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6 **REVISION 1**

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8 **Abiotic and biotic processes that drive carboxylation and decarboxylation reactions**

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42 **Abstract**

43 Carboxylation and decarboxylation are two fundamental classes of reactions that impact
44 the cycling of carbon in and on Earth’s crust. These reactions play important roles in both long-

45 term (primarily abiotic) and short-term (primarily biotic) carbon cycling. Long-term cycling is
46 important in the subsurface and at subduction zones where organic carbon is decomposed and
47 outgassed or recycled back to the mantle. Short-term reactions are driven by biology and have the
48 ability to rapidly convert CO₂ to biomass and vice versa. For instance, carboxylation is a critical
49 reaction in primary production and metabolic pathways like photosynthesis, in which sunlight
50 provides energy to drive carbon fixation, whereas decarboxylation is a critical reaction in
51 metabolic pathways like respiration and the tricarboxylic acid cycle. Early life and prebiotic
52 chemistry on Earth likely relied heavily upon the abiotic synthesis of carboxylic acids. Over time,
53 life has diversified (de)carboxylation reactions and incorporated them into many the facets of
54 cellular metabolism. Here we present a broad overview of the importance of carboxylation and
55 decarboxylation reactions from both abiotic and biotic perspectives to highlight the importance of
56 these reactions and compounds to planetary evolution.

57 **Keywords:** Decarboxylation, carboxylation, subduction recycling, biological carbon cycling,
58 early Earth

59 **Introduction**

60 Earth's carbon cycle consists of a complex interplay of biological and abiological processes
61 that together contribute to making Earth unique among the planets in our Solar System. For life
62 living on and in Earth's crust, carbon is a primary element and represents approximately 48% of
63 the dry weight biomass of almost all organisms. Life's tight coupling to the carbon cycle is an
64 important feature of our planet, and organisms have evolved many complex mechanisms for
65 scavenging and transforming carbon in the environment. The contemporary carbon cycle stands in
66 stark contrast to how the carbon cycle likely operated over much of Earth's history. Prior to the
67 emergence of microbial life sometime before 3.5 Ga, the abiological carbon cycle consisted of

68 simple organic molecules, including amino acids (Miller 1953). However, as life on Earth began
69 to diversify and the redox states of Earth's oceans and atmospheres began to change, the carbon
70 cycle diversified in tandem.

71 **Sources and transformation of carbon on Earth**

72 As Earth's carbon cycle evolved, several core biological and abiological reactions were
73 key, and these were heavily affected by the prevailing oxidation state of carbon. Due to carbon's
74 intrinsic properties (*e.g.* bond lengths and strengths in combination with other elements, large
75 number of oxidation states, tetravalency, etc.), there are estimated to be as many as 10^{80} possible
76 constitutional isomers of C under 500 atomic mass units (amu) (Lipinski and Hopkins 2004). Few
77 of these are likely to be actualized. Remarkably, terrestrial life relies on a relatively small
78 proportion of these, 10-100s of primary metabolites (Dobson 2004; Smith and Morowitz 2004)
79 and few hundred thousand secondary metabolites (Ji et al. 2009). Carbon compounds occupy
80 various metastable oxidation states, ranging from +4 to -4, for CO₂ and CH₄ respectively.

81 At the heart of the modern carbon cycle is the reduction of CO₂ (inorganic) to form reduced
82 organic compounds and the stepwise oxidation of organic C back to CO₂ (Figure 1). Many of these
83 reactions can be generalized as *carboxylations*, *i.e.*, the addition of CO₂ to an existing organic
84 molecule, yielding carboxylic acids, and *decarboxylations*, the loss of carboxylic acid groups and
85 their release as CO₂. Carboxylations and decarboxylations can occur either biotically through
86 enzymatic mechanisms or abiotically for example via the influence of UV light (Getoff 1965; Chiu
87 et al. 2017), thermal chemistry (see Papineau et al. 2017, and below), or the influence of mineral
88 interactions (Weber 2004).

89 Carboxylation reactions can be generalized into long-term (or slow; >100,000 yrs)
90 processes and short-term (or fast; decades or less) C cycle processes (Berner 2003). With the

91 advent and diversification of life, the rate of short-term, biological C cycling has increased relative
92 to the rate of long-term, primarily abiotic processes. Short-term biological cycling operates at rates
93 many orders of magnitude faster than long-term cycling. Throughout Earth's history, the balance
94 between these carboxylation and decarboxylation reactions has influenced Earth's climate (Archer
95 2008; Lacy et al. 2010).

96 **Long-term carbon cycle.**

97 The long-term cycle is primarily *abiotic* and likely has occurred over most of Earth's
98 history (Rothman et al. 2003), and is dependent upon plate tectonics, as the subducting plates bring
99 both organic and inorganic carbon fixed at the surface into the solid Earth. The subducted carbon
100 has at least two major fates, it can be emitted as volatiles through volcanism, or it can be returned
101 to the mantle where it can be converted to various forms. The phases of carbon in the mantle (fluid,
102 solid, or dissolved) depend on the pressure, intensity and rate of heating, as well as the ambient
103 redox conditions, but graphite, carbon dioxide, and methane are likely abundant species (Figure
104 2). Ultimately, the quantity and ratio of reduced to oxidized carbon and its proxies in the
105 subducting material are influenced by the redox state of both the deep Earth and atmosphere
106 (Hayes and Waldbauer 2006).

107 **Short-term carbon cycle.**

108 The short-term cycle is essentially *biotic* and therefore unique to Earth, at least as far as
109 current knowledge extends. Biological C cycling is a balance between 1) autotrophic CO₂ fixation
110 (carboxylation), which occurs primarily through the Calvin-Benson-Bassham Cycle (CBB) used
111 by diverse groups of bacteria, archaea, and photosynthetic eukaryotes (Erb and Zarzycki 2018),
112 and 2) the biological degradation of organic matter to CO₂ through fermentation and respiration
113 (decarboxylation). Biogenic and thermogenic methane serve as additional important shunts

114 between biologically derived and non-biologically derived carbon pools, as methane may be
115 converted first to CO₂ then incorporated to biomass via the Calvin-Benson-Bassham cycle
116 (Murrell and Jetten 2009; Chistoserdova 2011). Finally, fixed biological C is also oxidized via
117 anthropogenic combustion of biomass and fossil fuels.

118 Microorganisms were the major life forms on Earth in terms of biomass until relatively
119 recently, having been dethroned by terrestrial plants in the last few hundred million years (Bar-On
120 et al. 2018; McMahon and Parnell 2018). Interestingly, there are few primary biological C fixation
121 pathways (Fuchs 2011) relative to the diversity of microbial life (Hug et al. 2016). Nonetheless,
122 as life diversified and evolved, so too did cellular biochemistry resulting in the evolution of wholly
123 new classes of organic compounds, such as lignin produced by vascular plants and some algae
124 (Martone et al. 2009).

125 **Abiotic processes**

126 **Abiotic carboxylation reactions**

127 Fischer-Tropsch Type (FTT) synthesis is a primary pathway for abiotic carboxylic acid
128 formation. This class of reactions can plausibly occur under hydrothermal conditions via the
129 reduction of CO₂, CO, or HCO₃⁻ with H₂ in the presence of transition metal catalysts (*i.e.* Ni, Cr,
130 and Fe), which produces short-chained alkanes and simple carboxylic acids, like formate, oxalate
131 and short fatty acids (Berndt et al. 1996; Foustoukos and Seyfried 2004). A large range of
132 carboxylic acids have been experimentally synthesized in FTT reactions using formate as a
133 reactant, which also easily forms under these hydrothermal conditions (McCollom and Simoneit
134 1999), though it should be pointed out the widespread occurrence of such abiotic synthesis on
135 Earth has been recently called into question (Fiebig et al. 2019). Furthermore, phosphatized fatty
136 acids (also called phospholipids) can self-assemble into micelles in aqueous solutions (Chen and

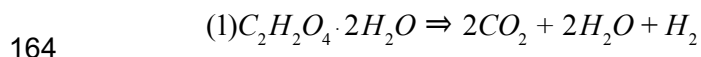
137 Walde 2010). As the complexity of cells evolved, fatty acids would evolve specific roles in cellular
138 membranes, giving them unique properties, such as self-assembly into micelles (Segré et al. 2001).
139 FTT synthesis has been used to explain the origin of carboxylic acids and various other
140 hydrocarbons in hydrothermal vents (Rushdi and Simoneit 2006; McCollom and Seewald 2007),
141 deep crustal settings (Sherwood Lollar et al. 2002), and meteorites (Sephton 2002; Steele et al.
142 2016).

143 **Decarboxylation reactions at high-pressures and temperatures**

144 As part of the long-term cycling of carbon, subduction zones play a direct role in returning
145 surface carbon to the mantle (Kelemen and Manning 2015). Near-continent sediments can be very
146 rich in organic matter (up to 60 wt%, Mayer 1994), for instance sediments formed during anoxic
147 events or located in near-shore environments with highly productive overlying waters (Hartnett et
148 al. 1998). Once these sediments are subducted, their organic carbon content can be transported
149 down to the upper mantle, as suggested by the ^{12}C -enriched isotopic signature of diamonds derived
150 from the upper mantle, the transition zone and lower mantle (Stachel et al. 2002; Cartigny et al.
151 2004; Palot et al. 2016). Kelemen and Manning (2015) revised upward the maximum flux of
152 carbon subducted in sediments from 17 Mt C/y to 23 Mt C/y, adding the previously neglected
153 contribution of organic carbon averaged to around 1 wt%. However, the fate of organic matter in
154 subduction zones is still debated, and present knowledge relies mostly on experimental studies
155 focused on the interaction between crystalline graphite and aqueous fluids (Poli et al. 2009;
156 Tumiati et al. 2012, 2017) or melts (Duncan and Dasgupta 2017).

157 Organic matter can have an impact on CO_2 degassing in subduction zones, thus
158 understanding pre-graphite states of organic matter may provide valuable insight subduction zone
159 carbon decomposition. Organic acids have been used since the 1960's as sources of C-O-H fluids

160 in high-pressure (HP)/high-temperature (HT) experiments (SI Table 1) because they decompose
161 at relatively low temperatures generating complex carbon-bearing fluid mixtures depending on
162 their bulk composition (i.e., the C/O/H ratios). For instance, oxalic acid dihydrate decomposes to
163 CO₂ at 150–200°C (Pernert 1952; Morgan et al. 1992; Tiraboschi et al. 2016) through the reaction:

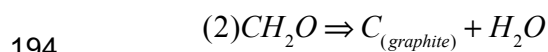


165 Since the pioneering work of Eugster and Skippen (1967), it is known that a range of C–O–H
166 fluids can be generated in high pressure (HP)/high temperature (HT) experiments, producing
167 aqueous fluids containing both CO₂ (i.e., oxidized C⁴⁺ carbon) and CH₄ (i.e., reduced C⁴⁻ carbon)
168 depending on the starting organic compound (Figure 3).

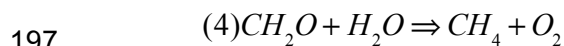
169 In principle, oxalic acid dihydrate (OAD) should decompose at HP/HT conditions to
170 produce graphite and a mixed H₂O–CO₂ fluid, while anthracene (ATR) should evolve to give
171 nearly pure methane, glucose and water (Figure 3). In fact, the disproportionation reaction of oxalic
172 acid and glucose to graphite + aqueous fluids bearing CH₄ and CO₂ has been observed
173 experimentally (Yamaoka et al. 2002; McCubbin et al. 2014). However, in real-world fluid–rock
174 systems, the redox state of the environment plays a major role (Tumiati and Malaspina 2019), such
175 that the oxidation state of carbon in fluids in equilibrium with graphite is dependent on the oxygen
176 fugacity conditions buffered by the rocks. Thus, C–O–H fluids will be CO₂-bearing under
177 oxidizing conditions and CH₄-bearing (C⁰ + C⁻⁴) under reducing conditions, and is independent of
178 the source organic compound's bulk composition.

179 Pressure and temperature are crucial to define whether a buffering mineral assemblage will
180 impose oxidizing or reducing conditions. For instance, at oxygen fugacity conditions constrained
181 by the equilibrium fayalite + O₂ = magnetite + quartz (FMQ), often taken as a reference redox
182 state for subducting slab–mantle interfaces (e.g., Tumiati and Malaspina 2019), C-O-H fluids in

183 equilibrium with graphite should change their composition from nearly pure water containing
184 traces of CH₄ and CO₂ at 1–2 GPa and 200–450°C, to mixed H₂O–CO₂ fluids at 3–4 GPa and 700–
185 800°C (Figure 3). As a consequence, while graphite should be somewhat refractory to dissolution
186 at forearc depths, it becomes readily soluble in aqueous fluids at subarc depths, which can dissolve
187 up to 80 mol% CO₂ in silicate-bearing systems (Tumiati et al. 2017). Moreover, although
188 experimental data are not available yet, thermodynamic calculations suggest the dissolution of
189 graphite in HP fluids may produce at certain pH–*f*O₂ conditions carboxylic acids such as acetate,
190 formate, propionate and/or bicarbonate and carbonate (Sverjensky et al. 2014; Pan and Galli 2016;
191 Tiraboschi et al. 2018). Nevertheless, as a general conclusion we can infer that the fate of organic
192 matter, represented below by CH₂O, is broadly governed in the deep Earth by the following
193 reactions:



195 which accounts for the graphitization of organic matter (Beysac and Rumble 2014) and



198 which accounts for the oxidation (combustion) (Equation 3) and for the reduction (abiotic
199 methanation) (Equation 4) of organic matter. At the high pressures and temperatures characterizing
200 subduction zones, organic matter is therefore expected to be converted into carbon polymorphs
201 (graphite, diamond) and aqueous fluids bearing carbon dissolved species that are constrained by
202 the redox state and pH of the local environment.

203 **Roles of abiotic decarboxylation in subsurface carbon stores**

204 Decarboxylation reactions play important roles in the maturation of buried organic material
205 on Earth. Carboxylic acid functional groups represent a significant portion of biological

206 macromolecules (*i.e.*, lipid membrane fatty acids, amino acids, and metabolic intermediates).
207 Carboxylic acids thus constitute one of the main carbon reservoirs in living organisms and are
208 heavily represented among the carbon species altered during diagenesis. In early stages of organic
209 matter decomposition cell lysis releases free carboxylic acid containing metabolites which are
210 readily taken up by other organisms. As decomposition progresses, membrane lipids undergo
211 saponification, which liberates fatty acids. These free often convert to anions or salts in
212 sedimentary environments. As diagenesis progresses, carboxylic acids decompose abiotically via
213 decarboxylation reactions to form alkanes and a mix of CO_2 , CO_3^{2-} and HCO_3^- depending on
214 factors such as pH and temperature. Decarboxylation reaction rates depend on pH,
215 temperature/depth, the concentration of reactants such as inorganic acids and oxidants, and the
216 presence of catalysts (Weber 2004). The rate of decarboxylation influences on the formation of
217 petroleum and its migration in the subsurface.

218 Thermocatalytic decarboxylation is a major abiotic pathway for the removal of carboxyl
219 groups from organic molecules. Simple one carbon carboxylic acids, such as formic acid,
220 decompose slowly in the absence of catalysts but rates can be accelerated at higher temperatures
221 ($>225^\circ\text{C}$) and controlled by equilibrium with CO_2 (McCollom and Seewald 2003). As an example,
222 acetic acid is thought to be formed in oil field brines, via decomposition of long-chain carboxylic
223 acids to form alkanes. Once the alkanes form, they are oxidized to alkenes, alcohols and aldehydes
224 to eventually form carboxylic acids. The resultant carboxylic acids can then decarboxylate to form
225 CO_2 and a n-1 alkane (SI Figure 1).

226 Likewise, the thermal decomposition of amino acids, which contain a carboxyl group as
227 well as an amine group, is influenced by mineral assemblages, oxidation state, and dissolved
228 species concentrations (Estrada et al. 2017). Some abiotic decarboxylation reactions may be

229 oscillatory, a laboratory example of such a reaction is the Belousov-Zhabotinsky reaction, which
230 produces CO₂ from carboxylic acids with strong halogen oxidants, such as iodate and bromate,
231 and strong acids like sulfuric and phosphonic acids (Zhabotinsky 1991; Papineau et al. 2017).
232 During catagenesis and metagenesis, abiotic decarboxylation typically occurs in sedimentary
233 layers below microbial habitability zones (e.g. of aerobic heterotrophy, sulfate reduction,
234 fermentation, and methanogenesis), which can contribute to the production of porewater carbonate
235 and the formation of concretionary structures (Irwin et al. 1977). Under higher temperature
236 regimes, thermal cracking of alkanolic acids produces n-alkanes, which is a dominant pathway for
237 the formation of petroleum (Kissin 1987). Production of heavy n-alkanes in kerogen occurs during
238 the decarboxylation of fatty acids and esters, and is slow under geological conditions (Kissin 1987;
239 Weber 2004). Kerogen composition is described by its O and H composition, with kerogen type
240 III having high O/C and low H/C atomic ratios, and kerogen type I having low O/C and high H/C
241 atomic ratios; kerogen type II is intermediate between types I and III (SI Figure 2).

242 Coal is thought to be formed similarly by thermal decomposition: peat and lignite initially
243 form during diagenesis and have the highest carboxylic acid content. Loss of CO₂ from these
244 source materials yields bituminous-grade coal, which contains progressively less volatiles
245 (hydrocarbon gases) and carboxylic acids as it undergoes catagenesis. Eventually nearly complete
246 decarboxylation leads to the formation of anthracite coal through metagenesis and metamorphism
247 (SI Figure 3).

248 **Biology's role in short-term carbon cycling**

249 Carboxylation and decarboxylation reactions are necessary for the building and
250 deconstruction of cellular metabolites. As mentioned above, within the cell, carboxylic acid groups
251 are prominent features of lipids, amino acids, and primary metabolites. Thus, carboxylic acids

252 serve many structural roles as building blocks for cells. Additionally, carboxyl groups help
253 maintain intracellular pH by acting as weak acids.

254 **Biotic Carboxylation reactions**

255 Biotic carboxylation reactions are crucial for photoautotrophic and chemoautotrophic life,
256 and thus are necessary for biogeochemical cycling of all elements. Without carbon entering the
257 system, cells would not be able to synthesize cellular material, including the enzymes that drive
258 elemental cycling. Thus, carboxylation reactions are important not only for the cell itself, as they
259 obviate the need for exogenous complex carbon compounds for growth, but they also likely drive
260 micro and macro community assembly and dynamics.

261 Ecologically, primary producers, *i.e.* organisms that fix CO₂, are keystone drivers of
262 ecosystems, enabling biomass transfer to higher trophic levels. Thus, the carboxylation reactions
263 they perform are essential for making carbon biologically active. Carbon fixation is an energy-
264 consuming process for cells and microbes have evolved several ways to fix carbon and offset the
265 metabolic costs (Fuchs 2011). For example, phototrophs, like cyanobacteria or chloroplasts in
266 plants, use the light energy to drive carbon fixation, while chemolithotrophs use energy stored in
267 reduced chemical substrates (*e.g.* H₂, H₂S, NH₃ or Fe²⁺). Here we will focus on a subset of these
268 reactions.

269 One of the most widespread modes of carboxylation is the Calvin-Benson-Bassham cycle,
270 using the enzyme ribulose-1,5-bisphosphate carboxylase/oxygenase (RuBisCO). For both aquatic
271 and terrestrial systems, this enzyme is commonly associated with photosynthesis as the RuBisCO
272 and the CBB pathway accounts for the fixation of over 99.5% of all biological carbon on Earth
273 (Raven JA 2009). In fact, RuBisCO is often considered to be the most abundant enzyme on Earth
274 (Ellis 1979; Raven 2013). At the heart of the reaction, RuBisCO adds CO₂ to a five-carbon sugar

275 (ribulose-1,5-bisphosphate), creating an unstable six-carbon molecule that decomposes to give two
276 3-phosphoglycerate molecules (Figure 4). Interestingly, despite being a leaky and energetically
277 costly pathway, the CBB cycle operates in many energy-limited environments, *e.g.* subsurface
278 environments (Lau et al. 2016), the deep ocean (Swan et al. 2011) and energy non-limited
279 hydrothermal vents (Anantharaman et al. 2013; Sheik et al. 2014). The prominence of this enzyme
280 and fixation cycle has altered the surface of Earth and its carbon isotopic signature over geological
281 time scales.

282 The reverse Tricarboxylic Acid Cycle (rTCA), also known as the Arnon-Buchanan cycle,
283 is considered the modern version of a proposed prebiotic autocatalytic cycle fueled by the
284 formation of the highly insoluble mineral pyrite in sulfur-rich hydrothermal environments
285 (Wächtershäuser 1988, 1990). Discovered in the anaerobic green phototrophic sulfur bacterium
286 *Chlorobium limicola*, the pathway is a reversal of the citric acid cycle (the Krebs or TCA cycle),
287 generating acetyl-CoA from two CO₂ molecules. Widespread among anaerobic and
288 microaerophilic bacteria, including all Aquificae, chemolithoautotrophic Epsilon proteobacteria,
289 Chlorobi, and Nitrospirae (Berg 2011; Fuchs 2011; Hügler and Sievert 2011) in addition to few
290 other bacterial strains, the rTCA cycle is energetically more favorable than the CBB cycle and
291 appears to be widespread in subsurface and extreme environments. The dependence of two out of
292 three carboxylation reactions on reduced ferredoxin has relegated this pathway to
293 anaerobic/microaerophilic conditions due to the difficulty of keeping ferredoxin reduced in the
294 presence of oxygen (Fuchs 2011). Two main variants of the cycle exist, symmetric and
295 asymmetric, although the evolutionary reasons for their differentiation are not well understood
296 (Aoshima et al. 2004; Braakman and Smith 2012; Giovannelli et al. 2017). In the asymmetric
297 version of this pathway isocitrate is formed directly via the NADPH-dependent carboxylation of

298 2-oxoglutarate. Interestingly, recent work has shown that several carboxyl intermediates of the
299 rTCA cycle can be mediated by simple metal ions (Muchowska et al., 2017), suggesting a possible
300 very early evolution of carbon fixation metabolism.

301 The Wood-Ljungdahl (WL) pathway is thought to be one of the most ancient microbial
302 metabolisms for carbon fixation (Weiss et al. 2016). At its core are two carbon reduction reactions
303 that form one carbon carboxyl intermediates that are then converted to the acetate (Figure 4), hence
304 this pathway is commonly referred to as acetogenesis. Some extant microorganisms, like sulfate-
305 reducing Delta proteobacteria, are able to use the WL pathway for carbon fixation or for carbon
306 mineralization by reversing the pathway (Ragsdale 2008). The WL pathway only operates in strict
307 anaerobes, *i.e.* microbes that grow only under reducing conditions. Since anaerobic conditions
308 were almost certainly dominant throughout Earth's history and most subsurface environments are
309 reducing, the WL pathway is generally considered more ancient than the CBB pathway. This
310 pathway may also be important in subsurface environments, due to its energetic efficiency (Cotton
311 et al. 2018). Additionally, the reversibility of the WL pathway facilitates carbon mineralization
312 making this an interesting, multifaceted pathway (Ragsdale 2008).

313 **Biotic decarboxylation reactions**

314 Biological decarboxylation is essential for the recycling of fixed carbon and for creating new
315 metabolic intermediates. Both autotrophic and heterotrophic organisms use decarboxylation
316 reactions to interconvert metabolites among biosynthetic pathways. For many heterotrophic
317 organisms, respiration fed by the Tricarboxylic Acid Cycle (TCA, also known as the Krebs cycle)
318 is a central metabolic pathway which distributes metabolites among various biosynthetic pathways
319 and enables the remineralization organic carbon (Figure 4). Carbon enters this pathway via acetyl-
320 CoA, is converted into various carboxylic acid intermediates and finally released as CO₂. The TCA

321 cycle is capable of complete carbon mineralization, ultimately converting organic carbon to CO₂,
322 thus this pathway also generates electrons that are directed to electron acceptors like, oxygen,
323 nitrate, iron, and sulfate or methane. Alternatively, decarboxylation can occur through various
324 fermentative pathways. Fermentation produces both CO₂ and a reduced carbon compound, some
325 of which are carboxylic acids, that can be further utilized by other organisms for growth. For
326 instance, the breakdown and fermentation of organic matter to short-chain carboxylic acids by
327 microorganisms in the gut is vital for human metabolism. At the ecosystem scale, the interplay
328 between heterotrophy and autotrophy shapes and drives the short-term carbon cycle.

329 **Earth's evolving biotic carbon cycle**

330 As biological metabolism diversified over geologic time, it is likely that carbon and
331 carboxyl compound diversity also increased. In extant organisms, carboxylation and
332 decarboxylation reactions are ubiquitous and key for producing many metabolic intermediates and
333 secondary metabolites. Several prominent, punctuated evolutionary developments have likely
334 helped spur carbon diversity by providing new habitats and niches for life to expand into, such as
335 meteorite impacts, plate tectonics, and the biologically driven oxygenation of the oceans and
336 atmosphere. It is likely short-term carbon cycling has changed considerably over geological time.

337 The proliferation of oxygenic photosynthetic microorganisms during the Palaeoproterozoic
338 Great Oxidation Event (GOE, between about 2.5 and 2.0 Ga) also coincided with widespread
339 fixation of CO₂ into biomass through CBB-mediated carboxylation. With copious amounts of O₂
340 in the environment, new forms of metabolism such as aerobic respiration and new forms of life,
341 including multicellular organisms, emerged (Han and Runnegar 1992; El Albani et al. 2019).
342 Higher plants have contributed significantly to organic C diversity through production of myriad
343 plant secondary compounds, *i.e.* lignin and phenolic acids (Renault et al. 2017). Concomitantly,

344 as life developed the means to make new carboxyl compounds emerged, new decarboxylation
345 pathways also evolved to degrade organic matter back to CO₂.

346 **Implications**

347 Carboxylation and decarboxylation are an important class of chemical reactions that have
348 helped shape life on the surface of Earth for billions of years. These reactions are driven through
349 biotic and abiotic processes at different locations in and on Earth's crust. Prebiotic carboxylic acid
350 synthesis may have been critical for the origin and development of life, which has been using this
351 class of reactions for energy generation and reproduction for billions of years. Where these
352 processes occurring in the environment (*i.e.* subduction zones, coal deposit, aerobic water column,
353 tropical forests) dictates the rates of reaction and the diversity of carboxylic acids produced.
354 Whether produced biotically or abiotically, these individual simple reactions are at the heart of the
355 complex and elegant pathways that cycle carbon on Earth and potentially on extraterrestrial
356 planets.

357 **Acknowledgments**

358 The authors would like to thank the participants of the Deep Carbon Observatory's Earth
359 in Five Reactions workshop that were originally part of the brainstorming session that served, in
360 part, to form the basis of this manuscript. Specifically, we would like to acknowledge Dan Colman,
361 Tony Hartshorn, Adrienne Hoarfrost, and Gunter Wegener for their early input. A special thanks
362 and to Josh Wood for help conceptualizing and illustrating figure two. We are also appreciative of
363 the Deep Carbon Observatory for providing support for the workshop.

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595

596 **Figure 1.** An example of an abiotic carboxylation (top) and a reversible biotic decarboxylation
597 (left to right) and carboxylation (right to left) reaction. Abbreviations: S-CoA=Co-enzyme A,
598 which is used to create thioesters of carboxylic acids, and NAD⁺/NADH₂=oxidized and reduced
599 forms of Nicotinamide Adenine Dinucleotide, a commonly used electron donor/acceptor in cells.
600

601 **Figure 2.** A simplified depiction of key carboxylation and decarboxylation reactions and their
602 association with short and long-term terrestrial carbon cycling. Long-term processes are depicted
603 by blue boxes and short-term processes in orange.

604

605

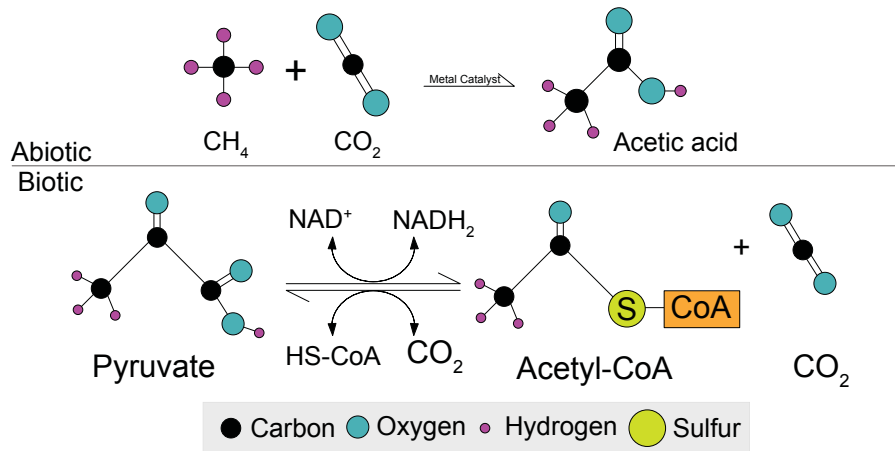
606 **Figure 3.** Ternary COH diagram showing a selection of organic compounds explored in high-
607 pressure/high-temperature experiments. Glucose composition is emphasized in yellow. Solid black
608 line: all possible compositions of COH fluids in equilibrium with pure crystalline graphite,
609 calculated by thermodynamic modelling at PT conditions characterizing the subduction slab
610 surface (Syracuse et al. 2010). Red and blue fields indicate respectively the presence of oxidized
611 (CO₂) and of reduced (CH₄) volatile carbon species in COH fluids. Dashed lines join graphite with
612 the expected composition of COH fluids in closed systems. White dots: calculated composition of
613 COH fluids at different P(GPa), T(°C) in systems open to oxygen, where its fugacity is buffered
614 externally at FMQ (fayalite/ferrosilite + magnetite + quartz/coesite) conditions. For every dot,
615 fO_2^{FMQ} was first calculated using the *Perple_X* package (cf. Connolly, 2005) and the
616 thermodynamic database (2002 update) of Holland and Powell (1998). Then, the composition of
617 the fluid has been calculated using the *Perple_X* routine "fluid" and the MRK equation of state of
618 graphite-buffered C-O-H-S fluids (eq. 7; Connolly and Cesare, 1993), and it has been proven to
619 be consistent with the COH fluid model of Zhang and Duan (2009). The dashed lines in Figure 3
620 join graphite (or diamond, depending on pressure–temperature conditions) with the predicted
621 composition of fluids in equilibrium with it (intercepts with the thick black line; (Connolly and
622 Cesare 1993; Zhang and Duan 2009), considering the decomposition of different organic
623 compounds.

624

625

626 **Figure 4.** Examples of central carbon metabolisms microorganisms use to fix CO₂ to biomass
627 (CBB= Calvin-Benson-Bassham, rTCA= reverse Tricarboxylic Acid, WL= Wood Ljungdahl).
628 Carbon mineralization to CO₂ commonly occurs through fermentations or through the
629 Tricarboxylic Acid (TCA) cycle. Green arrows represent carboxylation reactions, the red arrows
630 represent decarboxylation steps, and orange arrows represent reductive carboxylation.
631
632

Figure 1



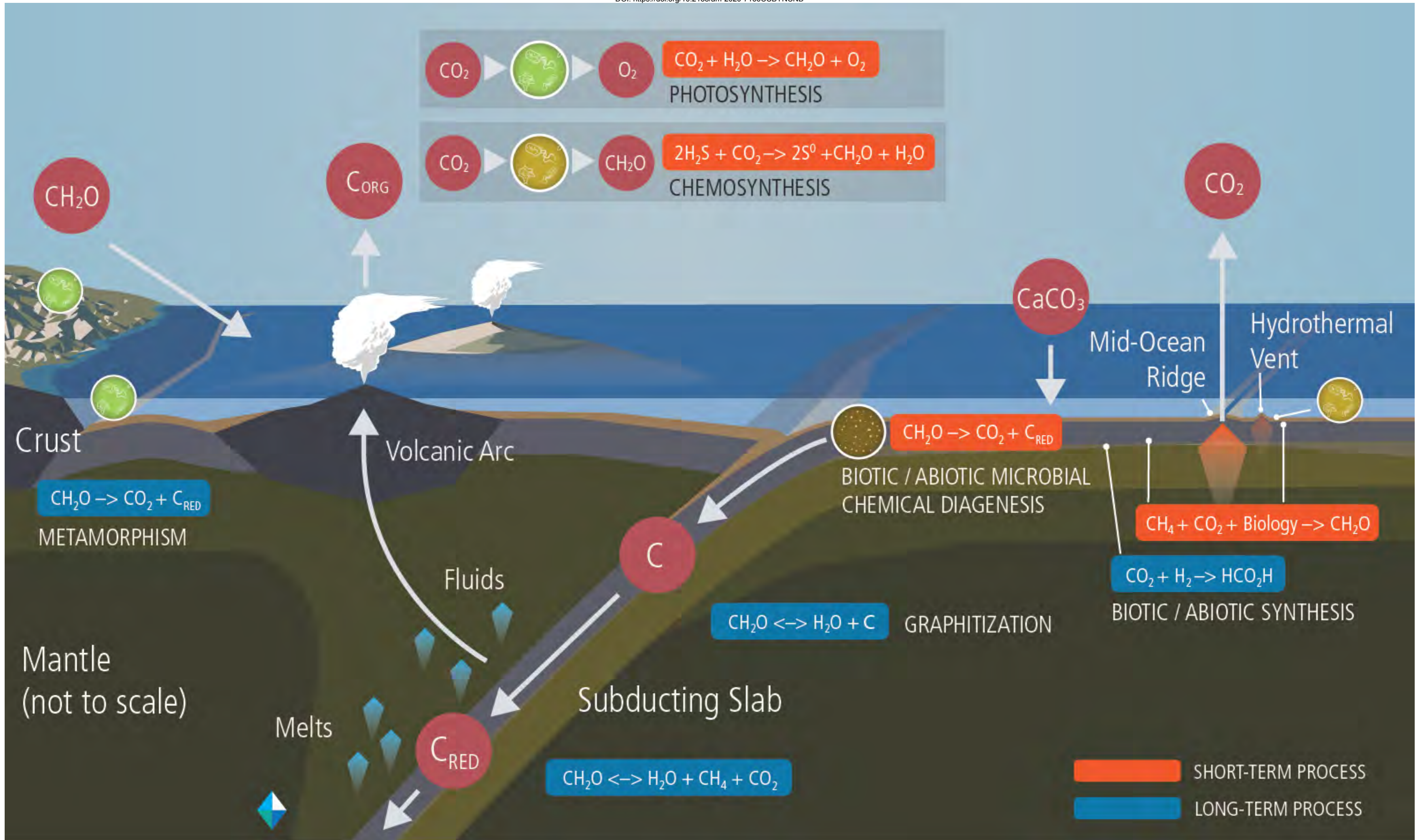


Figure 3

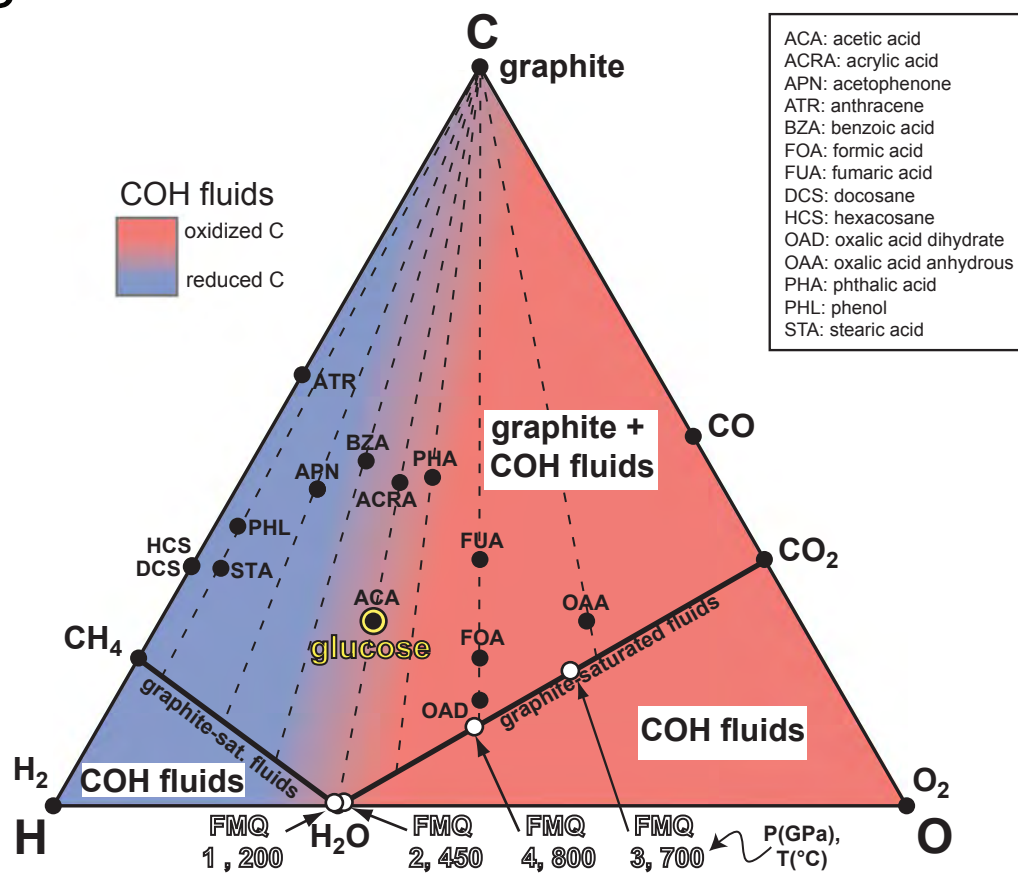


Figure 4

