on 0.22 μ m 85 filters (Millex®-GV, Merck) and injected in a Dionex Ultimate 3000 LC system coupled to an 86 LTQ-OrbiElitemass spectrometer (Thermo Fisher Scientific) fitted with a heated electrospray 87 ionization source (HESI) operating in negative ionization mode. Voltage optimization was 88 described elsewhere[2]. The chromatographic separation was achieved using a Thermo Fisher Syncronis[™] C₁₈ column (50 mm length, 2.1 mm inner diameter, 1.7 µm particle size) and 89 90 carried out at 30°C with a flow rate of 0.5 mL/min using NH₄OAc buffer (10 mM; pH 7 91 adjusted with NH₄OH) as solvent A and MeCN as solvent B. The gradient started at 20% B 92 for 1 min, followed by a linear gradient at 100% B for 7 min and remained 2 min at 100% B. 93 The system returned to the initial solvent composition in 2 min and was re-equilibrated under 94 these conditions for 2 min.

95 *LC-MS 2*

96 LC-MS analyses were performed using the instrument, column and ionization parameters 97 described in material and methods section. The gradient started with a flowrate of 0.5 98 mL/min, at 12% B for 2 min, followed by a linear gradient at 20% B for 2 min, a second 99 linear gradient at 100% B for 6 min and remained 2 min at 100% B. The system returned to 100 the initial solvent composition in 4 min and was re-equilibrated under these conditions for 2 101 min. Carboxyindane references were purchased from Enamine Ltd. (1-1H-indane carboxylic 102 acid, 2-1H-indane carboxylic acid, 4-1H-indane carboxylic acid, 5-1H-indane carboxylic 103 acid), injected separately in LC-MS and mixed together.

104 Chloride analyses

Samples of 4 ml were collected, centrifuged (3050 x g), filtred (Millex®-GS, (0.22 μ m, Merck) and sent for chloride analysis to the French public analysis laboratory of Drome (LDA26).

108 **Purification methods**

109 Preparative Medium Pressure liquid Chromatography

- 110 Purification was carried out using a Combi Flash® Companion® (Teledyne ISCO) instrument
- 111 equipped with a 40 g prepacked silica gel column (GraceResolv Flash Silica, Grace Davison,
- 112 Discovery Sciences).
- 113 Preparative High Performance Liquid Chromatography (HPLC)
- 114 A Shimadzu LC-20 AP preparative HPLC system coupled with an ultraviolet/visible detector
- 115 (Shimadzu SPD-20AV) and an evaporative light scattering detector (Shimadzu ELSD-LT II)
- 116 at 70°C and 3.4 bar was used. Chromatographic separation was achieved using a preparative
- 117 HPLC column (X-Bridge Prep C18 5 μm OBD (150 x 19 mm) equipped with a security guard
- 118 column (Cartridge Holder, 19 x 10 mm) and a prefilter (Guard Cartridge 19 x 20 mm).
- 119 Nuclear Magnetic Resonance (NMR) analyses
- 120 Nuclear magnetic resonance spectroscopy (NMR) (¹H-NMR; ¹³C-NMR; ¹H-¹H-COSY; ¹H-
- 121 ¹³C-HSQC; ¹H-¹³C-HMBC) was conducted on a Bruker Advance Ultra Shield 600 MHz
- spectrometer (UEVE, Evry, France). ¹³C-enriched B1 was analyzed on a Bruker Advance III
 800 MHz spectrometer (ICSN, Gif-sur-Yvette).
- 124 Chemical protocols

125 Compound ¹³C-enriched B1

- 126 To a solution of ¹³C-chlordecone (${}^{13}C/{}^{12}C = 80$ %; 11 mg, 4.2 10⁻⁵ mol, 1 eq.) and vitamin B₁₂
- 127 (4 mg, 2.9 10⁻⁶ mol, 0.14 eq.) in degassed H₂O/MeOH 7:3 (25 mL), was added titanium(III)
- 128 citrate (5 mL, 3.2 10⁻⁴ mol, 15 eq.) basified to pH 12.7 with NaOH (3 M). Reaction mixture
- 129 was stirred under N₂ atmosphere at room temperature for 80 min and quenched by contact
- 130 with O₂. Extraction with pentane (5 x 30 mL) followed by concentration under reduced

- pressure gave rise to a white crude solid. After purification (similar to TP B1) ¹³C-enriched
 B1 (2.7 mg; 9.2 10⁻⁶ mol; 22%) was obtained as a white solid. NMR data are available in
 Figures S33-S34.
- 134 Transformation products D1, D2, D3 and D4

To a solution of chlordecone (100 mg, $1.9 \ 10^{-4}$ mol, 1 eq.) and vitamin B₁₂ (60 mg, $5.8 \ 10^{-5}$ 135 136 mol, 0.3 eq.) in degassed H₂O/MeOH 1:4 (125 mL), was added titanium(III) citrate (25 mL, 1.6 10⁻³ mol, 8.4 eq.) basified to pH 12.7 with NaOH (3 M). Reaction mixture was stirred 137 under N₂ atmosphere at room temperature for 4 h and, quenched by contact with O₂. 138 139 Extraction with pentane (5 x 250 mL) followed by concentration under reduced pressure gave 140 rise to a white crude solid. A first purification step was carried out through PLC (Merck, PLC Silica gel, 1 mm, F254, 20 x 20) (cyclohexane/EtOAc 9:1); D1-4 retardation factor was 141 0.53.D1-D4 were then purified using a preparative HPLC system. Elution was performed at a 142 143 flowrate of 25 mL/min using H₂O as solvent A and MeCN as solvent B. Elution started at 144 50% B for 8 min, followed by a linear gradient reaching 60% B within 4 min, followed by a 145 plateau within 43 min and a last linear gradient reaching 100% B within 7 min. Fractions 146 containing respectively D1-D2 and D3-D4 (from 39 to 43 and 50 to 53 min respectively) were pooled separately, extracted 3 times with pentane and concentrated under reduced pressure to 147 give the title compounds D1-D2 (3 mg; $9.7 \ 10^{-6}$ mol; 5%) and D3-D4 (1.5 mg; $5.4 \ 10^{-6}$ mol; 148 149 3%) as white solids. All NMR, GC-MS and LC-HRMS analyses for D1-D2 and D3-D4 are available in Figures 18-19, 22, 46-56. 150

- 151 Derivatization of C compounds
- To a solution of C1-C2 or C3-C4 (250 μ g, 9 10⁻⁷ mol) in ethanol or methanol (1 mL) was added concentrated sulfuric acid (10 μ L). The reaction mixture was stirred for 1 h and

- 154 quenched with K₂CO₃ (sat). Aqueous phase was extracted with CH₂Cl₂ and analysed in GC-
- 155 MS. D1-D2 was obtained from C1-C2, D3-D4 from C3-C4 in presence of methanol. E1-E2
- 156 was obtained from C1-C2, E3-E4 from C3-C4 in presence of ethanol (Figure S6).

157 Degradation of CLD in presence of DTT or zero-valent zinc

158 CLD degradation tests were performed as described in Ranguin et al., 2017[4]. To a solution of chlordecone (0.25 mg, $5.0 \ 10^{-7}$ mol, 1 eq.) in degassed ethanol (25 mL) were added either 159 1,4-dithiotreitol (231 mg, 1.5 10⁻³ mol, 3000 eq.) or zerovalent zinc (65.9 mg, 1.0 10⁻³ mol, 160 2015 eq.) and vitamin B_{12} (13.56 mg, 1.0 10⁻⁵ mol, 20 eq.). The reaction mixtures were carried 161 out in triplicate, in a glove box (N₂ 98 %, H₂ 2 %) at room temperature (rt) for 4 h. Aliquots 162 163 were acidified with HCl (6 M) to quench the reaction, filtered on 0.22 µm filters (33 mm, Millex-GV, Merck), then analyzed in LC-MS and GC-MS according to the protocol GC-MS 164 165 2 (Figure S7).

166 Degradation of CLD in presence of hematin sodium and dithionite or sodium sulfide or zero-167 valent iron

To a solution of hematin (31.2 mg, 4.9 10^{-5} mol, 5 eq.) in degassed sodium carbonate solution (0.1 M, 2.5 mL) was added sodium dithionite or sodium sulfide or zero-valent iron (220 mg, 1-3 10^{-3} mol, 100 eq.) or titanium (III) citrate basified to pH 12.7 with NaOH (3 M) (5 mL, 3.3 10^{-4} mol, 32 eq.), and degassed water (30 mL). The solution was then added to chlordecone (5.0 mg, 9.9 10^{-6} mol, 1 eq.). The reaction mixture was stirred under N₂ atmosphere at room temperature (rt) for several days; and followed by GC-MS and LC-MS analysis.

175 Degradation of CLD in presence of chlorophyll a and sodium sulfide

To a solution of sodium sulfide (220 mg, 1-3 10^{-3} mol, 100 eq.) in degassed water (30 mL) was added chlorophyll a (33.3 mg, 3.7 10^{-2} mol, 4 eq.). The solution was then added to chlordecone (5.0 mg, 9.9 10^{-6} mol, 1 eq.), and stirred under N₂ atmosphere at room temperature (rt) for several days; and followed by GC-MS and LC-MS analysis.

- 180 Degradation of CLD adapted from Jablonski et al., 1996[5]
- 181 To a solution of chlordecone (0.3 mg, 5.9 10^{-7} mol, 1 eq.) and vitamin B₁₂ (0.05 mg, 3.7 10^{-8} ,

182 0.06 eq) or rumen (0.05 mL), in degassed HEPES (50 mM, 5 mL) and ethylene glycol (0.8 183 mL), was added titanium (III) citrate basified to pH 12.7 with NaOH (3 M) (1.3 mL, 8.7 10^{-5} 184 mol, 145 eq.). The reaction mixture was stirred under N₂ atmosphere at rt for 15 min and at 50 185 °C for 4h. TLC was done on both reactions (Hexane/Acetone 3:1) and four spots were 186 observed under UV lights with respective retardation factors (0.87, 0.97, 0.14, 0). These 187 reactions were followed by GC-MS and LC-MS analysis.

188 Degradation of CLD according to Schrauzer & Katz, 1978[6]

To a solution of chlordecone (0.5 mg, 9.9 10^{-7} mol, 1 eq.) and vitamin B₁₂ (20 mg, 1.5 10^{-5} , 15 eq.) in degassed methanol (5 mL) was added acetoin (1.12 mL, basified to pH 12.7 with NaOH (3 M) (1.3 mL, 1.3 10^{-3} mol, 1000 eq.). The reaction mixture was stirred under N₂ atmosphere at room temperature (rt) for several days and followed by GC-MS and LC-MS analysis.

- 194 Pd/C catalyzed hydrodehalogenation of C compounds (adapted from Chevallier et al,195 2018)[3]
- 196 To a solution of C1-C2 or C3-C4 (1 mg, $3-4 \ 10^{-6}$ mol, 1eq.) in degassed EtOH/THF/H2O (8 :
- 197 2 : :1; 5 mL) was added palladium on carbon (10 % wt.; 1 mg; 9 10^{-7} mol, 0.25 eq.). The N₂
- 198 atmosphere of the round flask was removed by bubbling 3 volumes of H₂. The reactor was

then filled with H₂. The reaction was finally stirred for 3 hours under H₂ atmosphere; 3
samplings were performed (T0, T 1h, T 3h), and analyzed in LC-MS Orbitrap according to
the protocol LC-MS 2 (Figure S8).

202 Anoxic microbiological experiments

All bacterial cultures were performed at room temperature, anaerobically in a glove box (Unilab mBraun) under an N₂/H₂ (98/2; V/V) atmosphere in MM+ medium[2] (otherwise mentioned) containing Na₂S (0.4g/L) as reducing agent.

206 Chlordecone bacterial degradation in presence of Consortium 86 and *Citrobacter* sp. 86

207 Modifications of previously described MM+ medium[2] were the following: MgCl₂ and 208 NH₄Cl were replaced by MgS0₄ and (NH₄)₂SO₄, respectively. NaCl, CaCl₂, KCl were 209 removed. Two liters of freshly prepared medium were inoculated with active consortium 86 210 culture (1/100 V/V). After 8h of incubation, three glass bottles were filled up with 650 ml of 211 this growing bacterial culture and one filled up with medium as negative control. 162.5 µl of a 212 chlordecone solution (200 mg of chlordecone in 1 ml dimethylformamide) were added to 213 reach a final chlordecone concentration of 50 mg/L in each bottle. The same procedure was 214 carried out for Citrobacter sp. 86. Cultures were incubated and monitored during 250 days 215 using GC-FID, GC-MS, LC-HRMS and chloride analysis.

216 Chlordecone biodegradation in soil/liquid microcosms from Guadeloupe Island

Soil microcosm were prepared from 0.5 g dry weight (dw) of andosol or 0.5 g dw of nitisol sampling in Guadeloupe Island. Each sample was inoculated under oxygen-free nitrogen athmosphere containing 5% H₂, at 25 °C, in 12 mL M9 mineral medium or M9 mineral medium supplemented with vitamin B_{12} (2 mg/L). Duplicate samples were taken over 36 months, using a sacrificial approach. A microcosm pair (duplicate samples) was retrieved from each serie and analysed at t = 0, t = 10 months, and t = 36 months. Two negative control series were conducted for each serie (again in duplicate, but only for t = 0 and t = 36 months), and consisted of (a) soil samples incubated aerobically and (b) irradiated soil samples soil samples (30 ± 1.5 kGy).

Duplicate samples from t = 0, t = 10 and t = 36 months were first basified to pH 12 with NaOH (1M), extracted with pentane (6 x 15 mL) after vortex and decantation. Aqueous phases were then acidified to pH 1 with HCl (1 M) and extracted with CH₂Cl₂ (12 x 15 mL) after vortex and decantation. Organic layers were then pooled, concentrated *in vacuo* and analyzed in duplicate using GC-MS and LC-MS protocols (**Figure S3**, **Table S2Table S3**).

231 Chlordecone degradation in presence of Martinique soil/liquid microcosms and liquid cultures

5 g of soil samples (Andosol 914, Nitisol 918, Ferralsol 919) were incubated in 50 mL MM+
medium[2] and amended with chlordecone at 40 mg/L. After one month, 500 µl of each
Martinique liquid supernatant was used to inoculate 50 mL of medium MM+ with
chlordecone at 40 mg/L. Liquid cultures were monitored by GC-MS analysis (Figure S9).

236 Soil biodiversity analysis: 16S rRNA gene pyrosequencing and analysis

237 DNA extraction of 9 soils was performed on 10 g of sample using PowerMax® Soil DNA 238 Isolation Kit (MoBio) according to the manufacturer's instructions. V4-V5 regions of 16S 239 rRNA gene were amplified with the primers 515F GTGYCAGCMGCCGCGGTAA and 926R 240 CCGYCAATTYMTTTRAGTTT then DNA libraries were sequenced using 250 bp paired 241 end reads chemistry on a MiSeq Illumina sequencer. Illumina sequencing adapters and primer 242 sequences were removed from the reads by in-house-designed software based on the FastX 243 package. Reads were merged using usearch v9.2.64 -fastq mergepairs command[7]. The 244 merged reads were cleaned, dereplicated and clusterised using usearch v9.2.64 (-fastq filter, -

245 derep fullength and -cluster otus commands with default settings). The taxonomic assignation was performed on 200,000 sequences randomly selected with SortMeRNA 246 247 v2.1[8] Greengenes database gg 13 5 using (http://greengenes.secondgenome.com/downloads/database/13 5) or SILVA128 database[9] 248 249 with --id 0.97 and --coverage 0.97. 16S sequences of bacterial species included in the 250 consortia 86 and 92[2] were added to the databases. The OTU table was generated using 251 usearch v9.2.64 -usearch global command with -id 0.97.

252 Statistical methods

253 Estimation of CLD and B1 extraction efficiency in andosol and nitisol

Extractions of one andosol (914) and one nitisol (918) were followed to determine CLD and

255 B1 extraction efficiency. Samples were prepared as described before (4 g of crude sample and

256 15 mL of milliQ water acidified to pH 1) and were extracted with CH₂Cl₂ (12 x 15 mL). Each

extraction was injected in GC-MS (500 μL) (Figure S10, Table S5).

258 Calibration curves

Chlordecone (CLD), chlordecol (CLD-OH), A1, B1 concentrations were estimated through
GC-MS calibration curves (Figure SI). They were made using pure metabolites separately
diluted in Hexane/Acetone (4/1). C1-C2 and C3-C4 concentrations were estimated through
LC-MS calibration curves (Figure SI). They were made using pure metabolites separately
diluted in NH₄OAc buffer (10 mM; pH 7) /MeCN (4/1) (Figure S11).

Six successive dilutions enabled to build calibration curves with a concentration range from 0.01 to 25 mg/L. Each sample was injected in duplicate. B1, C1-C2, C3-C4 purity was estimated through NMR analysis (¹H) by addition of chloroform as internal standard. Linear regression was chosen as the calibration model. LOD (limit of detection) and LOQ (limit of
quantification) were assumed according the AFNOR NF-T-90-210 requirement.

$$LOD = \frac{y_0 + 3 \times \sigma_{y_0}}{a}$$
$$LOQ = \frac{y_0 + 10 \times \sigma_{y_0}}{a}$$

269 Where a is the slope of the calibration curve, y_0 the intercept and σ_{y0} the standard deviation on 270 the intercept (**Table S6**).

271 Repeatability and uncertainty of the method

Uncertainty of measure was assumed according to the NF ISO/CEI GUIDE 98-3 (2014) requirement. This uncertainty included three parameters: (1) the uncertainty of reproducibility of extraction $U_{reproducibility}$, (2) the uncertainty of GC-MS or LC-MS instrument $U_{instrument}$ and (3) the mean bias U_{bias} (corresponding to the difference between the calculated concentration and theoretical value) according to NF T 90-210 (2009) requirement (**Table S6**).

277
$$U_{reproducibility} = \sqrt{\frac{\sum_{i}^{n} (x_{i} - \bar{x})^{2}}{n-1}} \text{ with } n = 2$$

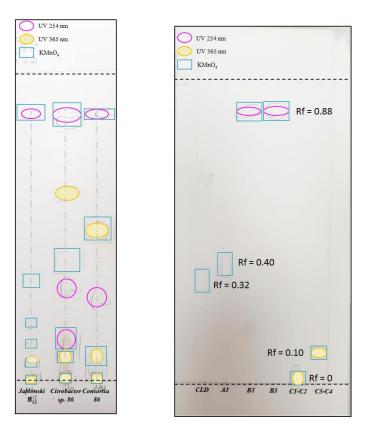
278
$$U_{instrument} = \sqrt{\frac{\sum_{i}^{n} (m_i - \bar{m})^2}{n-1}} \text{ with } n = 2$$

279 $U_{bias} = \sqrt{\frac{\sum_{i}^{n} b_{i}^{2}}{n}}$ with n = 5; and b_i, the difference between the calculated concentration with 280 the calibration model and the theoretical value

The uncertainty U was a function of the three previous uncertainties with a confidence levelof 68%.

$$U = \sqrt{U_{reproducibility}^{2} + U_{intrument}^{2} + U_{bias}^{2}}$$

284 S2 Supporting Figures



285

Figure S1. Thin layer chromatography of extract cultures from *Citrobacter* sp. 86 and Consortia 86 in (hexane : acetone 3:1). After three months in culture, 1L was acidified each time, extracted three times with DCM (3 x 1L) and concentrated *in vacuo*. Retardation factors were then compared with purified standards.

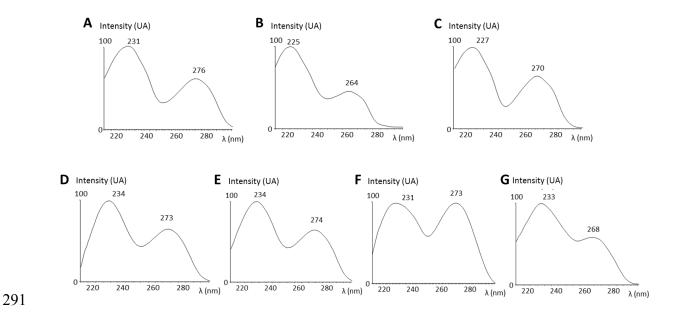
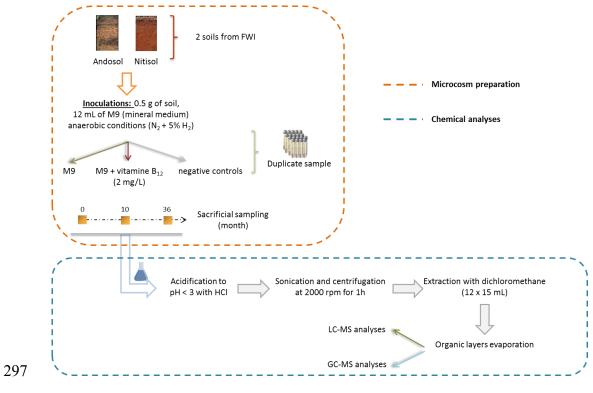
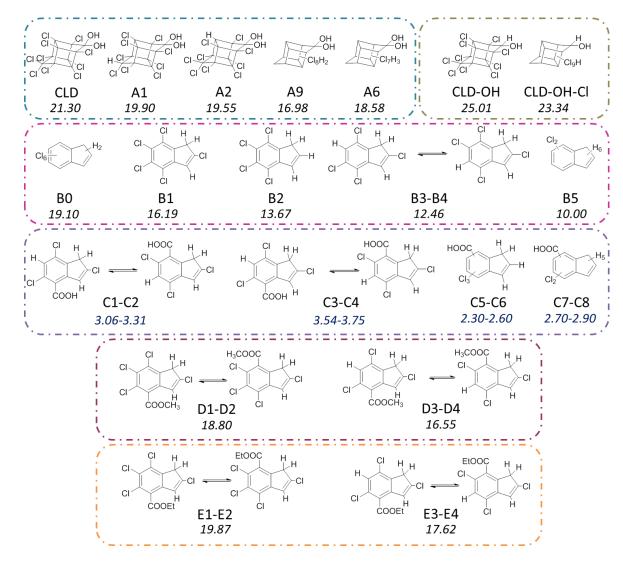


Figure S2. UV spectra of chlordecone bacterial metabolites extracted from liquid chromatography analysis. (A) UV spectrum of B1, (B) UV spectrum of B2, (C) UV spectrum of B3-B4, (D) UV spectrum of C1, (E) UV spectrum of C2, (F) UV spectrum of C3, (G) UV spectrum of C4

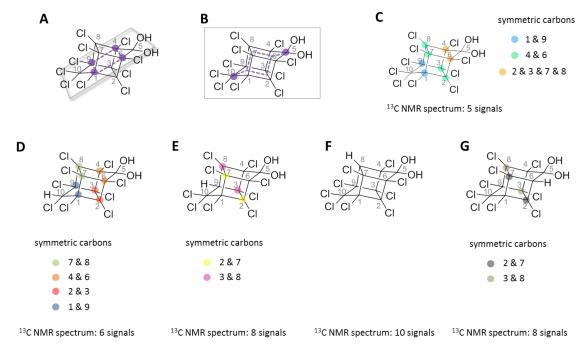


298 Figure S3. Microcosm preparation and extraction procedure enabling to find chlordecone TPs



300

Figure S4. Structure of the 29 chlordecone TP detected in environmental samples and/or in
biodegradation processes. GC-MS retention time are indicated in black, in italics, in minute.
LC-MS retention time are indicated in blue, in italics, in minute. Chromatographic methods
are described in material and methods section.



307 Figure S5. Chlordecone planes of symmetry and hydrochlordecone symmetric carbons according to the Hydrogen position. Number of expected signals in ¹³C NMR are also 308 309 indicated. (A) Chlordecone horizontal plane of symmetry, (B) Chlordecone vertical plane of 310 symmetry, Symmetric carbons for (C) chlordecone compared (D) 10to 311 monohydrochlordecone (E) 9 monohydrochlordecone (F) 8 monohydrochlordecone and (G) 312 6-monohydrochlordecone.

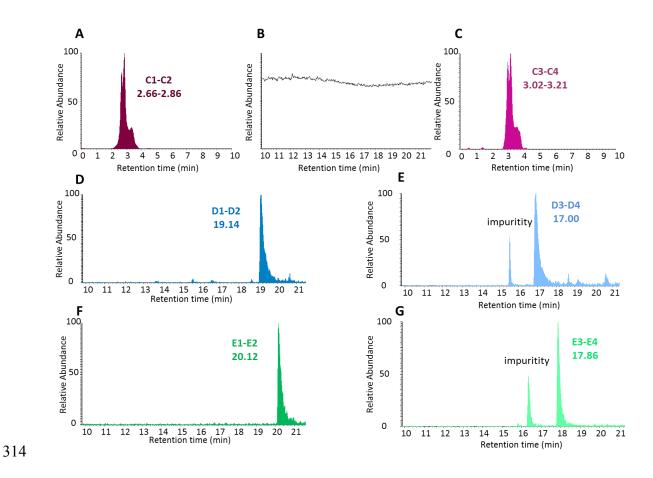
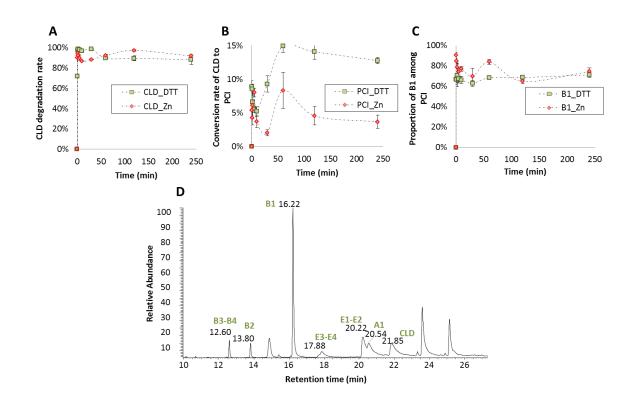


Figure S6. Derivatization of C compounds into D and E compounds and analysis of reactions by GC and LC-MS. Each figure represent extract ion chromatograms (A) LC-MS analysis of C1-C2 before reacting; (B) GC-MS analysis of C1-C2 and C3-C4 before reacting, no D or E compounds were present at this stage; (C) LC-MS analysis of C3-C4 before reacting, (D) GC-MS analysis of C1-C2 esterification in MeOH after 1h (E) GC-MS analysis of C3-C4 esterification in MeOH after 1h (F) GC-MS analysis of C1-C2 esterification in EtOH after 1h (G) GC-MS analysis of C3-C4 esterification in EtOH after 1h



323

Figure S7. Degradation of chlordecone with vitamin B_{12} and DTT or zero valent zinc, according to Ranguin et al., 2017 protocol. (A) Degradation rate of CLD with vitamin B_{12} and DTT (squares) or vitamin B_{12} and zerovalent zinc (diamonds); (B) Conversion rate of CLD to PCI (polychloroindene) calculated as PCI peak area (C) Proportion of B1 among PCI (B1, B2, B3, B4) calculated as B1 peak area/PCI peak area (D) GC-MS (full scan) analysis of DTT experiment after 4h (C1-C2 were also observed in LC-MS Orbitrap, negative mode)

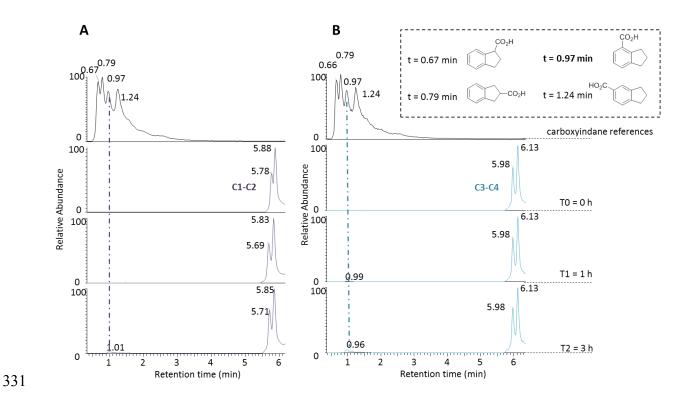
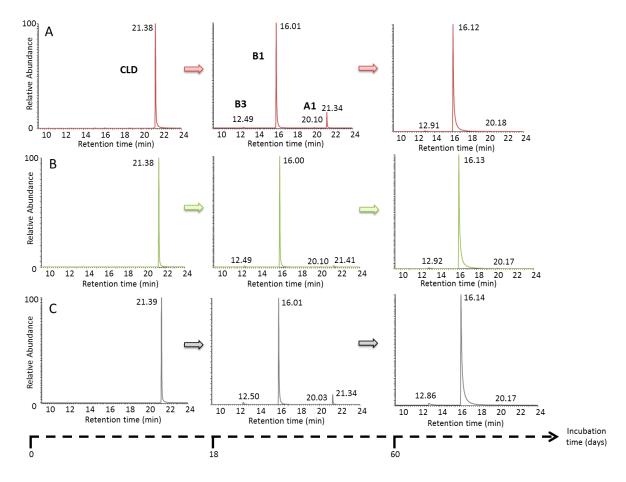


Figure S8. Hydrodehalogenation of C compounds and identification of the carboxyindane 332 333 transformation product by LC-MS-Orbitrap (A) LC-MS spectra of C1-C2 Pd-catalyzed 334 hydrodehalogenation over time (B) LC-MS spectra of C3-C4 Pd-catalyzed 335 hydrodehalogenation over time (ESI, negative mode). All chromatograms correspond to 336 extract ion chromatograms for C1, C2, C3, C4 and carboxyindane references.



338

Figure S9. Liquid cultures from three soils (A) 914, (B) 918, (C) 919 supplemented with MM+ medium (Chaussonnerie et al., 2016), sodium sulfide as reding agent and chlordecone. Three samplings were taken for each soil at t = 0 day, t = 18 days and t = 60 days, extracted with isooctane and analyzedby GC-MS analysis.

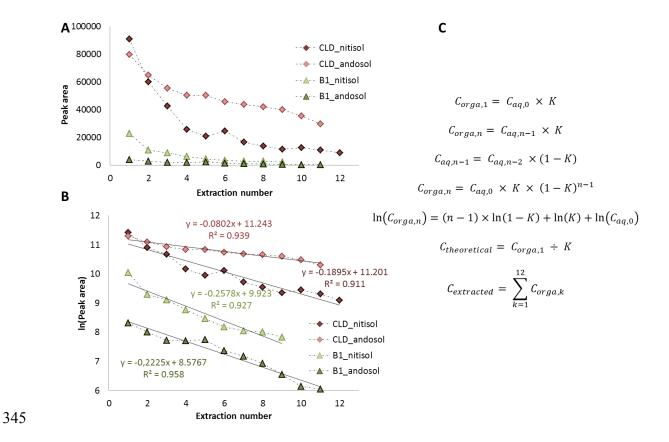


Figure S10. Estimation of CLD and B1 extraction efficiency by GC-MS analysis of each 346 347 extraction fraction (A) CLD and B1 peak area obtained from GC-MS analysis as a function of extraction number and type of soil. (B) logarithm of CLD and B1 peak area as a function of 348 349 extraction number (C) Equations used to assume extraction efficiency of both metabolite; 350 with $C_{x,y}$ the concentration of CLD or B1 in organic (orga) or aqueous (aq) layer at nth extraction; K the partition coefficient and n the final number of extraction (12). These 351 352 equations were assessed considering that organic and aqueous layers contained the same 353 volume, thus the extracted fraction of B1 and CLD could be compared to the theoretical 354 concentration.

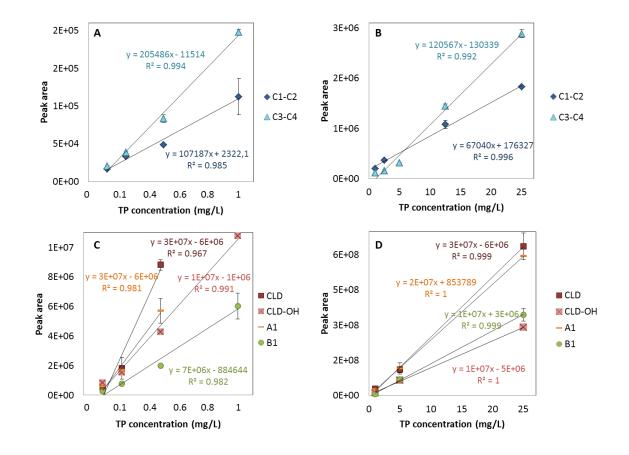
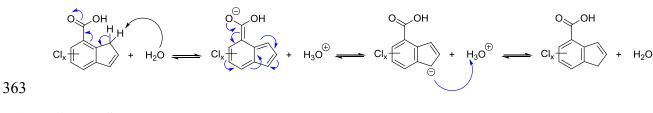
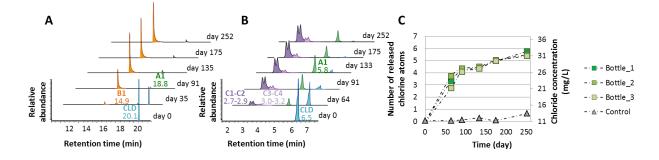


Figure S11. Calibration curves of CLD, CLD-OH, A1, B1, C1-C2 and C3-C4. (A) Calibration curve for 0.25-1 mg/L concentration range of C1-C2 and C3-C4 using LC-MS Orbitrap; (B) Calibration curve for 1-25 mg/L concentration range of C1-C2 and C3-C4 using LC-MS Orbitrap; (C) -Calibration curve for 0.25-1 mg/L concentration range of CLD, CLD-OH, A1 and B1 using GC-MS; (D) Calibration curve for 1-25 mg/L concentration range of CLD, CLD-OH, A1 and B1 using GC-MS.

355



364 Figure S12. Proposed keto-enol mediated isomerization for polychloroindene carboxylic365 acids



367

368 Figure S13. TP production during chlordecone degradation by consortium 86 during 250 369 days examined by GC-MS analysis, LC-MS analysis and chloride concentration. (A) GC-MS 370 monitoring of chlordecone degradation and detection of CLD (blue), A1 (green), B1 (orange) 371 and B3 (yellow) TP. (B) TP production during chlordecone degradation by APW using LC-372 MS-Orbitrap analysis and detection of CLD (blue), A1 (green), C1 (purple), C2 (purple), C3 373 (magenta) and C4 (magenta) metabolites. Extract ion chromatograms for quasimolecular ion 374 [M-H]- of m/z = 506.6797; 472.7187; 296.8852; 260.9271 (C) Chloride concentration 375 monitoring during chlordecone degradation. The average number of released chlorine atoms 376 per consumed chlordecone molecule is indicated.

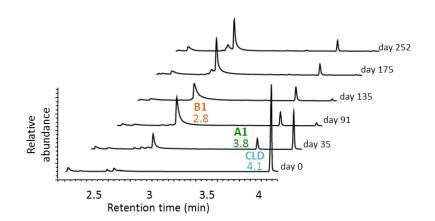


Figure S14. TP production during chlordecone degradation by consortium 86 during 250
days examined by GC-FID analysis.

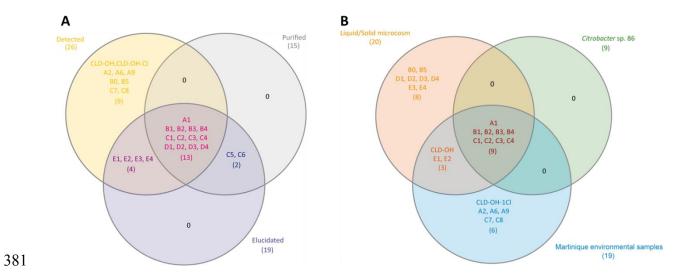


Figure S15. Venn diagrams representing the distribution of each observed TP. (A) Venn diagrams representing detected, purified and structure elucidated TP. (B) Venn diagrams distinguishing TP detected in liquid/solid microcosms, produced by Citrobacter sp. 86 (or consortium 86) and observed in Martinique environmental samples.

387 S3 Supporting Tables

Metal complex	Vitamin B ₁₂ ^{vi}	Hematin ""	Chlorophyll a	Rumen ^{vi}	Without		
Reductant							
Acetoin ⁱ <i>(Methanol)</i>	A, B, C, D (week)	/	/	/	X (week)		
Sodium dithionite ⁱⁱ	Traces of A (week)	Traces of A (days)	/	/	X (week)		
Sodium sulfide	A1, B1, C (days)	A1, B1, C (days)	Traces of A (days)	/	X (days)		
1,4-dithiothreitol <i>(Ethanol)</i> ⁱⁱⁱ	A, B, C, E (hours)	/	/	/	X (hours)		
Zero valent iron (<i>H</i> ₂ <i>O</i> -acetone) ^{iv}	A, B, C (week)	A (weeks)	/	/	A (weeks)		
Zero valent zinc <i>(Ethanol)</i> ⁱⁱⁱ	A, B, C, E (hours)	/	/	/	A, B, C, E (week)		
Titanium citrate ^v	B, C (hours)	A (hours)	/	Traces of A (hours)	A (hours)		
Titanium citrate <i>(H₂O-methanol)</i>	B, C, D (hours)	/	/	/	A (hours)		
Titanium citrate (<i>H</i> ₂ O-ethanol)	B, C, E (hours)	/	/	/	A (hours)		

388 Table S1. Screening of CLD transformation chemical conditions. Green cases meaning that 389 the transformation was complete, orange cases that the transformation was partial and red 390 cases that no transformation occurred. The slash / means that combinations have not been 391 tested, and the green boxes show conditions retained for chemical synthesis of A1, B C, D 392 and E compounds. Typical reaction time is indicated in brackets. Without reductant, no CLD 393 transformation occurred. The solvent generally used is water otherwise indicated in brackets. 394 This screening of degradation conditions has been established according to several already 395 published protocols (i) Schrauzer & Katz 1978[6]; (ii) Holmstead et al., 1976[10]; (iii) 396 Ranguin et al., 2017[4] (iv) Belghit et al., 2015[11]; (v) Chevallier et al., 2018[3]; (vi) 397 Jablonski et al., 1996[5].

Со	nditions	Sample number	Incubation time (month)	B5	B3-B4	B2	B1	B0	D1-D2	D3-D4	E1-E2	E3-E4	CLD-OH	CLD	A1
		57N	0	-	++	++	++	nd	+	+	nd	nd	+		nd
	_	65N	10	-	++	++	++	nd	+	+	nd	nd	+	nd	nd
So		69N	36		+	++	+++	-	+	-	nd	nd	+	nd	nd
Nitisol	Vitamin	71N	0	nd	+	++	++	-	+	+	nd	nd	+		nd
-		79N	10	nd	-	nd	+	nd	nd	nd	nd	nd	+	nd	nd
	B ₁₂	83N	36	-	++	+++	+++	+	++	+	-	+	-	nd	nd
		1A	0	nd	-	nd	+	nd	nd	nd	nd	nd	+	-	nd
-	_	9A	10	nd	+	nd	+	nd	nd	nd	nd	nd	+	-	
SS		13A	36	nd	+	nd	+	nd	nd	nd	nd	nd	+		nd
Andosol	Vitamin	15A	0	nd	nd	nd	-	nd	nd	nd	nd	nd	+		nd
4		23A	10	nd	+	nd	+	nd	nd	nd	nd	nd	+		nd
	B ₁₂	27A	36	nd	+	nd	+	nd	nd	nd	nd	nd	+	nd	
	$\boxed{\text{nd}} < 510^4 - 510^4 - 410^5 \qquad - 510^5 - 410^6 \qquad + 510^6 - 410^7 \qquad + + 510^7 - 410^8 \qquad + + + 510^8 - 410^9$														

Table S2. Analysis by GC-MS of soil/liquid microcosms. No quantification was done on this

401 set of data, only peak area from full scan mode is represented here. (nd means not detected)

Conditions		Sample number	Incubation time (month)	C1-C2	C3-C4	A1	CLD
		57N	0	nd	nd	nd	+
	_	65N	10	nd	nd	-	-
<u>s</u>		69N	36	+	++	-	-
Nitisol		71N	0	nd	nd	++	++
	Vitamin P	79N	10	nd		nd	nd
	B ₁₂	83N	36	nd	nd	nd	nd
		1A	0	-	nd	-	++
_	_	9A	10	nd	+	-	-
Andosol		13A	36	nd	nd	-	-
And	\/!+!	15A	0	nd	nd	-	++
	Vitamin B	23A	10	nd	+	-	-
	B ₁₂	27A	36	nd	nd	+	+
< 1	0 ² -	- 10 ² - 1	LO ³ - 1	0³ - 5 10	³ +	5 10 ³	³ - 10 ⁴

Table S3. Analysis by LC-MS (extract ion chromatograms, negative mode) of soil/liquid
405 microcosms. No quantification was done on this set of data, only peak area is represented
406 here. (nd means not detected)

Sample number	Туре	Origin	Plot	GPS N	GPS W
number					
914	Andosol	Maison rouge	Cacao 1	N 014°46.038	W 061°08.356
915	Andosol	Maison rouge	Cacao 2	N 014°45.802	W 061°08.323
918	Nitisol	Trinité	Montvert 3	N 014°41.626	W 060°57.507
920	Nitisol	Trinité	Montvert 4	N 014°41.774	W 060°57.505
919	Ferralsol	Trinité	Montvert 2	N 014°41.652	W 060°57.296
921	Ferralsol	Trinité	Montvert 1	N 014°41.635	W 060°57.154
922	Ashes and pumice stone	Basse pointe	H. Leyrits	N 014°37.272	W 060°58.112
923	Ashes and pumice stone	Basse pointe	H. Leyrits	N 014°50.598	W 061°07.258
926	Nitisol	Lamentin	R.LZ.Belème	N 014°39.924	W 061°00.361
927	Bed sediments	Trinité	Mangrove	N 014°43.250	W 060°56.462
928	Bed sediments	Trinité	Mangrove	N 014°43.260	W 060°56.480
929	River water	G. Trinité	Galion	N 014°42.008	W 060°57.067
930	River water	G. Trinité	Galion	N 014°42.008	W 060°57.067
931	Mangrove water	Trinité	Mangrove	N 014°43.250	W 060°56.462
932	Mangrove water	Trinité	Mangrove	N 014°43.260	W 060°56.480

Table S4. Environmental samples used for chlordecone TP investigation.

	Andosol	Nitisol
CLD B1	52 % 94 %	64 % 65 %
DI	94 %	05 %

Table S5. Estimated extraction efficiency of CLD and B1 in andosol and nitisol.

	CLD	A1	CLD-OH	B1	C1-C2	C3-C4
U _{bias}	33%	49%	13%	18%	28%	25%
LOD (m/L)	0.12	0.12	0.16	0.80	0.04	0.10
LOQ (mg/L)	0.66	0.65	0.85	0.15	0.21	0.50

Table S6. Mean bias, LOD (limit of detection) and LOQ (limit of quantification) of main

414 chlordecone TPs.

Sample type	Sample	CLD	CLD-OH	A1	B1	C1-C2	C3-C4	Other detected TP
	number							
Andosol	914	0.11 ± 0.03	0.03 ± 0.01	0.28 ± 0.22	2.07 ± 1.84	nd	0.01 ± 0.004	A6, A9, A2, Monohydro-CLD-OH, B2, B3-B4
Andosol	915	0.15 ± 0.04	0.04 ± 0.01	0.07 ± 0.06	5.08 ± 2.20	0.45 ± 0.18	0.99 ± 0.50	A6, A9, A2, B2, B3-B4, C7-C8
Nitisol	918	0.04 ± 0.01	0.04 ± 0.01	nd	0.06 ± 0.01	0.01 ± 0.001	1.20 ± 1.18	Monohydro-CLD-OH, B2, B3-B4
Nitisol	920	0.05 ± 0.01	0.04 ± 0.01	nd	0.04 ± 0.01	nd	0.01 ± 0.002	B2, B3-B4
Ferralsol	919	0.08 ± 0.01	0.07 ± 0.01	nd	0.04 ± 0.01	0.02 ± 0.01	0.01 ± 0.002	<i>B3-B4</i>
Ferralsol	921	0.03 ± 0.01	0.03 ± 0.01	nd	0.17 ± 0.13	nd	nd	A9, B3-B4
Ashes and	922	0.03 ± 0.01	0.05 ± 0.01	nd	0.04 ± 0.01	nd	nd	
pumice stone								
Ashes and	923	0.03 ± 0.01	0.03 ± 0.01	nd	0.12 ± 0.07	0.04 ± 0.02	0.30 ± 0.29	A9, B2, B3-B4
pumice stone								
Nitisol	926	nd	nd	nd	nd	nd	nd	B3-B4
Bed	927	nd	nd	nd	0.02 ± 0.01	nd	nd	B3-B4
sediments Bed	928	nd	nd	nd	0.02 ± 0.01	nd	nd	
sediments								
River water*	929	1.02 ± 0.35	nd	nd	1.81 ± 0.32	nd	nd	A6, A9, B2, B3-B4, E1-E2
River water*	930	0.44 ± 0.15	nd	nd	0.12 ± 0.02	0.04 ± 0.01	nd	B2, B3-B4
Mangrove	931	0.22 ± 0.20	nd	nd	0.13 ± 0.03	nd	nd	B3-B4
water*								
Mangrove water*	932	nd	nd	nd	0.14 ± 0.03	nd	nd	B3-B4

Table S7. Es	stimation of	chlordecone	e TP c	concentration	in enviro	nmental	samples	from Martinique.	Other ch	lordecon	e nd m	eans "not
detected".	*means	that	result	s are	reported	l in	μg/L	otherwise	they	are	in	mg/kg

S4 Supporting Mass Spectra

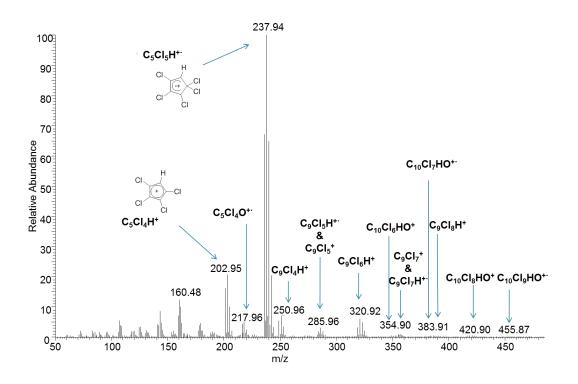


Figure S16. Mass spectrum analysis of A1 by GC-MS

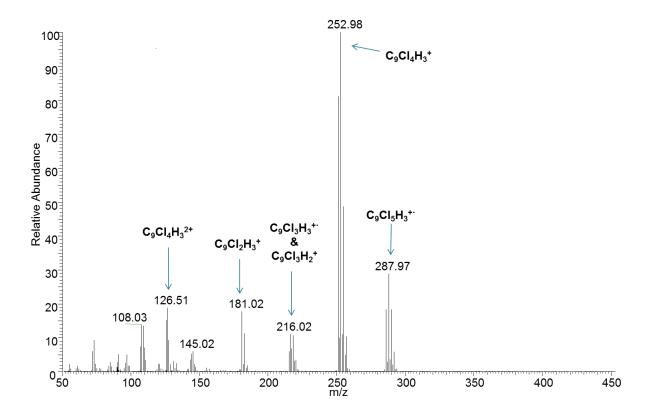


Figure S17. Mass spectrum analysis of B1 by GC-MS

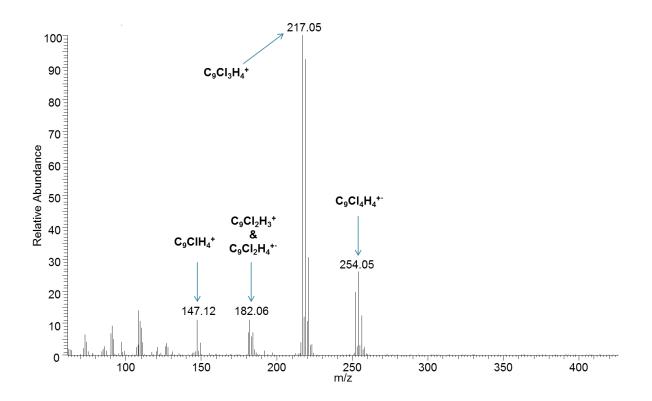


Figure S18. Mass spectrum analysis of B2 and B3-B4 by GC-MS

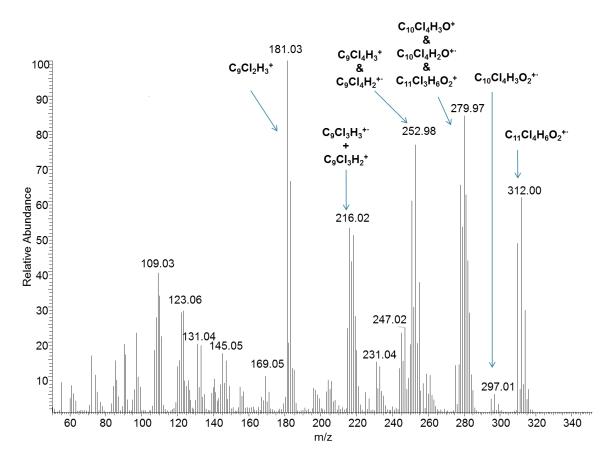


Figure S19. Mass spectrum analysis of D1-D2 by GC-MS

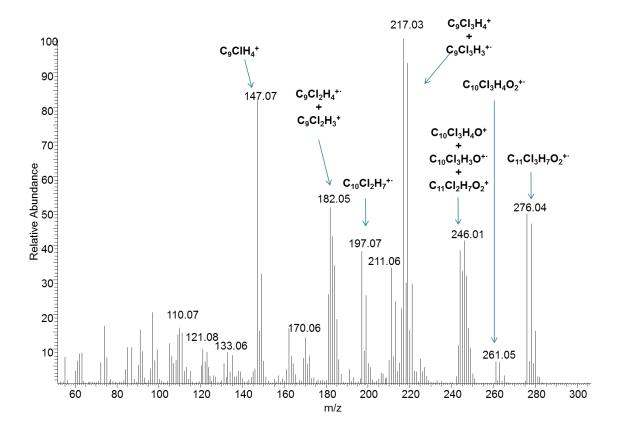


Figure S20. Mass spectrum analysis of D3-D4 by GC-MS

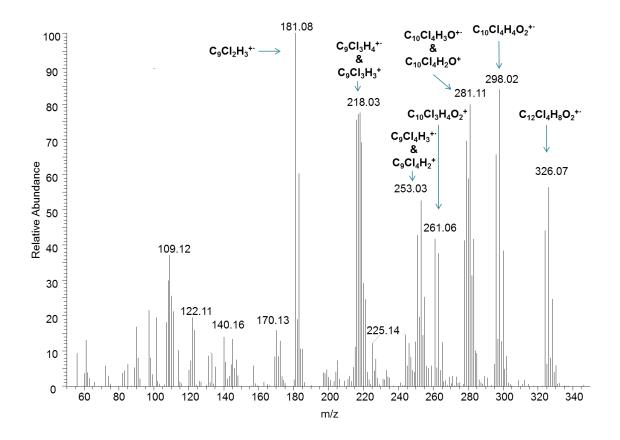


Figure S21. Mass spectrum analysis of E1-E2 by GC-MS

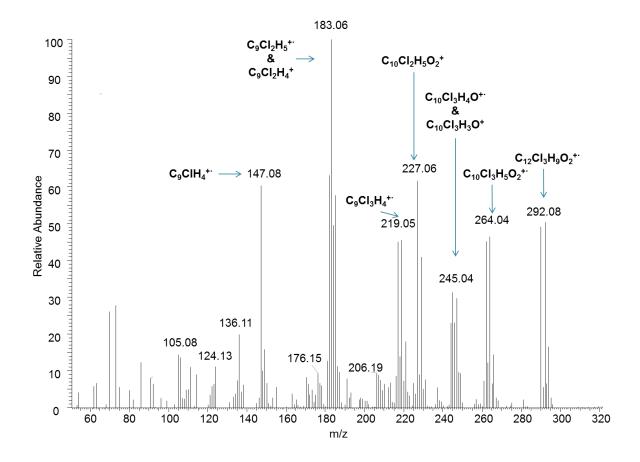


Figure S22. Mass spectrum analysis of E3-E4 by GC-MS

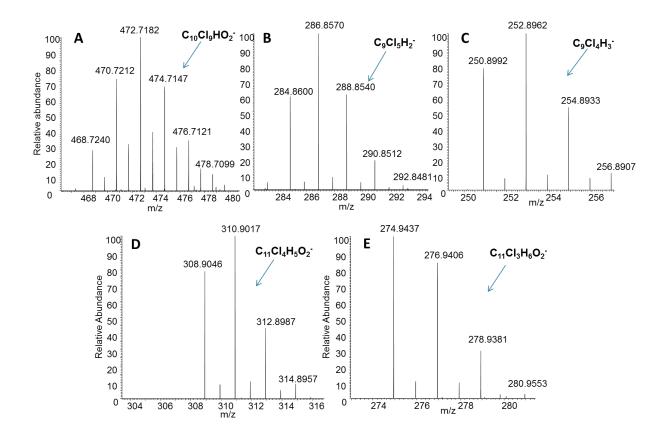


Figure S23. Mass spectra analysis of (A) A1, (B) B1, (C) B2 and B3-B4, (D) D1-D2 and (E) D3-D4 by LC-MS Orbitrap (negative mode)

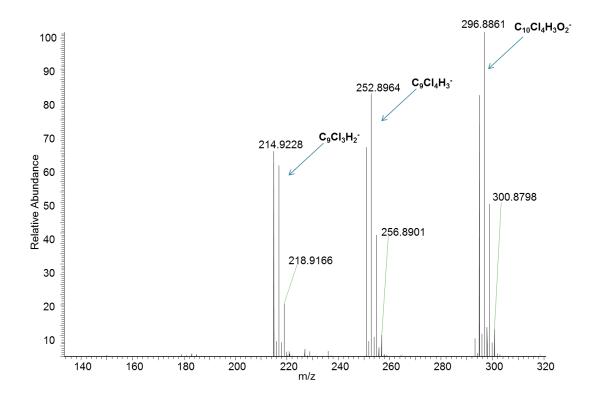


Figure S24. Mass spectrum analysis of C1-C2 by LC-MS Orbitrap (negative mode)

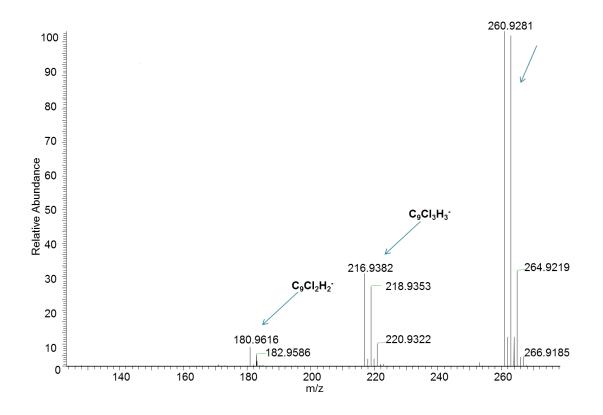


Figure S25. Mass spectrum analysis of C3-C4 by LC-MS Orbitrap (negative mode)

S5 Supporting NMR Spectra

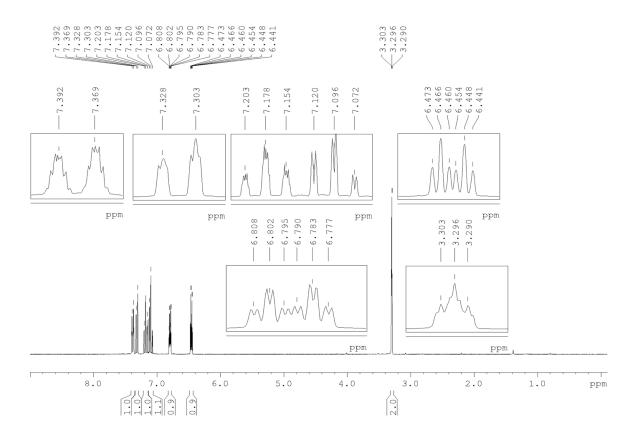


Figure S26. ¹H-NMR spectrum of indene in CDCl₃ (300 MHz)

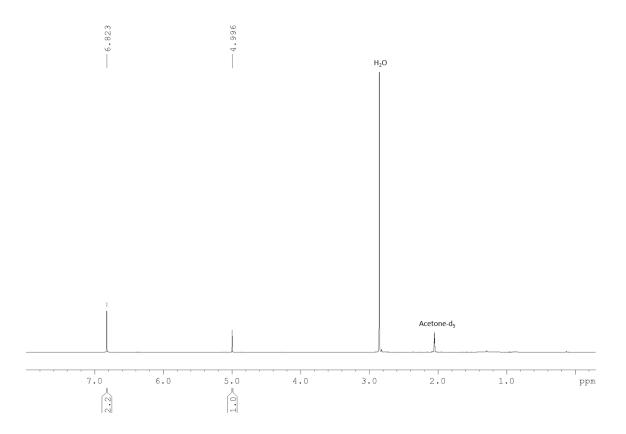


Figure S27. ¹H-NMR spectrum of A1 in $(CD_3)_2CO$ (600 MHz). ¹H NMR (600 MHz; $(CD_3)_2CO$): δ 6.85 (s, 2 H, C₅- $(OH)_2$), 5.00 (s, 1 H, C₁₀-*H*). [M-H]⁻ calcd. for C₁₀Cl₉O₂H₂, 468.7246; found, 468.7242. TLC: Rf (hexane : acetone 3/1) = 0.32

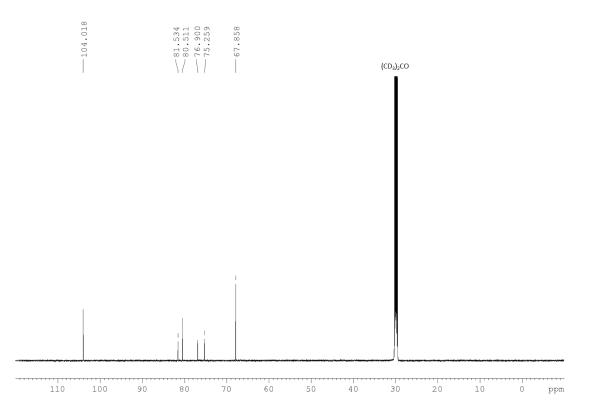


Figure S28. ¹³C-NMR spectrum of A1 in $(CD_3)_2CO$ (600 MHz). ¹³C NMR (150 MHz; $(CD_3)_2CO$): δ 103.0 (C₅), 81.5 (C₄₋₆), 80.5 (C₁₋₉), 76.9 (C₇₋₈), 75.2 (C₂₋₃), 67.8 (C₁₀).

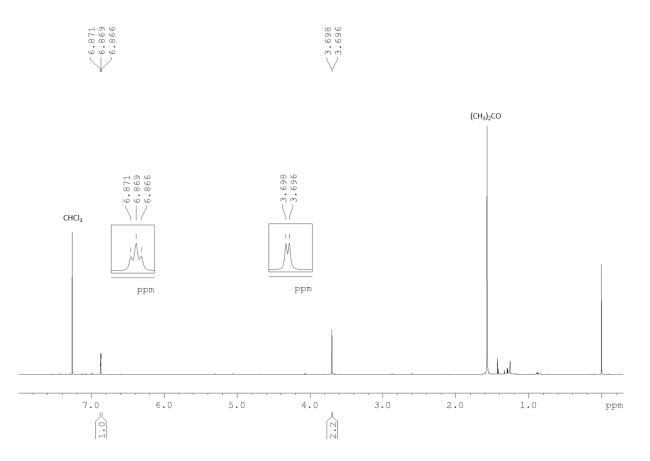


Figure S29. ¹H-NMR spectrum of B1 in CDCl₃ (600 MHz)

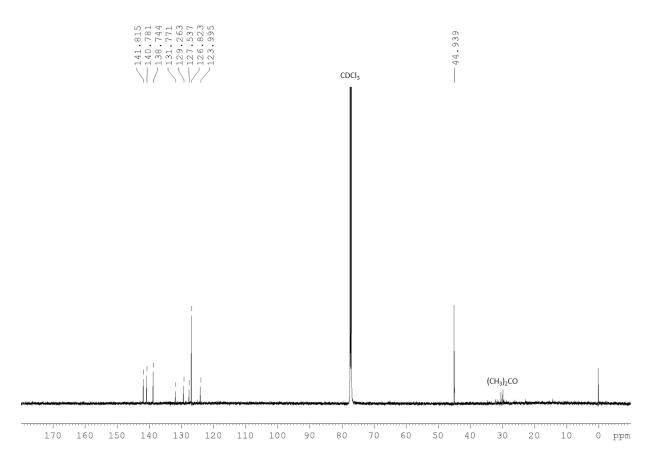


Figure S30. ¹³C-NMR spectrum of B1 in CDCl₃ (600 MHz)

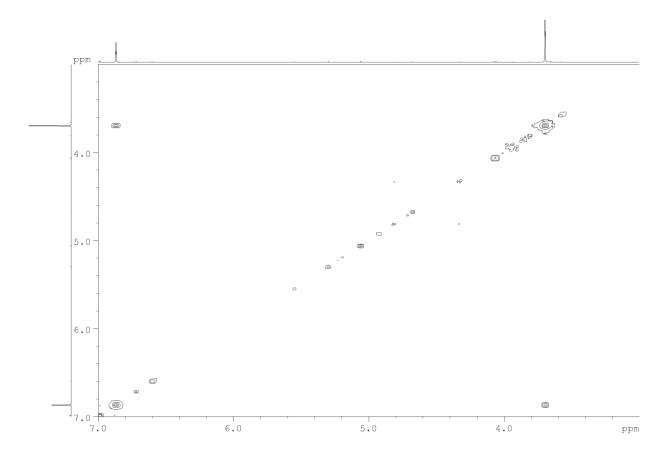


Figure S31. COSY spectrum of B1 in CDCl₃ (600 MHz)

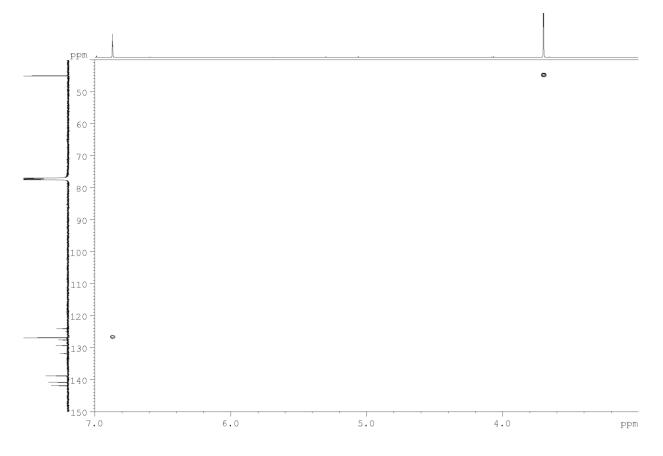


Figure S32. HSQC spectrum of B1 in CDCl₃ (600 MHz)

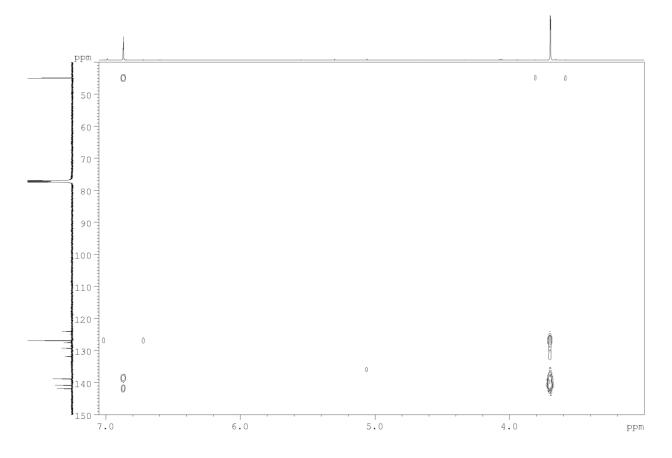


Figure S33. HMBC spectrum of B1 in CDCl₃ (600 MHz)

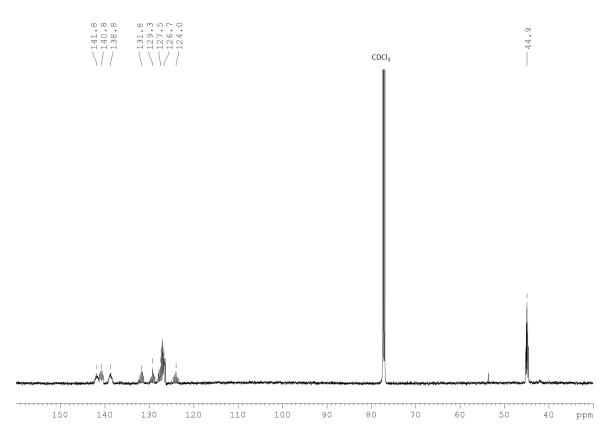


Figure S34. ¹³C-NMR spectrum of ¹³C-B1 in CDCl₃ (600 MHz)

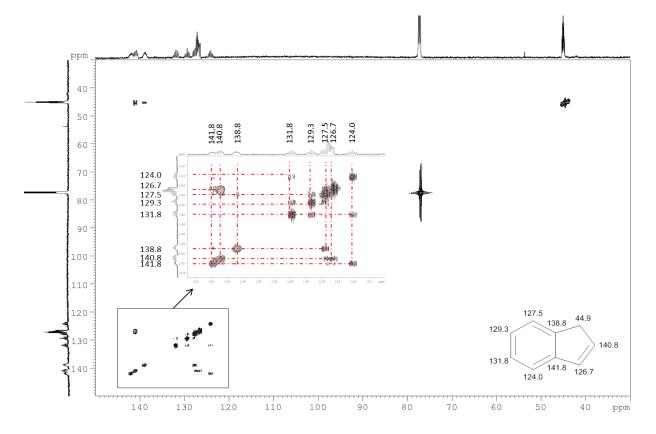


Figure S35. ¹³C-¹³C COSY spectrum of ¹³C-B1 in CDCl₃ (800 MHz). ¹³C NMR (150 MHz; CDCl₃): δ 141.9 (C₉), 140.9 (C₂), 138.8 (C₈), 131.9 (C₅), 129.4 (C₆), 127.6 (C₇), 127.0 (C₃), 124.1 (C₄), 45.0 (C₁). HRMS (m/z): [M-H]⁻ calcd. for C₂[13]C₇Cl₅H₂, 291.8828; found, 291.8838.

	CI 6 7 8 1 5 4 19 CI		_	COSY HMBC			
#	¹³ C(∂)	HSQC	¹H(<i>δ</i>)	J Coupling (Hz)	Protons	НМВС	COSY
1	44.9	CH_2	3.70	d, 1.4	2H	C2, C3, C8	H-3
2	140.8						
3	126.8	СН	6.87	t <i>,</i> 1.4	1H	C1, C4, C8, C9	H-1
4	124.0						
5	131.8						
6	129.8						
7	127.5						
8	138.7						
9	141.8						

Table S8. NMR data assignment of B1 in $CDCl_3$ HRMS (m/z): [M-H]⁻ calcd. for $C_9Cl_5H_2$, 284.8594; found, 284.860. TLC: Rf (cyclohexane : ethyl acetate 9/1) = 0.78

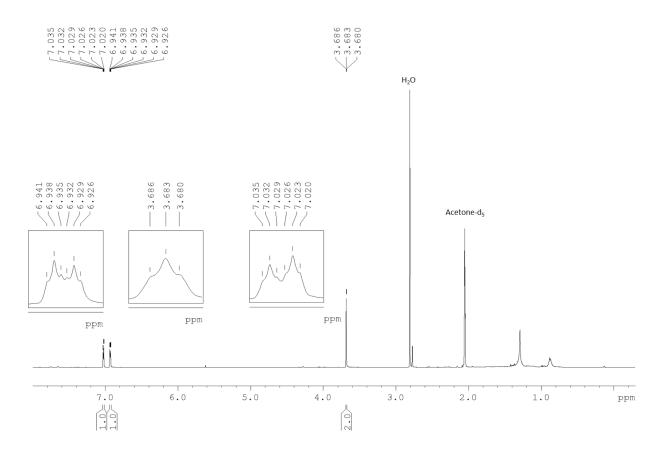


Figure S36. ¹H-NMR spectrum of B2 in (CD₃)₂CO (600 MHz)

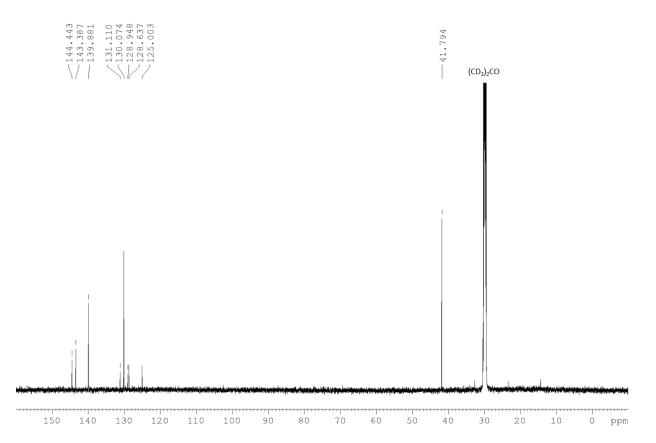


Figure S37. ¹³C-NMR spectrum of B2 in (CD₃)₂CO (600 MHz)

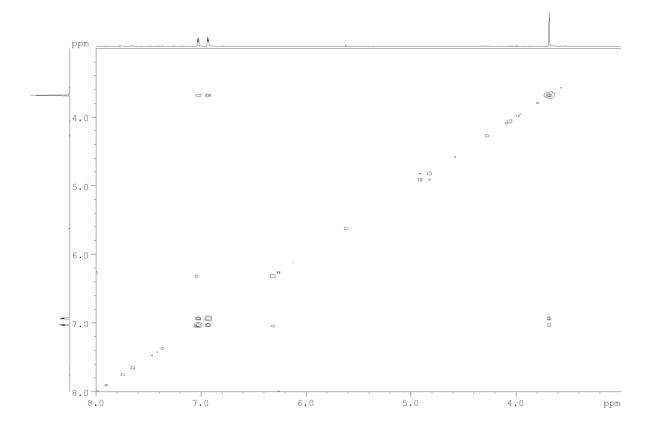


Figure S38. COSY spectrum of B2 in (CD₃)₂CO (600 MHz)

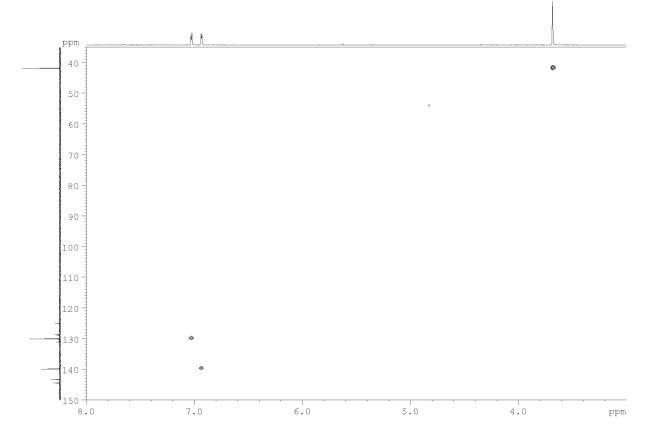


Figure S39. HSQC spectrum of B2 in (CD₃)₂CO (600 MHz)

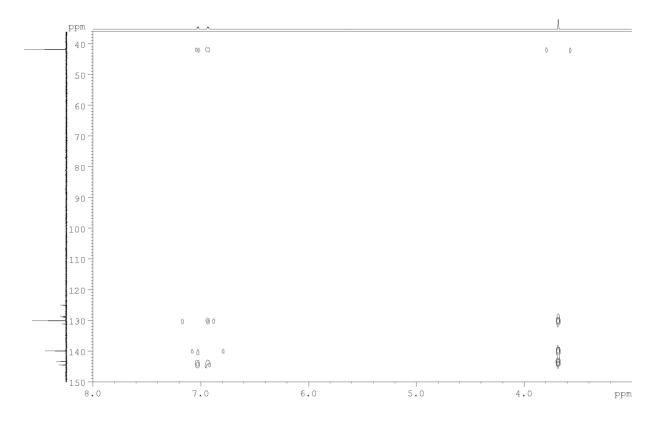


Figure S40. HMBC spectrum of B2 in (CD₃)₂CO (600 MHz)

CI 6 7 8 CI 5 4 9 CI 5 4 9 CI			COSY				
#	¹³ C(∂)	HSQC	¹H(<i>δ</i>)	J Coupling (Hz)	Protons	НМВС	COSY
1 2	41.8	CH ₂	3.68	t. 1.9	2H	C2, C3, C8	H-2, H-3
	139.9	СН	6.93	dt, 5.5, 1.9	1H	C1, C3, C9	H-1
3	130.1	CH	7.03	dt, 5.5, 1.7	1H	C1, C2, C9	H-1
4	125.0						
5	131.1						
6	128.9						
7	128.6						
8	143.4						
9	144.4						

Table S9. NMR data assignment of B2 in $(CD_3)_2CO$. HRMS (m/z): $[M-H]^-$ calcd. for $C_9Cl_4H_3$, 250.8983; found, 250.8993. TLC: Rf (cyclohexane : ethyl acetate 9/1) = 0.68

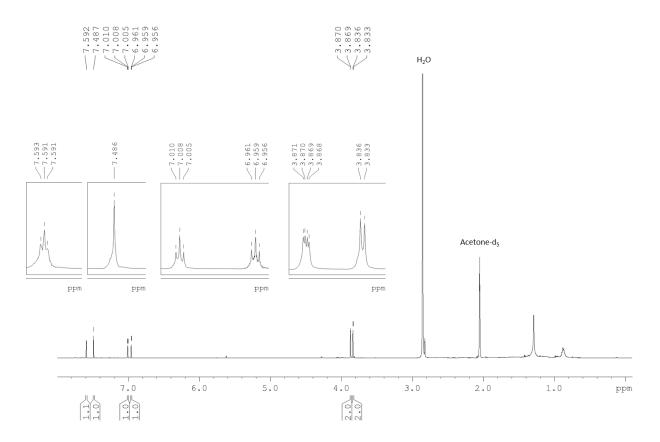


Figure S41. ¹H-NMR spectrum of B3-B4 in (CD₃)₂CO (600 MHz)

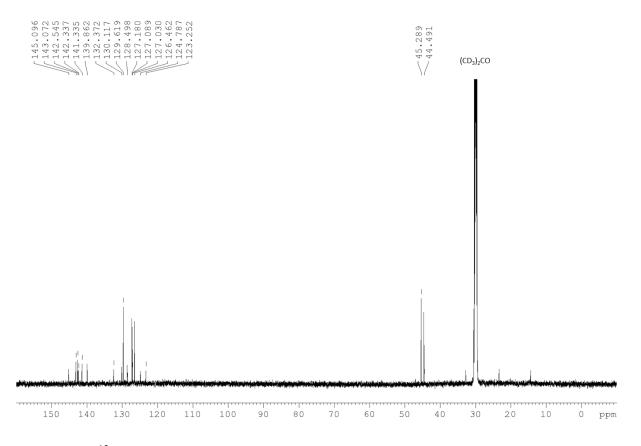


Figure S42. ¹³C-NMR spectrum of B3-B4 in (CD₃)₂CO (600 MHz)

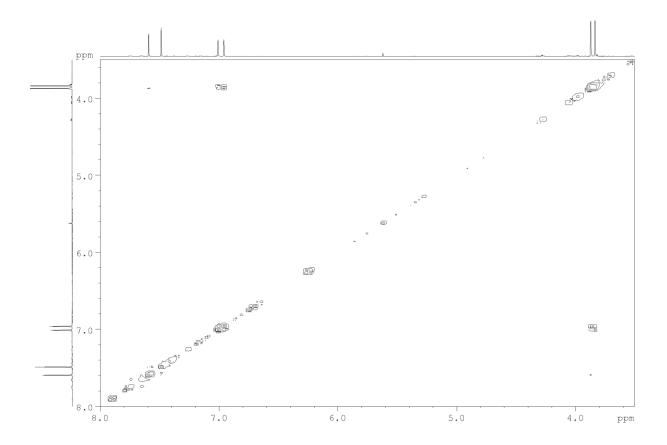


Figure S43. COSY spectrum of B3-B4 in (CD₃)₂CO (600 MHz)

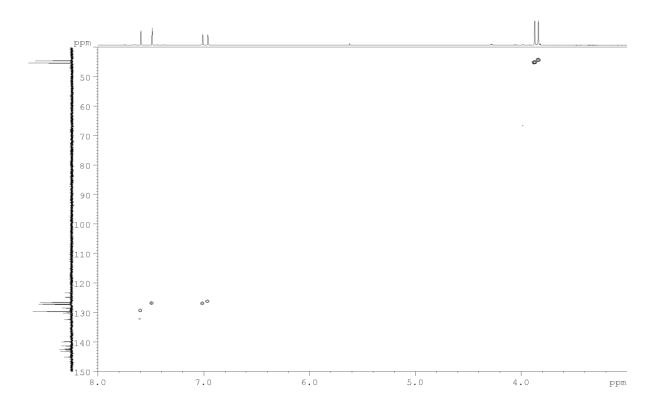


Figure S44. HSQC spectrum of B3-B4 in (CD₃)₂CO (600 MHz)

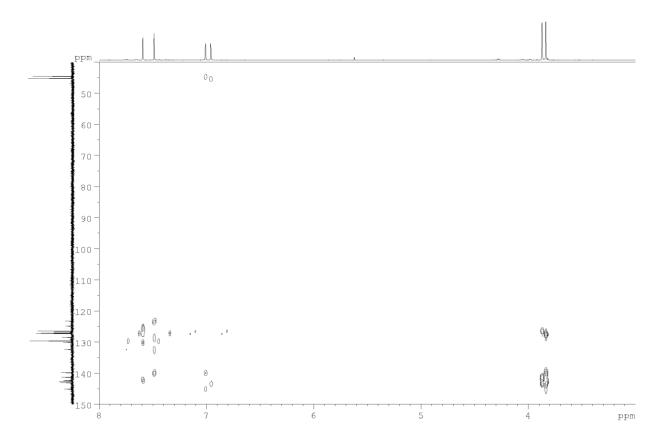
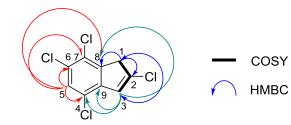


Figure S45. HMBC spectrum of B3-B4 in (CD₃)₂CO (600 MHz)

$ \begin{array}{c} & & \\ & & $										
#	¹³ C(∂)	HSQC	¹H(<i>δ</i>)	J Coupling (Hz)	Protons	НМВС	COSY			
1 2 3	45.3	CH2	3.87	dd, 1.3, 0.6		C2, C3, C8	H-3, H-6			
2	143.1									
	126.5	СН	6.96	t, 1.4		C1, C4, C8, C9	H-1			
4	124.8									
5*	130.1									
6*	129.6	СН	7.59	t <i>,</i> 0.6		C5, C9	H-1			
7*	127.0									
8	141.3									
9	142.3									

Table S10. NMR data assignment of B3 in $(CD_3)_2CO$. HRMS (m/z): $[M-H]^-$ calcd. for $C_9Cl_4H_3$, 250.8983; found, 250.8993. TLC: Rf (cyclohexane : ethyl acetate 9/1) = 0.88 * means that carbon position is assumed but not confirmed



#	¹³ C(∂)	HSQC	¹H(<i>δ</i>)	J Coupling (Hz)	Protons	НМВС	COSY
1	44.5	CH2	3.83	d. 1.4	2H	C2. C3. C8	H-3. H-5
2	142.5						
3	127.2	CH	7.01	t, 1.5	1H	C1, C4, C8, C9	H-1
4	123.3						
5*	127.1	СН	7.49	S	1H	C4, C6, C7, C8	
6*	132.4						
7*	128.5						
8	139.9						
9	145.1						

Table S11. NMR data assignment of B4 in $(CD_3)_2CO$. HRMS (m/z): $[M-H]^-$ calcd. for $C_9Cl_4H_3$, 250.8983; found, 250.8993. TLC: Rf (cyclohexane : ethyl acetate 9/1) = 0.88 * means that carbon position is assumed but not confirmed

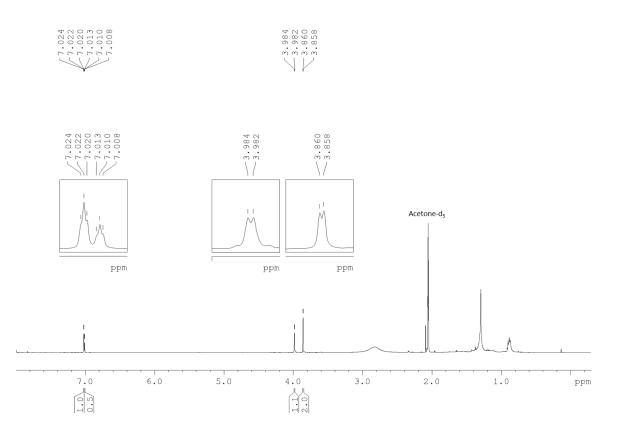


Figure S46. ¹H-NMR spectrum of C1-C2 in $(CD_3)_2CO$ (600 MHz)

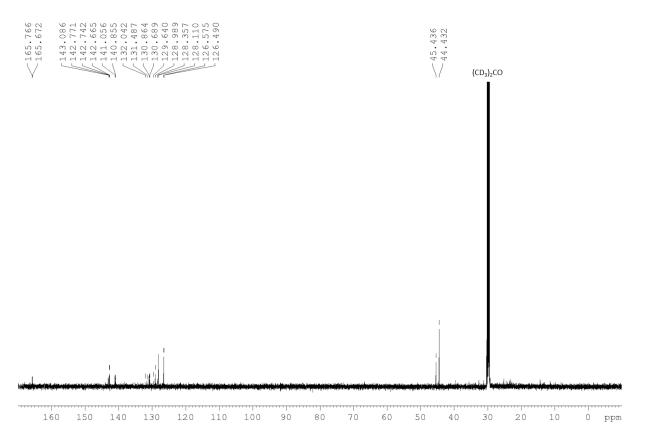


Figure S47. ¹³C-NMR spectrum of C1-C2 in (CD₃)₂CO (600 MHz)

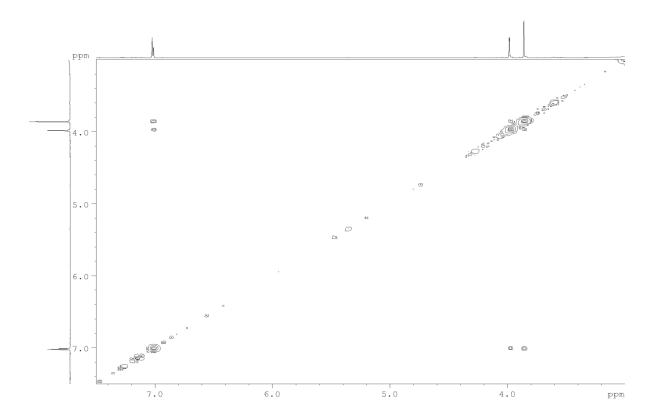


Figure S48. COSY spectrum of C1-C2 in (CD₃)₂CO (600 MHz)

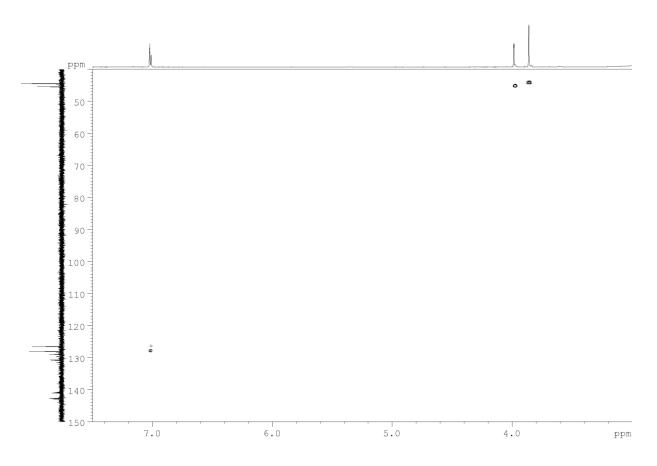


Figure S49. HSQC spectrum of C1-C2 in (CD₃)₂CO (600 MHz)

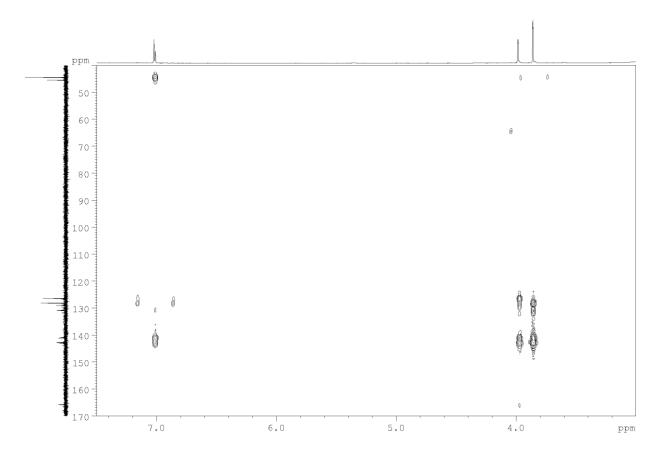


Figure S50. HMBC spectrum of C1-C2 in (CD₃)₂CO (600 MHz)

CI CI CI 6 CI 5 4 HOOC		\cap	COSY HMBC				
#	¹³ C(∂)	HSQC	¹H(<i>ð</i>)	J Coupling (Hz)	Protons	HMBC	COSY
1 2	44.4		3.86	d. 1.4	2H	C2. C3. C7. C8	H-3
2	142.7		7.02	± 4 4	411	C1 C1 C0	11.4
3	128.4		7.02	t, 1.4	1H	C1, C4, C9	H-1
4	130.7						
5*	132.0						
6*	131.5						
7*	130.9						
8	140.9						
9	142.7						
10	165.7						

Table S12. NMR data assignment of C1 (major isomer) in $(CD_3)_2CO$. HRMS (m/z): [M-H]⁻ calcd. for $C_{10}Cl_4O_2H_3$, 294.8882; found, 294.8888. TLC: Rf (methanol : dichloromethane : formic acid 1/9/0.05) = 0.5

			COSY HMBC				
#	¹³ C(∂)	HSQC	¹H(<i>ð</i>)	J Coupling (Hz)	Protons	HMBC	COSY
1 2	45.4		3.98	d. 1.5	2H	C2. C3. C7. C8	H-3
2 3	142.8 126.5		7.01	+ 1 5	1H	C1 C4 C0	Ц 1
5 4	126.5		7.01	t, 1.5	ΤΠ	C1, C4, C9	H-1
5*	128.1						
6*	129.0						
7*	129.6						
8	141.1						
9	143.2						
10	165.8						

Table S13. NMR data assignment of C2 (minor isomer) in $(CD_3)_2CO$. HRMS (m/z): [M-H]⁻ calcd. for $C_{10}Cl_4O_2H_3$, 294.8882; found, 294.8888. TLC: Rf (methanol : dichloromethane : formic acid 1/9/0.05) = 0.5

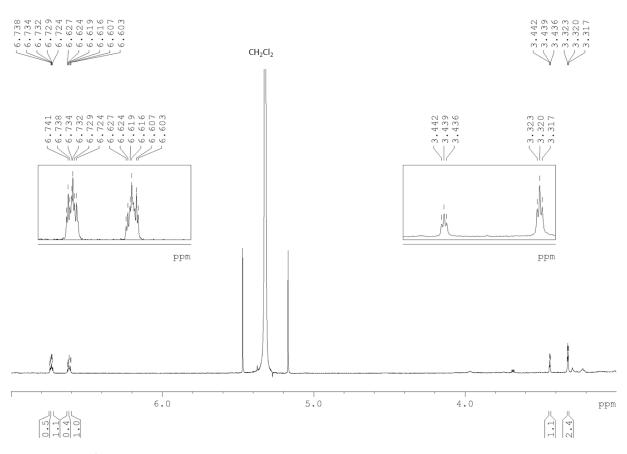


Figure S51. ¹H-NMR spectrum of C5-C6 in CD₂Cl₂ (600 MHz)

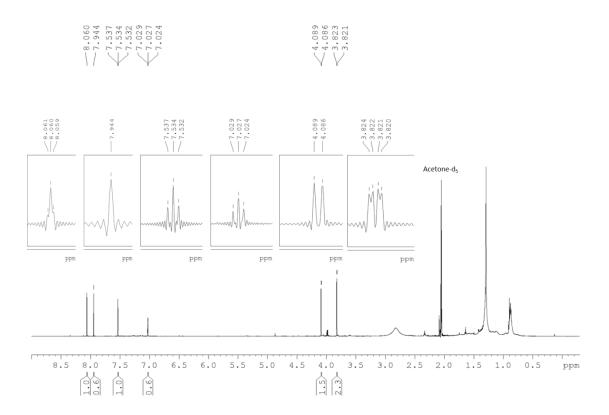


Figure S52. full ¹H-NMR spectrum of C3-C4 in $(CD_3)_2CO$ (600 MHz, exponential apodization parameter: Hz) including enlargement windows (exponential apodization parameter: -0.3 Hz).

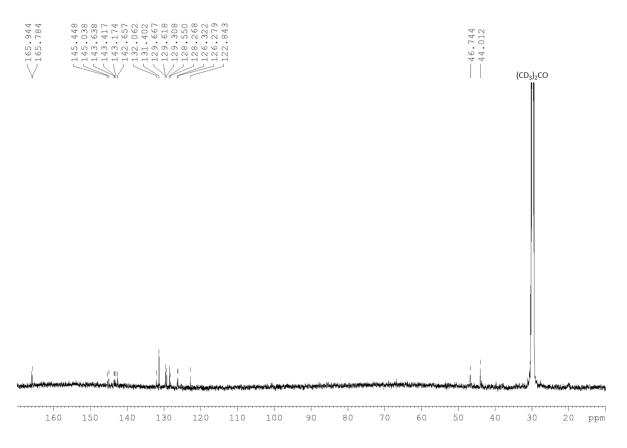


Figure S53. ¹³C-NMR spectrum of C3-C4 in (CD₃)₂CO (600 MHz)

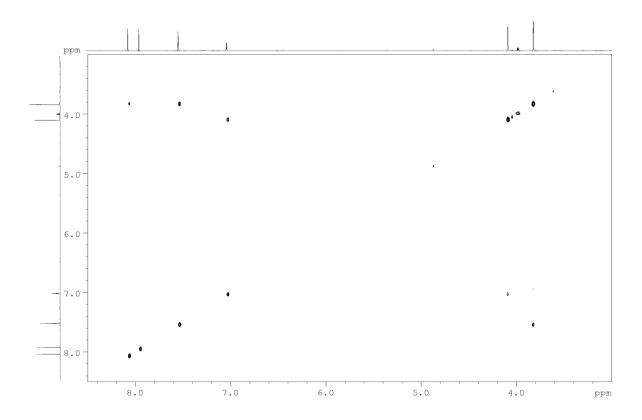


Figure S54. COSY spectrum of C3-C4 in (CD₃)₂CO (600 MHz)

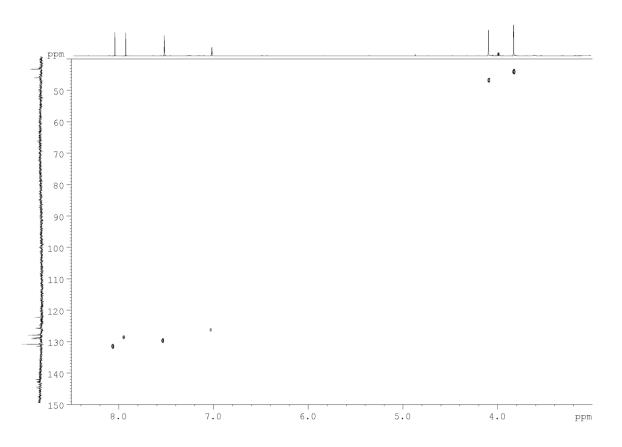


Figure S55. HSQC spectrum of C3-C4 in (CD₃)₂CO (600 MHz)

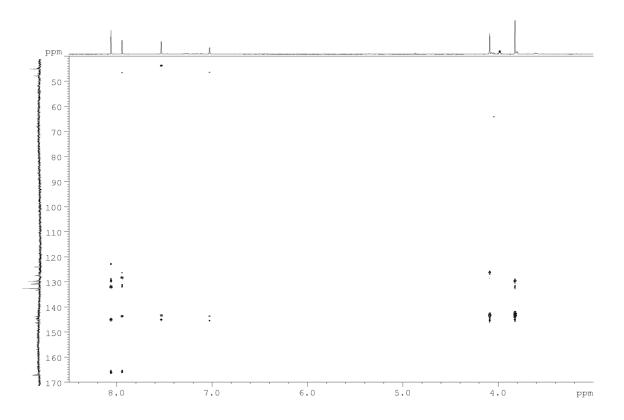
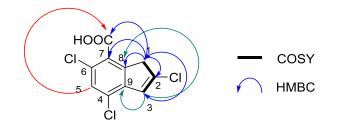


Figure S56. HMBC spectrum of C3-C4 in (CD₃)₂CO (600 MHz)



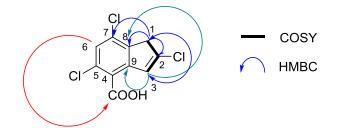
#	¹³ C(∂)	HSQC	¹H(<i>δ</i>)	J Coupling (Hz)	Protons	НМВС	COSY
1	46.7		4.09	d. 1.5	2H	C2. C3. C7. C8. C10	H-3
2	145.4						
3	126.3		7,03	t, 1.5	1H	C1, C8, C9	H-1
4*	126.3						
5*	128.6		7,94	S	1H	C10 ⁱ	
6*	129.7						
7*	128.3						
8	143.2						
9	143.6						
10	165.8						

Table S14. NMR data assignment of C3 in $(CD_3)_2CO$. HRMS (m/z): $[M-H]^-$ calcd. for $C_{10}Cl_3O_2H_4$ 260.9271; found, 260.9278. TLC: Rf (methanol : dichloromethane : formic acid

1/9/0.05) = 0.69

* means that carbon position is assumed but not confirmed

ⁱ means that others HMBC couplings were observed but were not attributed due to the incertitude on carbon position



#	¹³ C(∂)	HSQC	¹H(<i>δ</i>)	J Coupling (Hz)	Protons	НМВС	COSY
1 2	44.0		3.82	dd. 1.6. 0.6	2H	C2. C3. C7. C8	H-3. H6
	143.4						
3	129.3		7.53	t, 1.5	1H	C1, C8, C9	H-1
4*	132.1						
5*	131.4						
6*	122.8		8.06	t <i>,</i> 0.6	1H	C10 ⁱ	H-1
7*	129.6						
8	142.7						
9	145.0						
10	165.9						

Table S15. NMR data assignment of C4 in $(CD_3)_2CO$. HRMS (m/z): $[M-H]^-$ calcd. for $C_{10}Cl_3O_2H_4$ 260.9271; found, 260.9278. TLC: Rf (methanol : dichloromethane : formic acid

1/9/0.05) = 0.69

* means that carbon position is assumed but not affirmed

ⁱ means that others HMBC couplings were observed but were not attributed due to the incertitude on carbon position

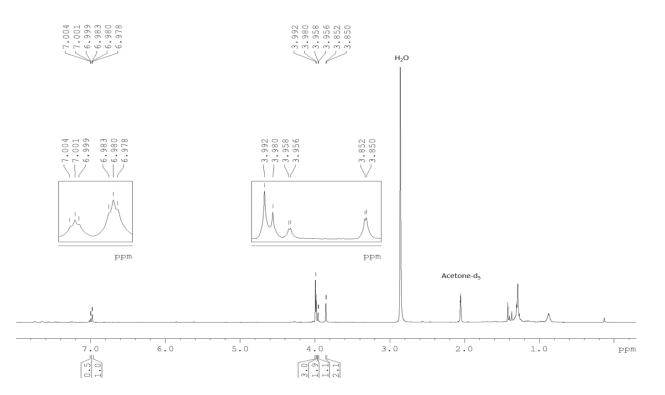


Figure S57. ¹H-NMR spectrum of D1-D2 in (CD₃)₂CO (600 MHz)

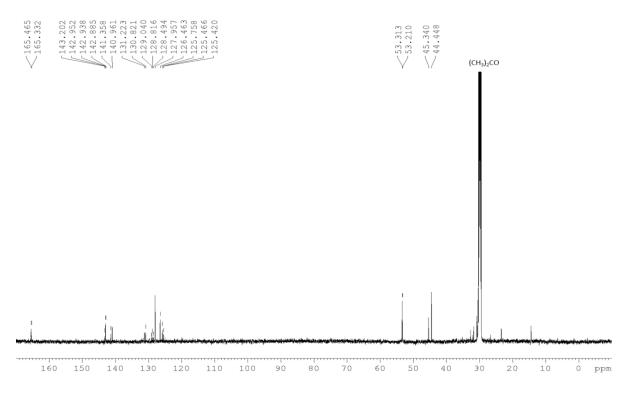


Figure S58. ¹³C-NMR spectrum of D1-D2 in (CD₃)₂CO (600 MHz)

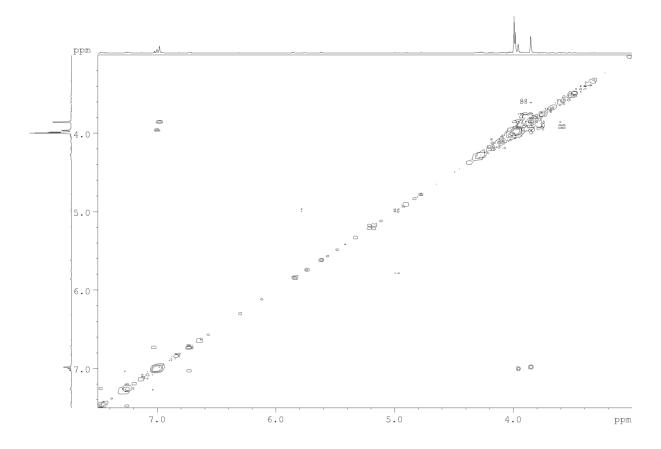


Figure S59. COSY spectrum of D1-D2 in (CD₃)₂CO (600 MHz)

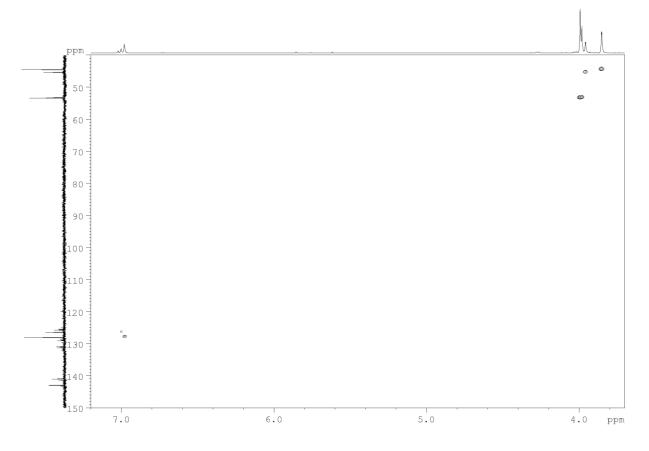


Figure S60. HSQC spectrum of D1-D2 in (CD₃)₂CO (600 MHz)

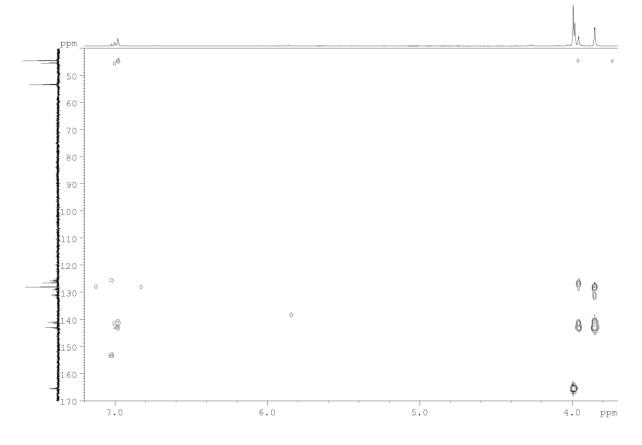
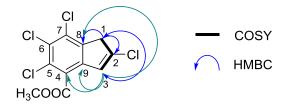


Figure S61. HMBC spectrum of D1-D2 in (CD₃)₂CO (600 MHz)

H_3COOC CI CI CI 5 4 CI CI 5 4 CI CI 5 4 CI		$\widehat{}$	COSY HMBC				
#	¹³ C(∂)	HSQC	¹H(<i>δ</i>)	J Coupling (Hz)	Protons	НМВС	COSY
1 2	45.3		3.95	d. 1.5	2H	C2. C3. C8	H-3
	143.2						
3	126.5		7.00	t, 1.5	1H	C1, C9	H-1
4*	125.5						
5*	129.0						
6*	128.8						
7*	128.5						
8	141.4						
9	143.2142.9						
10	165.3						
11	53.2		3.98	S	3H		

Table S16. NMR data assignment of D1 in $(CD_3)_2CO$. HRMS (m/z): $[M]^-$ calcd. for $C_{11}Cl_4O_2H_5$, 308.9038; found, 308.9046.



#	¹³ C(∂)	HSQC	¹H(<i>δ</i>)	J Coupling (Hz)	Protons	HMBC	COSY
1	44.4		3.85	d. 1.4	2H	C2. C3. C7. C8	H-3
2	142.9						
3	128.0		6.98	t, 1.4	1H	C1, C9	H-1
4*	130.7						
5*	125.8						
6*	125.4						
7*	131.2						
8	141.1						
9	142.9						
10	165.5						
11	53.3		3.99	S	3H		

Table S17. NMR data assignment of D2 in $(CD_3)_2CO$. HRMS (m/z): $[M]^-$ calcd. for $C_{11}Cl_4O_2H_5$, 308.9038; found, 308.9046.

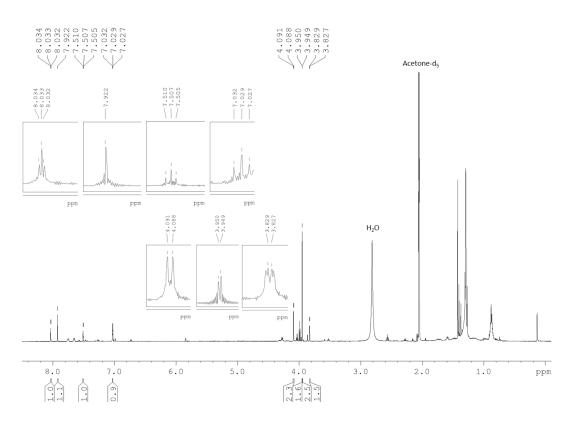


Figure S62. ¹H-NMR spectrum of D3-D4 in $(CD_3)_2CO$ (600 MHz, exponential apodization parameter: Hz) including enlargement windows (exponential apodization parameter: -0.3 Hz).

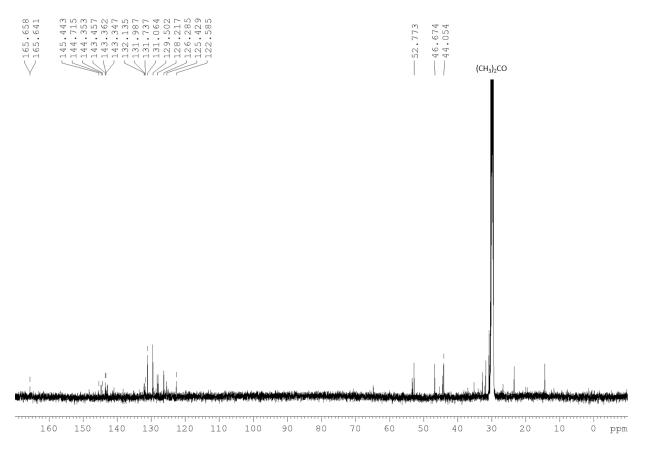


Figure S63. ¹³C-NMR spectrum of D3-D4 in (CD₃)₂CO (600 MHz)

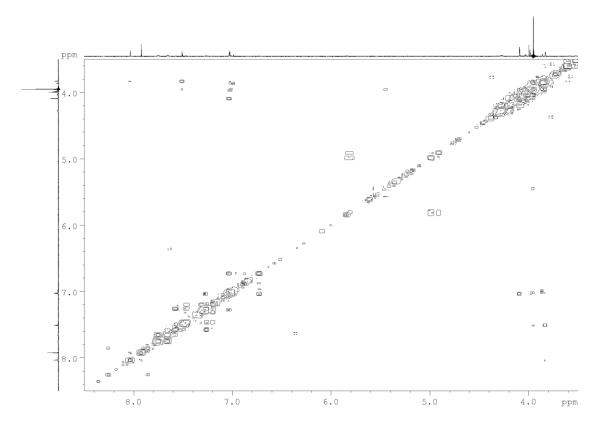


Figure S64. COSY spectrum of D3-D4 in (CD₃)₂CO (600 MHz)

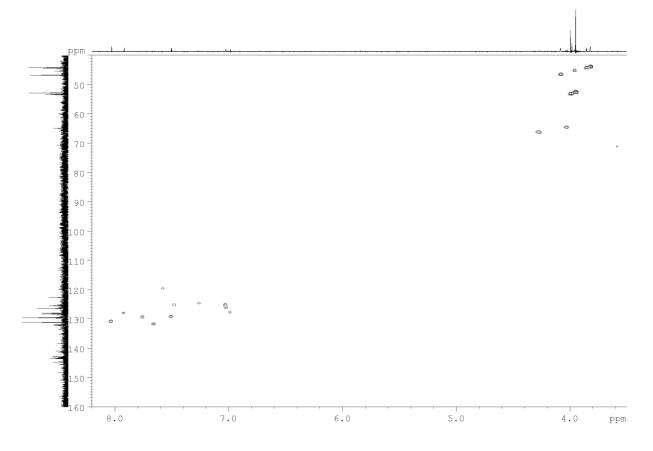


Figure S65. HSQC spectrum of D3-D4 in (CD₃)₂CO (600 MHz)

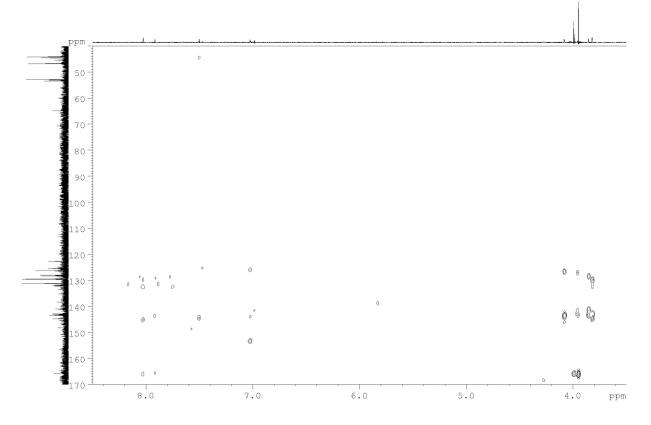
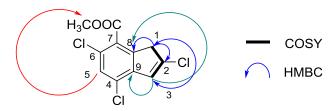


Figure S66. HMBC spectrum of D3-D4 in (CD₃)₂CO (600 MHz)



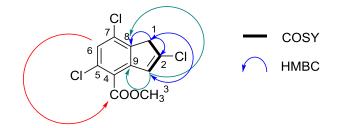
#	¹³ C(∂)	HSQC	¹H(<i>δ</i>)	J Coupling (Hz)	Protons	НМВС	COSY
1 2	46.7 143.3		4.08	d. 1.6	2H	C2. C3. C8	H-3
2 3	145.5 125.4		7.02	t, 1.7	1H	C1, C8, C9	H-1
4*	128.2						
5*	128.3		7.92	S	1H	C10 ⁱ	
6*	131.1						
7*	131.7						
8	143.5						
9	145.4						
10	165.6						
11	52.8		3.95	S	3H		

Table S18. NMR data assignment of D3 in $(CD_3)_2CO$. HRMS (m/z): [M]⁻ calcd. for $C_{11}Cl_3O_2H_6$ 274.9428; found, 274.9437.

* means that carbon position is assumed but not confirmed

ⁱ means that others HMBC couplings were observed but were not attributed due to the incertitude

on carbon position



#	¹³ C(∂)	HSQC	¹H(<i>ð</i>)	J Coupling (Hz)	Protons	НМВС	COSY
1 2	44.1		3.82	dd. 1.5 : 0.6	2H	C2. C3. C8	H-3. H6
2 3	143.4 129.5		7.5	+ 1 6	1⊔	C1, C8, C9	H-1
5 4*	129.5		7.5	t, 1.6	1H	CI, Co, C9	H-T
- 5*	132.1						
6*	132.0		8.03	t, 0.6	1H	C10 ⁱ	H-1
7*	129.7						
8	144.4						
9	144.7						
10	165.7						
11	52.8		3.95	S	3H		

Table S19. NMR data assignment of D4 in (CD₃)₂CO. HRMS (m/z): [M]⁻ calcd. for

 $C_{11}Cl_3O_2H_6$ 274.9428; found, 274.9437.

* means that carbon position is assumed but not confirmed

ⁱ means that others HMBC couplings were observed but were not attributed due to the incertitude

on carbon position

S6 Supporting References

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