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Article

Scum of the Earth: a hypothesis for prebiotic multi-compartmentalised environments

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Abstract: Compartmentalisation by bioenergetic membranes is a universal feature of life. The eventual compartmentalisation of prebiotic systems is therefore often argued to comprise a key step during the origin of life. Compartments may have been active participants in prebiotic chemistry, concentrating and spatially organising key reactants. However, most prebiotically plausible compartments are leaky or unstable, limiting their utility. Here, we develop a new hypothesis for an origin of life environment, that capitalises upon, and mitigates the limitations of, prebiotic compartments: multi-compartmentalised layers in the near surface environment — a ‘scum’. Scum-type environments benefit from many of the same ensemble-based advantages as microbial biofilms. In particular, scum layers mediate diffusion with the wider environment, favouring preservation and sharing of early informational molecules, along with the selective concentration of compatible prebiotic compounds. Biofilms are among the earliest traces imprinted by life in the rock record: we contend that prebiotic equivalents of these environments deserve future experimental investigation.

Keywords: Prebiotic chemistry; early Earth; origin of life

1. Introduction

Bulk geochemical environments have several deleterious properties as possible hosts for chemistry that may have given rise to life. Such environments are largely dilute, water-rich, and overly complex — all characteristics that can limit the efficacy of prebiotic reaction pathways. Take the so-called water paradox: the contrast of water as the universal solvent for life with its thermodynamic inhibition of similarly universal condensation reactions involved in e.g., nucleic acid formation [1]. Condensation reactions may be thermodynamically favoured by high concentrations of reactants and a low activity of water. However, high reactant concentrations are often challenging to obtain in dilute bulk solution. This concentration problem is compounded in multi-step reaction paths, whereby a series of linked reversible reactions will produce vanishing small amounts of a desired end-product at equilibrium [2]. Even assuming wholly high yielding, thermodynamically favourable, irreversible reactions, activation energy barriers to forward reaction may be prohibitive in dilute solution.

Extant life elegantly overcomes the water paradox, concentration problem, and kinetic barriers by harnessing enzyme-driven reactions to sustain electrochemical gradients across membranes. The cell membrane is built around an amphiphile bilayer that regulates diffusion, with inward facing hydrophobic groups and outward facing hydrophilic groups. This fundamental relationship between spatial structure, energy flow, and chemical behaviour in extant life draws attention to the possible role of structured environments in fostering prebiotic chemistry, which may be more analogous to extant cells than dilute bulk environments, e.g., mineral/rock pores, microdroplets, amphiphile-based vesicles, and so on [3–6]. Here, we propose a novel end-member prebiotic scenario of this kind:

35 near-surface multi-compartmentalised layers of amphiphile-rich material, which we term the scum
36 hypothesis.

37 2. The scum hypothesis

38 The key distinction of our hypothesis from previous protocell-centric scenarios for prebiotic
39 chemistry is environmental in nature. The prebiotic utility of electrochemical gradients across, and
40 low water activity (a_w) conditions in, free floating proto-cells and amphiphile-rich highly evaporated
41 solutions is long recognised [5,7–12]. Vesicles in particular are known to be valuable for prebiotic
42 chemistry: binding specific reactants and products, initiating chemically productive concentration
43 gradients with respect to the wider environment, and actively participating in key reactions [5,7,10].
44 However, when isolated, most prebiotically plausible vesicles are leaky, struggling to continuously
45 maintain concentration/free energy gradients sufficient to drive prebiotic chemistry [7]. Essentially,
46 the rate of productive reactions driven by the protocellular environment is outstripped by diffusive
47 and reactive sinks related to the bulk environment.

48 We suggest that numerous individual compartments may act together to mitigate such
49 deleterious loss of products to the wider environment, thereby aiding prebiotic chemistry. We
50 focus on multi-compartmentalised near surface environments, scums. Here, an ensemble of leaky
51 compartments achieves the chemical efficacy of individual more functional compartments (with some
52 means of active chemical transport to counter diffusion). We suggest that scum layers may mediate
53 molecular diffusion and water activity to help overcome the concentration problem, arithmetic demon,
54 water paradox, and kinetic barriers to prebiotic chemistry.

55

56 *Prebiotic plausibility of scum layers*

57

58 Many near-surface environments on Earth, including a large fraction of all lakes and oceans, are
59 characterised by a surface microlayer of distinct physical chemistry and composition to an underlying
60 bulk solution [13]. Amphiphiles may preferentially locate in freshwater and sea surface microlayer
61 due to a lower density of oil solutions compared to bulk water. Elevated surface tension at the
62 air-water interface may also energetically favours amphiphile partitioning into the surface microlayer
63 environment [14–16] (Figure 1a). Insolubility and surface-charge adherence can also cause organics to
64 aggregate in the surface microlayer [17]. At high concentrations, these organics can grow into a stable
65 film (Figure 1b-c). However, such organic-rich surface films remain apparently unstructured [17].
66 Stable aggregates of buoyant organics may be further promoted by the arrival of organic-laden bubbles
67 [17]. Surface foam layers may develop if the air-water interfaces of arriving bubbles are stabilised
68 by surfactant molecules (Figure 1d). Free-floating amphiphile-based structures may form in dilute
69 solutions at sufficient concentrations, forming e.g., vesicles [14,18] (Figure 1e).

70

71 Finally, scum layers, the focus of this work, are insoluble organic-rich environments of macroscopic
72 thickness at the water surface (Figure 1f) [19]. Here, we specifically focus on the prebiotic potential of
73 multi-compartmentalised scum layers, consisting predominantly of aggregated vesicles (Figure 1e-f).
74 Vesicles are not the only choice for multi-compartments, but are a plausible candidate. We hypothesise
75 that multi-compartmentalised scum layers may form either by re-organisation of a film or foam into
76 discrete compartments, perhaps in response to changes in bulk composition and/or environmental
77 conditions, e.g., wet-dry cycles, or by the surface accumulation of pre-organised structures, e.g.,
78 buoyant vesicles.

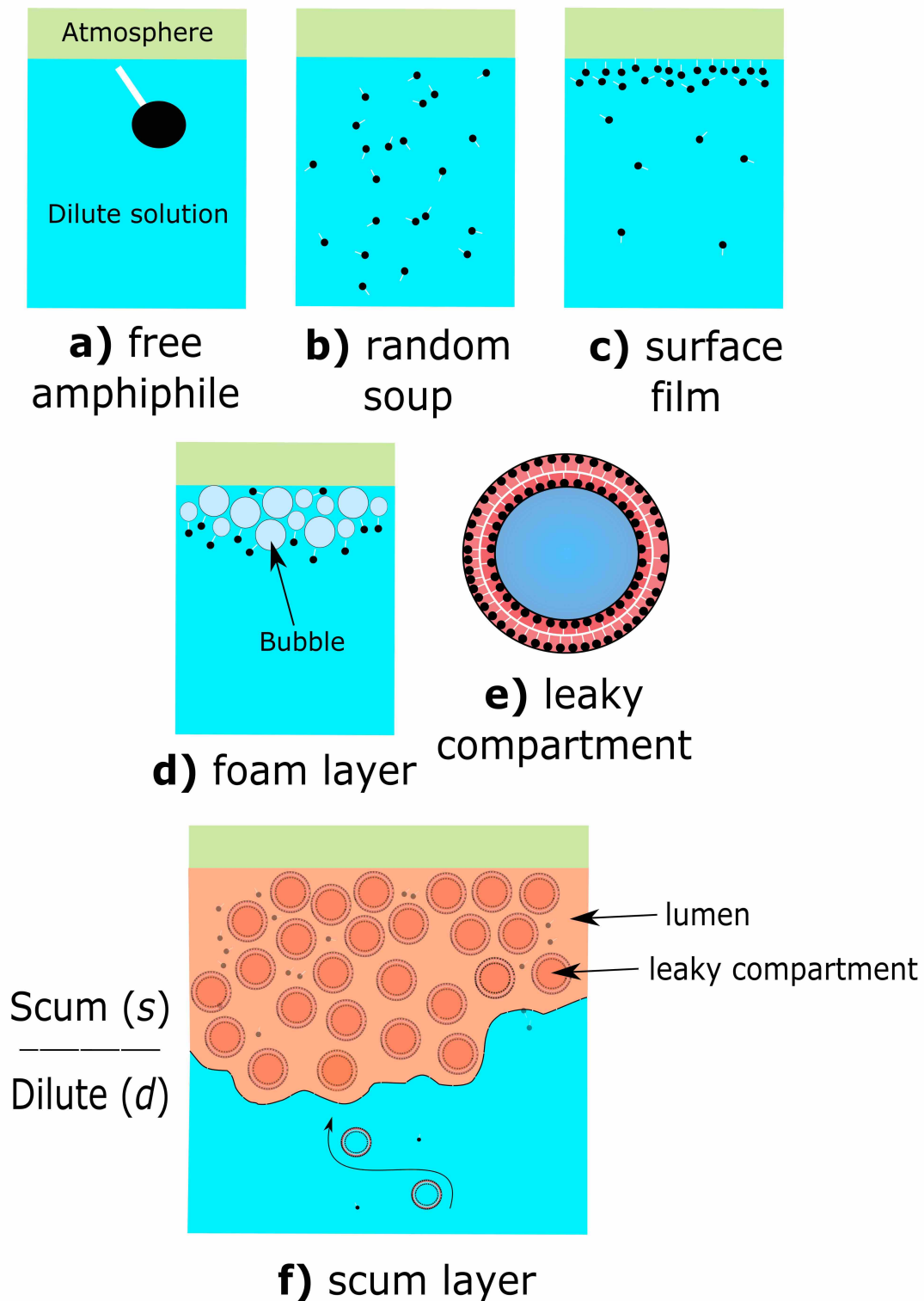


Figure 1. Possible arrangements of simple amphiphiles in a dilute pond. a) Free amphiphile molecule. Black circle represents hydrophilic head, and the white line represents a hydrophobic tail. b) Zoomed out view showing randomly oriented amphiphile molecules in an aqueous environment. c) Partitioning of amphiphiles into the near-surface environment, forming a single or multi-layered surface film. d) Foam layer, where amphiphiles are concentrated along complex air-water interfaces created by gas bubbles. e) Generalised amphiphile-based prebiotic compartment, e.g., vesicle, consisting of an amphiphilic membrane bilayer. f) Scum layer composed of multiple prebiotic compartments, and an interstitial lumen. Thick scum layers may differ noticeably from the wider dilute environment in terms of their overall composition.

78 There are several early Earth environments where scum layers could have accumulated via
 79 these mechanisms. Abiotic amphiphile sources include exogenous fatty acids [20] as well as in-situ
 80 synthesis e.g., HCN-centric activation chemistry leading to lipids [21]. Oceans would have largely
 81 diluted these sources. However, sea foams do form under agitating conditions in the modern ocean,
 82 (speculatively) offering one possible substrate that could have accumulated organics which gone on to
 83 host multi-compartments (Figure 2a-i). Evaporating saline solutions in tidal zones also represent an
 84 opportunity to concentrate amphiphiles (Figure 2a-ii). Only a specific subset of amphiphiles have been
 85 demonstrated to self-assemble in saline conditions [22]. These considerations restrict, but do not rule
 86 out, the possibility for multi-compartmentalised near-surface scum layers in marine settings.

87 A body of theoretical, experimental, and field observations suggest that subaerial restricted basins
 88 could have accumulated high amphiphile concentrations, which then self-assembled into prebiotic
 89 compartments (Figure 2a-iii) [8,23–26]. Accumulation could have occurred directly at the water surface,
 90 owing to the mechanisms described earlier in this paper, or across evaporative cycles [8,25,27]. This
 91 latter mechanism has been observed to produce budding rafts of vesicles [25] — perhaps the closest
 92 experimentally observed structure to the scum layer that we describe.

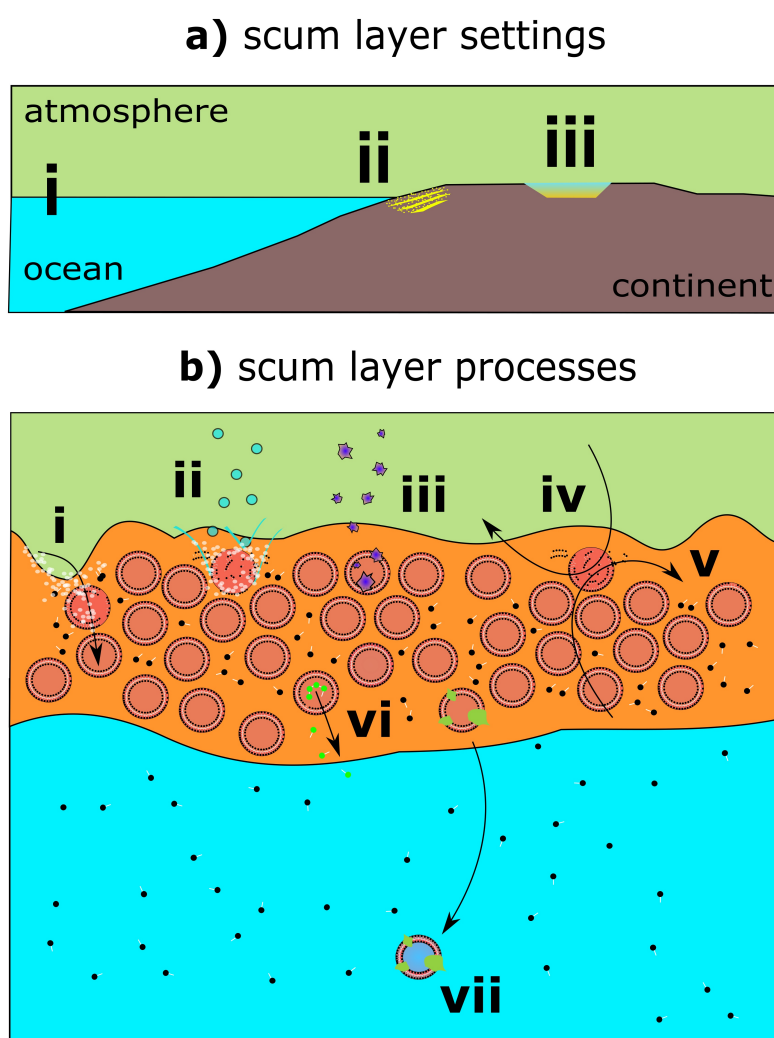


Figure 2. a) scum layer settings — i) open ocean, ii) tidal, iii) restricted basins. b) scum layer processes — i) waves, ii) precipitation, iii) cosmic dust settling, iv) air currents, v) water currents, vi) selective molecular diffusion, vii) mineral precipitation and compartment settling. Disruptive processes are shown intersecting open compartments (red circles, with escaping amphiphiles).

93 Although experimentation and modelling will be needed to test these possibilities, the formation
94 of scum layers on prebiotic Earth does appear plausible in several environmental settings. Indeed,
95 biologically active multi-compartmentalised scum layers, or biofilms, have been abundant on Earth
96 throughout its history [28]. Such biofilms derive competitive advantages from their ensemble structure
97 [29] — many of which point towards prebiotic utility for prebiotic analogue structures.

98 *Prebiotic utility of scum layers*

100
101 **Longevity:** The scum hypothesis posits that multi-compartmentalised and buoyant organic-rich
102 environments may have been stronger candidates than free floating individuals for driving certain
103 reaction pathways in prebiotic chemistry. Much like a modern biofilm, built from constituent cells, a
104 scum layer would have a longer lifespan in aggregate than that of its compartments — and therefore
105 much longer than that of free floating individuals [29]. Ensemble structures will also be more robust
106 to environmental perturbations than individual compartments [29]. Long-lived scum layers would
107 therefore reduce the chance of destructively dispersing prebiotic molecules into the wider dilute
108 environment.

109
110 **Conservation of genetic material:** Conservation of functional/informational polymers would
111 plausible occur, owing to slower overall diffusion between scum multi-compartments and their bulk
112 environment, than would be observed for isolated compartments (see section 3 for our quantitative
113 treatment of this important point). By analogy to the behaviour observed in biofilms, compatible
114 molecules released from a collapsed compartment would have their diffusion slowed by, or even
115 be incorporated into, surrounding compartments [29]. Indeed, biofilms are recognised as ideal
116 environments for the sharing of genetic material [29]. Scum layers would therefore similarly have
117 acted to preserve and disseminate informational polymers [30].

118
119 **Concentration of prebiotic reactants:** Modern surface scum layers display steep compositional
120 gradients with the underlying bulk solution, evidencing their ability to isolate molecules from
121 dilution. Many surface scum layers on lakes are enriched in particle-bound phosphorus by up to
122 100-fold versus dilute solution — a process that would appear to be abiotic in nature, related to
123 interfacial binding of particulates [17]. Such processes may be even more effective in high surface
124 area multi-compartmentalised scum layers. Given that high phosphorus-availability is so critical to
125 many proposed reaction pathways, yet thought to be challenging to obtain on prebiotic Earth [31–33],
126 the enrichment of phosphorus in near-surface scum layers is of potentially great prebiotic relevance.
127 This process could logically extend to the general capture of aerosols, volcanic ash, and fine-grained
128 exogenous materials (e.g., cosmic dust, spherules; Figure 2b-iii), which could then undergo leaching to
129 contribute essential elements for prebiotic chemistry, which are otherwise typically limiting, e.g., P,
130 transition metals, carbon, nitrogen, and sulfur [34].

131
132 **Irradiation exposure and attenuation:** Scum layers would have had access to the most intensely
133 ultraviolet(UV)-irradiated regions of an aqueous environment, which increasingly appears to be a
134 strict requirement for performing prebiotic chemistry on near-biological time-spans [35]. Conversely,
135 rapid UV-attenuation by organics in the scum layer could provide a steep irradiation gradient, i.e.,
136 providing shielding for the products of ongoing prebiotic chemistry [35,36]. The duality of UV
137 exposure and shielding offered by a scum layer may be ideally suited to hosting continuous synthesis
138 pathways that require both UV activity and absence within several reaction steps [33]. Water itself will
139 strongly attenuate UV irradiation intensity within several metres [37]. However, this spatial scale of
140 UV attenuation inherently requires a bulk environment. In contrast, by analogy to organic carbon
141 rich lakes, scum environments may provide shorter-range gradients in both concentration and UV
142 intensity [38]. Meanwhile, the wider environment can serve a different function, both supplying

143 a large reservoir of reactants and removing/diluting those wasteful side products of near-surface
144 chemistry which are not strongly bound to the scum layer.

145

146 **Opportunities for environmental cycling:** A viable environmental setting for prebiotic chemistry
147 also requires a mechanism of selective cycling, i.e., the earliest stages of evolution by natural
148 selection, wherein individual compartments in the scum represent a form of progenote [39]. This
149 competition-based selection may occur in scum layers in a number of ways. Cycling may occur in
150 restricted basin or tidal scenarios during wet-dry cycles. Universally, scum layers would be prone to
151 disruption during episodic agitation by waves (Figure 2b-i), precipitation (Figure 2b-ii), wind (Figure
152 2b-iv) or water currents (Figure 2bv). Length- and charge-dependent diffusion of molecules synthesised
153 in scum compartments into the bulk environment offers a mechanism to select for both certain product
154 molecules and certain types of compartment (Figure 2b-vi). Finally, mineralisation of compartments
155 could cause settling and burial of compartments (Figure 2b-vii). The alternative environment of
156 quiescent ponds would lack many, but not all, of these opportunities for selection. Meanwhile, scum
157 layers in open marine environments could experience any and all forms of disruptive cycling in Figure
158 2b. However, it is questionable whether these interferences would be too severe, with intense winds
159 and large waves instead entirely disrupting scum layers, or indeed prohibiting their initial stabilisation.

160

161 **Caveats:** Standing in the way of these apparent advantages are some important caveats. Modern
162 scum layers are apparently largely unstructured (that is, beyond component films, foams, and biological
163 structures) [17,40]. It therefore remains an open question as to whether prebiotic equivalents would be
164 truly able to self-organise beyond the level of water-air interface structures, and whether near-surface
165 enrichment of e.g., phosphorus can be entirely divorced from ongoing biological activity. The exact
166 degree of UV attenuation by plausible scum layer is also presently unknown, as is their specific ability
167 to concentrate reactants and hold onto products in comparison to isolated vesicles. Overall, there is
168 insufficient evidence at present for positing the scum hypothesis as a holistic scenario for the emergence
169 of life. However, the argument for a potentially constructive role for scum layers in prebiotic chemistry
170 already appears reasonable. In advance of experimental constraints on the detailed prebiotic efficacy
171 of multi-compartmentalised scum layers, we provide-first order calculations to suggest their prebiotic
172 utility by constructing a kinetic model of chemistry taking place in a scum layer.

173 3. A simple model of scum layer chemistry

174 We explore whether scum-type environments have any inherent utility in comparison to isolated
175 compartments for driving prebiotically relevant reactions. Condensation reactions provide a useful
176 case study for these purposes, being ubiquitous in prebiotic chemistry [41]. Take, for example, a
177 generic condensation reaction,



178 The equilibrium position of this condensation reaction is driven to the right by removal of water,
179 where further forward reaction must occur in order to re-attain equilibrium. Several mechanisms
180 may reduce water activity in naturally occurring prebiotic environments. Molecular crowding in a
181 water-poor solution can result in confining cavities of especially low a_w , where molecular desolvation
182 can energetically favour condensation [42]. Secondary effects, such as duplex pairing, may emerge to
183 favour condensation reactions involving nucleotides [43–45]. These effects are all observed for abiotic
184 nucleotide condensation reactions within a range of water activities from 1–0.01 — with confining
185 cavities having much lower water activities in otherwise more dilute bulk solutions [42].

186 We have previously defined multi-compartmentalised scum layers as buoyant accumulations
187 of structured organic molecules, e.g., vesicles, protocells. In addition to further opportunities for the
188 formation of confining spaces, such structured amphiphiles have additional promising properties
189 for driving prebiotic condensation chemistry [46–52]. Such structures have potential for actively

190 participating in prebiotic chemistry, or indeed as model protocells, and have already been studied as
 191 dispersed populations in this context [5,7].

192 It is not the intention of this article to prove that scum-type media are specifically more
 193 advantageous for driving prebiotic condensation reactions than free-floating prebiotic compartments.
 194 However, there are indications that film, foam, and scum layers all have some thermodynamic
 195 potential for hosting (at least) prebiotic condensation reactions. With this basic level of prebiotic
 196 utility justified, we next explore whether the macroscopic nature of a scum layer may be particularly
 197 advantageous for initiating and sustaining dynamic equilibrium — a defining attribute of living cells.

198
 199 *Dynamic equilibrium: the drive to life*

200
 201 Cells must protect biomolecules (e.g., DNA) against degradation by hydrolysis. This protection is
 202 ultimately achieved by cycling between the cellular environment and dehydrating microenvironments
 203 formed during enzyme-substrate binding [53]. Such continuous cycling results in cell-wide dynamic
 204 equilibrium, i.e., steady-state, but not thermochemical equilibrium with the wider environment [54].
 205 A key example of dynamic equilibrium is the concentration gradient that universally spans cell
 206 membranes. Fluxes between hydrolysis and dehydration in the cell are balanced overall, yet free
 207 energy is made available to exploit on local scales via microenvironmental shifts [55]. Without the
 208 intervention of enzymes and the bioenergetic membrane, a spatially and temporally homogeneous
 209 thermochemical equilibrium would be achieved — corresponding to the cessation of metabolism and
 210 other key regulatory activities. A prebiotic system that is incapable of sustaining dynamic equilibria
 211 therefore faces a major chemical distinction compared with what we know as life.

212 Total chemical energy available to do work (G_{scum} , J) in the system can be treated as the sum of
 213 the products of chemical potentials (μ , J mol⁻¹), stoichiometric coefficients (d), and total moles (n) of
 214 all species (i) undergoing reaction,

$$G_{scum} = \sum_i \mu_i d_i n_i \quad (2)$$

215 In systems that reach thermochemical equilibrium, equation 2 will sum to zero. However, other
 216 systems may drive balanced yet continuous fluxes of dehydrated and hydrolysed products into
 217 hydrolysing and dehydrating environments, respectively, resulting in non-zero values of G_{scum} , i.e.,
 218 thermochemical disequilibrium. At a steady state, this scenario may also be referred to as a dynamic
 219 equilibrium.

220 Consider the case of a selectively leaky prebiotic compartment, which can spontaneously drive
 221 forward condensation reaction, owing to e.g., a low internal water activity, and which is also
 222 undergoing diffusion-driven exchange with its surroundings. Thermochemical equilibrium in the
 223 compartment can only be achieved given either (1) that the compartment is a closed system, which
 224 lacks diffusive exchange with dilute surroundings, or (2) the surroundings are non-dilute: product
 225 can build up and diffuse back into the compartment at a rate such that an equilibrium position for the
 226 reaction is achieved. As the compartment becomes selectively leaky, maintaining the ability to induce
 227 forward reaction but more rapidly losing any product generated, a dynamic equilibrium is achieved.
 228 Compartments will approach a dilute limit once diffusion removes product as fast as it is generated
 229 in. At this dilute limit, the compartment is furthest from being at local thermochemical equilibrium:
 230 reactant and product concentrations will reflect those in the wider environment, and equation 2 will be
 231 at a maximum.

232 A valuable prebiotic environment has the the ability, like life, to exist in between these two
 233 states: in a steady-state where thermochemical equilibrium with the wider environment is never
 234 achieved, yet a moderate-to-high product concentration is maintained in the compartment in the
 235 face of diffusion. This scenario might be most easily found in semi-permeable environments that
 236 host exchanges of material between water-poor and water-rich micro-environments, e.g., in putative

237 functional semi-permeable protocells, or potentially across multi-compartmentalised scum layers
 238 built from individually leaky vesicles. A prebiotic system that recalls extant life is therefore defined
 239 by a delicate balance of forward and reverse reaction rates and molecular diffusion, which can be
 240 quantified.

241 We construct a simple kinetic model of condensation chemistry and molecular diffusion taking
 242 place in a scum layer (Figure 3). We solve for steady-state. We consider a wider environment (dilute,
 243 d) and a second multi-compartmentalised environment of prebiotic interest (scum, s). The change in
 244 concentration of products in environment s ($\Delta[\text{CX}]$, mol s^{-1}) must always be zero at steady state, and
 245 can be represented as the balance of product sources and sinks. Sources include forward reaction to
 246 form product (f_f , mol s^{-1}). Sinks include reverse reaction of product to form reactants (f_r , mol s^{-1}).
 247 Diffusion of product can occur either from scum to dilute environment, or vice versa, and may
 248 therefore represent either a source or a sink depending on the direction of diffusion being considered
 249 f_{diff} (mol s^{-1}).

$$\Delta[\text{CX}]^s = 0 = f_f - f_r \pm f_{diff}. \quad (3)$$

250 Product source and sink fluxes can be written as the product of environmental volumes (V^{env} , L)
 251 with rates of reaction ($r_{f,r}$, $\text{mol s}^{-1} \text{L}^{-1}$) and diffusion (r_{diff} , $\text{mol s}^{-1} \text{L}^{-1}$). We assume that forward
 252 reaction 1 can only proceed spontaneously in compartments (V_c^s). Reverse reaction can take place
 253 in both the compartments and interstitial lumen (Figure 3). A comprehensive treatment of scum
 254 diffusion and reaction behaviour is beyond the scope of this article. In order to simplify our model,
 255 we consider a selectively leaky scum where reactant concentrations are equal throughout scum and
 256 the dilute environment, allowing us to prescribe final reactant concentrations. Diffusive exchange
 257 fluxes will depend on the surface areas of contact between scum and dilute environment which, for
 258 a fixed thickness of scum, scales with scum volume (V^s). We can then explore how varying product
 259 forward and reverse reaction and diffusion rates affect steady-state concentrations of product in scum
 260 compartments, writing that

$$0 = r_f^s V_c^s - r_r^s V^s + V^s (r_{diff}^{d-s} - r_{diff}^{s-d}). \quad (4)$$

261 Even though there is no spontaneous forward reaction in environment d , there may be diffusion of
 262 product from environment s into environment d . Given that the dilute environment is of much greater
 263 volume than the scum layer, we assume that $f_r^d \gg f_{diff}^{s-d}$, such that the concentration of product in
 264 environment d is approximately zero at steady state. We can simplify our equation for product sources
 265 and sinks in the scum layer accordingly,

$$0 = r_f^s V_c^s - r_r^s V^s - f_{diff}^{s-d}. \quad (5)$$

266 At equilibrium, the rate of product diffusion from environment s in our model scenario cannot
 267 be greater than the rate of forward reaction, i.e., every product molecule produced is balanced via
 268 diffusion of a product molecule to environment d . In this case we can express product diffusion as
 269 some scaled value of the forward reaction rate,

$$f_{diff}^{s-d} = f_f K_{diff}. \quad (6)$$

270 The diffusion scaling constant K_{diff} can vary from 1 (a diffusion flux balancing forward reaction)
 271 to 0 (no diffusion from environment s). Finally, to obtain the forward and reverse reaction rates,
 272 for the general condensation reaction 1, we can write rate laws using model values for forward
 273 (k_f , $\text{L mol}^{-1} \text{s}^{-1}$) and reverse (k_r , s^{-1}) reaction rate constants,

$$r_f = k_f [\text{COH}] [\text{HX}], \quad (7)$$

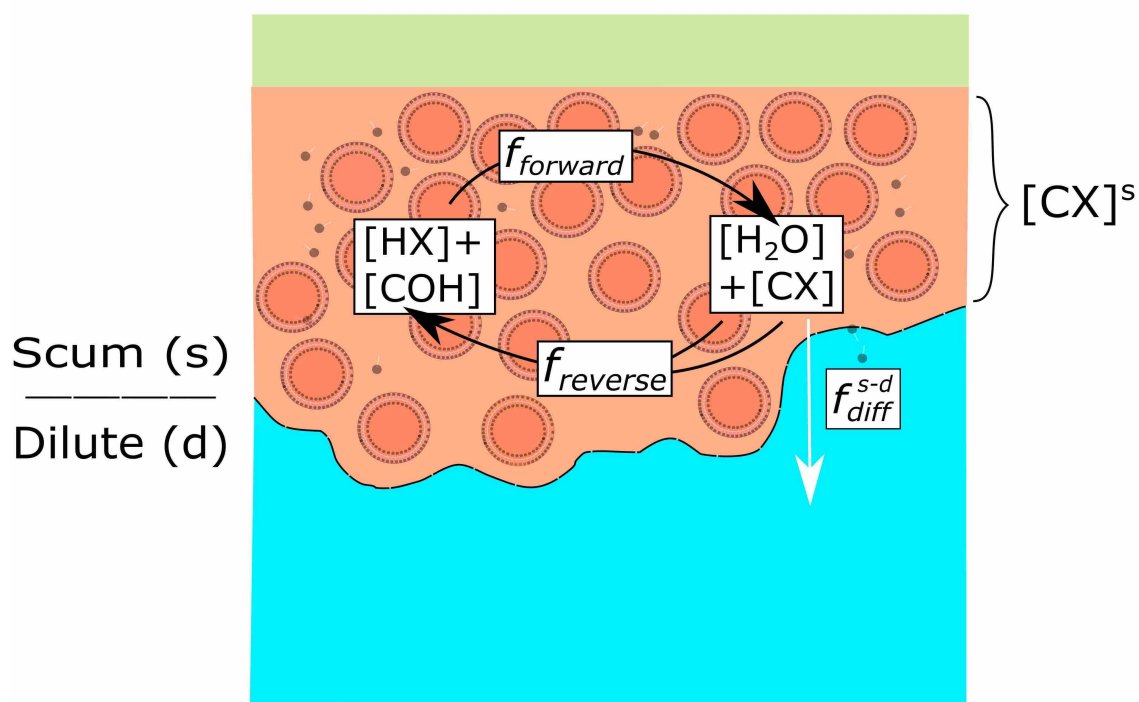


Figure 3. Schematic of fluxes affecting product concentrations in scum environment (s). Reverse reaction flux of product is the sum of the flux from reactions occurring in both compartments and interstitial fluid in the scum layer, and therefore varies as a function of total scum volume. Forward reaction only takes place in compartments.

$$r_r = k_r [\text{CX}] a_w. \quad (8)$$

274 Substituting for unknowns, we can then rearrange equation 5 and solve for the steady-state
275 concentration of product in the scum layer,

$$[\text{CX}] = \frac{k_f [\text{COH}] [\text{HX}] \frac{V_c^s}{V^s} (1 - K_{diff})}{k_r a_w}. \quad (9)$$

276 Using this information in conjunction with eq. 2, we can calculate available chemical energy in
277 environment *s* — our measure of whether the equilibrium achieved is more dynamic or thermochemical
278 in nature — as,

$$G_{scum} = \mu_{\text{CX}} V^s [\text{CX}] + \mu_{\text{HX}} V^s [\text{HX}] + \mu_{\text{COH}} V^s [\text{COH}] + \mu_{\text{H}_2\text{O}} V^s [\text{H}_2\text{O}]. \quad (10)$$

279 We use favorable values for rate constants and values for free energies of reaction that characterise
280 at least some condensation reactions in desolvating (low a_w) conditions [56,57].

281 We assume $k_f = 10 \text{ L mol}^{-1} \text{ s}^{-1}$, $k_r = 1 \text{ s}^{-1}$, μ_{COH} and $\mu_{\text{HX}} = 100 \text{ kJ mol}^{-1}$, $\mu_{\text{CX}} = -2.5 \text{ kJ mol}^{-1}$,
282 $a_w = 0.1$ (assuming open system water loss, with water produced during condensation immediately
283 lost to the environment), $[\text{COH}]$ and $[\text{HX}] = 1 \text{ mM}$, and volume = 1 L in environment *s*. We explore a
284 range of scum layer multi-compartmentalisation densities, from $\frac{V_c^s}{V^s} = 0.01$, i.e., relatively compartment
285 poor, with a high amount of interstitial fluid, to 0.8, i.e., almost entirely dominated by compartments
286 in which forward reaction can spontaneously occur. As water is present on both sides of the generic
287 condensation reaction being considered, water chemical potential is excluded from this calculation.

288 Using these assumptions, we calculate G_{scum} and equilibrium condensation product concentration
 289 for a range of diffusion rates between dilute solution (environment d) and a scum layer (environment
 290 s). The results of this calculation are shown in Figure 4. Boundary conditions of thermochemical
 291 equilibrium and the dilute limit are explicitly indicated.

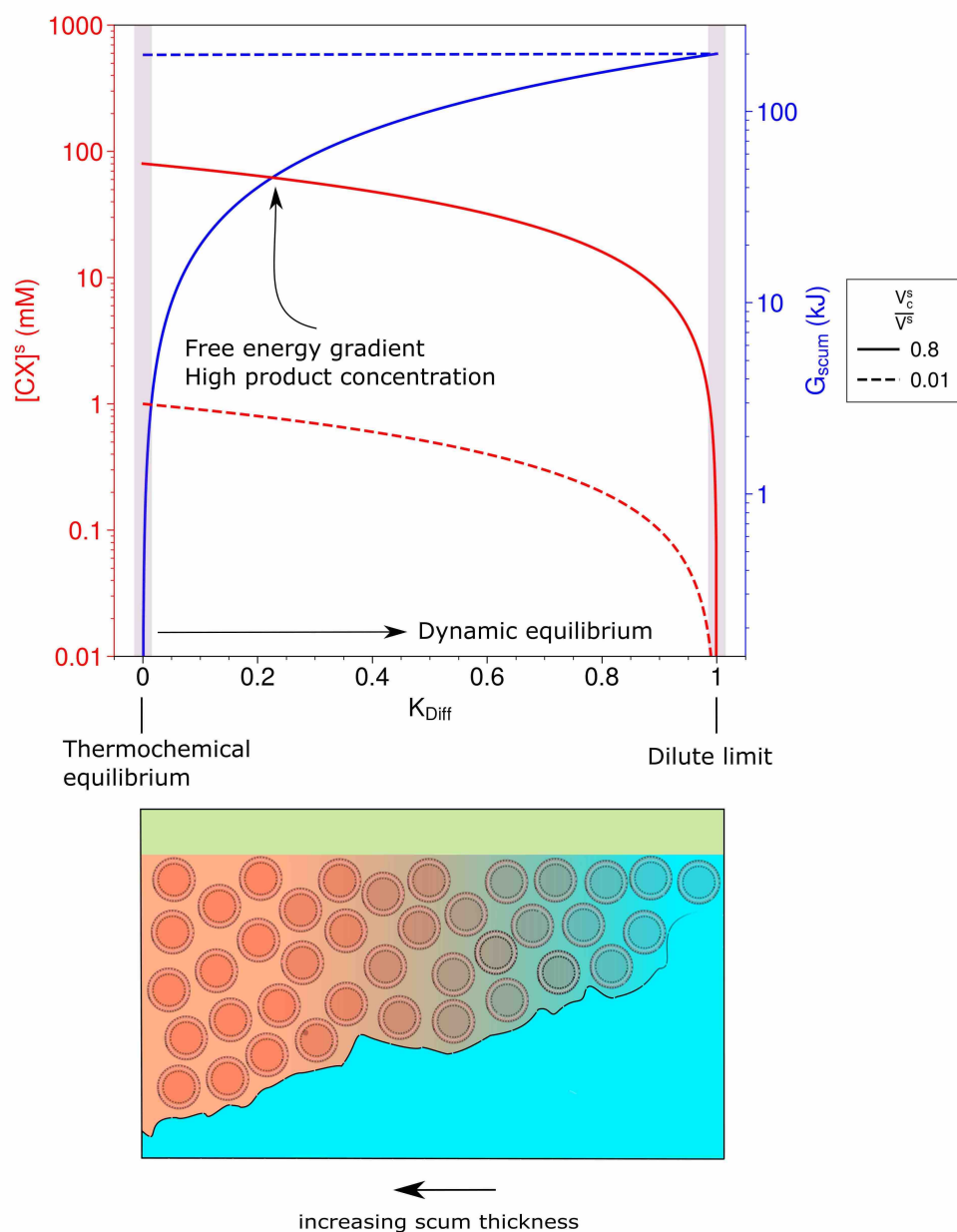


Figure 4. Effect of increasing diffusion scaling constant and reactive compartment volume on equilibrium concentration of product and available energy in environment s . Origin scenarios are mapped onto this scale, with both scum layers and functional semi-permeable protocells shown to bridge the gap between fully thermochemical and dynamic equilibrium.

292 If diffusion is rapid enough to remove the majority of product (K_{Diff} greater than 0.5), the dynamic
 293 equilibrium approached is one where the free energy available to do chemical work in environment s is
 294 at its maximum. As the rate of diffusion between environment s and d is set to lower values (Figure 4),
 295 available energy in environment s decreases but the equilibrium concentration of product rises. On the
 296 other extreme, no diffusion between the two environments will result in thermochemical equilibrium
 297 in environment s : zero available free energy, but maximum product availability. Maintaining dynamic

equilibrium alongside high product concentrations is achievable only given an intermediate diffusion flux from environments s to d . Such a system, which has access to an external environment and yet can maintain concentration gradients against it owing to (thermodynamic and/or kinetic) facilitation of forward reaction, may allow prebiotic reactions to proceed with constant supply — potentially helping to beat universal problems in prebiotic chemistry, such as the arithmetic demon [2].

The diffusion flux of product from a scum to dilute environment depends on factors such as the energy required to cross the interfaces between environment s and d , the surface area of that interface, the concentration of product in either environment, and the respective volume ratio of the two environments (Figure 4) [58]. Intermediate diffusion fluxes may be observed around a very small environment with a difficult-to-cross interface, e.g., a semi-permeable membrane, as in cells or functional protocells, or in a larger environment built from many individually leaky compartments, e.g., a multi-compartmentalised scum layer.

The reactive compartment density of the scum layer also plays a critical role in determining behavior, with low ratios of reactive compartment to total scum volume ($\frac{V_c^s}{V^s}$) failing to yield high product concentrations even at thermochemical equilibrium (Figure 4), i.e., the system remains detrimentally leaky. This requirement places another constraint on which types of scum layer may have prebiotic utility. However, assuming that leaky compartments are more feasible to assemble on early Earth than biology-adjacent protocells [7], utilising their enhanced aggregate properties via (compartment dense) scum layers would seem to be a viable route for prebiotic chemistry.

Our work shows that free-floating selectively leaky prebiotic compartments exist close to a dilute limit - very far from thermochemical equilibrium, but unable to sustain viable concentration gradients of any product molecules that they may help to spontaneously produce. Meanwhile, multi-compartmentalised scum layers, comprised of functional compartments and interstitial lumen, which together exhibit lower effective rates of diffusion with the dilute environment, provide a possible solution to this dilution problem from the perspective of physical chemistry, deserving of experimental exploration.

4. Summary and next steps

The origin of life involved linking metabolism, information, and compartmentalisation. In modern cells, bioenergetic membranes maintain the concentration gradients that underlie energy transfer and prevent information from diffusing away. It is possible that fully formed prebiotic analogues emerged with equivalents of these relationships. Alternatively, simpler structures, lacking the ability to actively regulate molecular transport, may have accumulated in near-surface multi-compartmentalised scum environments, benefiting in aggregate from a slower rate of diffusive exchange with the wider environment. We argue that such environments may plausibly have maintained dynamic equilibrium with their environments, without approaching the dilute limit of molecular exchange (Figure 4). In this scenario, dependent upon an external energy source, chemical work is continuously performed, yet useful product concentrations are also maintained against environmental dilution. Scum layers could therefore be a constructive interference with early prebiotic systems, potentially helping to reduce the challenge of regulating molecular diffusion in otherwise dilute prebiotic environments.

The formation of amphiphile-rich film/foam layers and free floating amphiphile-based structures on prebiotic Earth is already recognised as plausible [8,23–25,59]. Multi-compartmentalised scum layers could have emerged on prebiotic Earth via continuous growth and stabilisation of surface films/foams, followed by further self-organisation, or by direct surface/membrane-adherence of pre-formed buoyant structures. Experimental investigation will be required to determine which formation pathways are valid, and which scum types are of prebiotic utility.

Our hypothesis predicts that multi-compartmentalised scum-layers can form during either surface microlayer partitioning or buoyancy-driven aggregation of prebiotically plausible amphiphiles. Experimental work should first establish that prebiotic scum-type structures are plausible, by exploring the physical chemistry of various amphiphile-rich solutions (vesicle-bearing, initially unstructured,

and so on) under a range of conditions (quiescence, bubbling, currents, and so on). The formation of unstructured scum layers seems prebiotically plausible on current evidence: the requirements for multi-compartmentalisation, meanwhile, are less obvious. It could be argued that surface-charge interactions may act to repel close-packed prebiotic compartments that are chemically similar to one another. It is therefore possible that distinct membrane types may be required to overcome such electrostatic instabilities, or perhaps mixed systems will also be unstable, leading to unstructured systems of questionable prebiotic utility. We recommend, and intend to pursue, an experimental effort focused on the critical issue of scum formation pathways. If positive results are obtained for prebiotic plausibility of scum layers, future experiments can then begin to consider questions about prebiotic utility.

Scum-type media should be tested for their ability to concentrate reactants from a dilute bulk environment, promote forward reaction, and selectively concentrate product molecules of interest. Results should be compared to those obtained using a mixture of free-floating prebiotic compartments. Important tests include the ability of scum layers to (1) concentrate phosphate and drive phosphorylation reactions, (2) harness a combination of surface scum UV irradiation and juxtaposed UV shielding to drive prebiotic chemistry, (3) drive general condensation polymerisation chemistry, as well as provide selection mechanisms that favour accumulation of longer/more useful products [60], and (4) accumulate and activate fine-grained exogenous materials for use in prebiotic reactions, e.g., photocapture by meteoritic PAH pigments/quinone systems, coupled to nucleotide, amino acid, and lipid synthesis pathways [61–64].

Finally, the ability of scum layers to selectively concentrate molecules of increasing size should be considered. Longer molecules should diffuse more slowly across semi-permeable membranes [65], implying a steady approach towards local thermochemical equilibrium for longer polymers forming in scum-type environments. This behaviour could prove advantageous for maintaining high yields of long chain polymers without needing to invoke a closed system, i.e., as may characterised scum environments. It is currently unknown to what extent molecular diffusion rates and forward reaction rates compare between free floating protocells and scum-type environments. Experimental work is needed to place constraints on these crucial parameters.

Whilst we have pointed to the probable compatibility of scum layers with broadly important condensation reactions, there are many outstanding issues about the diversity, efficiency, and selectivity of prebiotic chemistry taking place in scum layers, as well as the overall placement of these environments in testable scenarios for the origin of life. Despite these outstanding issues, we note that buoyant bacterial communities have a deep ancestry on our planet — representing the earliest known biological structures in the rock record [28]. We contend that prebiotic equivalents of these environments deserve future experimental and theoretical investigation.

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6. References

- Deal, A.M.; Rapf, R.J.; Vaida, V. Water–Air Interfaces as Environments to Address the Water Paradox in Prebiotic Chemistry: A Physical Chemistry Perspective. *The Journal of Physical Chemistry A* **2021**, *125*, 4929–4942. doi:10.1021/acs.jpca.1c02864.
- Crispino, G.; Ho, P.; Sharpless, K. Selective perhydroxylation of squalene: taming the arithmetic demon. *Science* **1993**, *259*, 64–66. doi:10.1126/science.8418495.

- 393 3. Cartwright, J.H.E.; Russell, M.J. The origin of life: the submarine alkaline vent theory at 30. *Interface Focus*
394 **2019**, *9*, 20190104. doi:10.1098/rsfs.2019.0104.
- 395 4. Jia, T.Z.; Chandru, K.; Hongo, Y.; Afrin, R.; Usui, T.; Myojo, K.; Cleaves, H.J. Membraneless polyester
396 microdroplets as primordial compartments at the origins of life. *Proceedings of the National Academy of*
397 *Sciences* **2019**, *116*, 15830–15835. doi:10.1073/pnas.1902336116.
- 398 5. Bonfio, C.; Russell, D.A.; Green, N.J.; Mariani, A.; Sutherland, J.D. Activation chemistry drives the
399 emergence of functionalised protocells. *Chemical Science* **2020**. doi:10.1039/d0sc04506c.
- 400 6. Morasch, M.; Liu, J.; Dirscher, C.F.; Ianeselli, A.; Kühnlein, A.; Le Vay, K.; Schwintek, P.; Islam, S.; Corpinot,
401 M.K.; Scheu, B.; Dingwell, D.B.; Schwille, P.; Mutschler, H.; Powner, M.W.; Mast, C.B.; ; Braun, D. Heated
402 gas bubbles enrich, crystallize, dry, phosphorylate and encapsulate prebiotic molecules. *Nature Chemistry*
403 **2019**, pp. –. doi:doi.org/10.1038/s41557-019-0299-5.
- 404 7. Bonfio, C.; Godino, E.; Corsini, M.; Biani, F.F.d.; Guella, G.; Mansy, S.S. Prebiotic iron–sulfur peptide
405 catalysts generate a pH gradient across model membranes of late protocells. *Nature Catalysis* **2018**,
406 *1*, 616–623. doi:10.1038/s41929-018-0116-3.
- 407 8. Deamer, D.; Damer, B.; Kompanichenko, V. Hydrothermal Chemistry and the Origin of Cellular Life.
408 *Astrobiology* **2019**, *19*, 1523–1537. doi:10.1089/ast.2018.1979.
- 409 9. Deamer, D. Liquid crystalline nanostructures : organizing matrices for non-enzymatic nucleic acid
410 polymerization. *Chemical Society Reviews* **2012**, *41*, 5375–5379. doi:10.1039/c2cs35042d.
- 411 10. Milshteyn, D.; Cooper, G.; Deamer, D. Chemiosmotic energy for primitive cellular life: Proton gradients
412 are generated across lipid membranes by redox reactions coupled to meteoritic quinones. *Scientific Reports*
413 **2019**, *9*, 12447. doi:10.1038/s41598-019-48328-5.
- 414 11. Milshteyn, D.; Damer, B.; Havig, J.; Deamer, D. Amphiphilic Compounds Assemble into Membranous
415 Vesicles in Hydrothermal Hot Spring Water but Not in Seawater. *Life* **2018**, *8*, 11. doi:10.3390/life8020011.
- 416 12. Rajamani, S.; Vlassov, A.; Benner, S.; Coombs, A.; Olasagasti, F.; Deamer, D. Lipid-assisted Synthesis
417 of RNA-like Polymers from Mononucleotides. *Origins of Life and Evolution of Biospheres* **2008**, *38*, 57–74.
418 doi:10.1007/s11084-007-9113-2.
- 419 13. Wurl, O.; Wurl, E.; Miller, L.; Johnson, K.; Vagle, S. Formation and global distribution of sea-surface
420 microlayers. *Biogeosciences* **2011**, *8*, 121–135. doi:10.5194/bg-8-121-2011.
- 421 14. Walde, P. Surfactant Assemblies and their Various Possible Roles for the Origin(S) of Life. *Origins of Life*
422 *and Evolution of Biospheres* **2006**, *36*, 109–150. doi:10.1007/s11084-005-9004-3.
- 423 15. Engel, A.; Sperling, M.; Sun, C.; Grosse, J.; Friedrichs, G. Organic Matter in the Surface Microlayer:
424 Insights From a Wind Wave Channel Experiment. *Frontiers in Marine Science* **2018**, *5*, 182.
425 doi:10.3389/fmars.2018.00182.
- 426 16. Ćosović, B. Surface-Active Properties of the Sea Surface Microlayer and Consequences for Pollution in the
427 Mediterranean Sea. In *The Mediterranean Sea*; Saliot, A., Ed.; Springer Berlin Heidelberg: Berlin, Heidelberg,
428 2005; pp. 269–296. doi:10.1007/b107150.
- 429 17. Södergren, A. Role of aquatic surface microlayer in the dynamics of nutrients and organic compounds in
430 lakes, with implications for their ecotones. *Hydrobiologia* **1993**, *251*, 217–225. doi:10.1007/bf00007181.
- 431 18. Yanagawa, H.; Ogawa, Y.; Kojima, K.; Ito, M. Construction of protocellular structures under
432 simulated primitive earth conditions. *Origins of life and evolution of the biosphere* **1988**, *18*, 179–207.
433 doi:10.1007/bf01804670.
- 434 19. Clark, B.C.; Kolb, V.M. Comet Pond II: Synergistic Intersection of Concentrated Extraterrestrial
435 Materials and Planetary Environments to Form Procreative Darwinian Ponds. *Life* **2018**, *8*, 12.
436 doi:10.3390/life8020012.
- 437 20. Lawless, J.G.; Yuen, G.U. Quantification of monocarboxylic acids in the Murchison carbonaceous meteorite.
438 *Nature* **1979**, *282*, 396–398. doi:10.1038/282396a0.
- 439 21. Liu, Z.; Wu, L.F.; Xu, J.; Bonfio, C.; Russell, D.A.; Sutherland, J.D. Harnessing chemical energy
440 for the activation and joining of prebiotic building blocks. *Nature Chemistry* **2020**, *12*, 1023–1028.
441 doi:10.1038/s41557-020-00564-3.
- 442 22. Jordan, S.F.; Ramm, H.; Zheludev, I.N.; Hartley, A.M.; Maréchal, A.; Lane, N. Promotion of protocell
443 self-assembly from mixed amphiphiles at the origin of life. *Nature Ecology & Evolution* **2019**, *3*, 1705–1714.
444 doi:10.1038/s41559-019-1015-y.

- 445 23. Damer, B.; Deamer, D. Coupled Phases and Combinatorial Selection in Fluctuating Hydrothermal Pools:
446 A Scenario to Guide Experimental Approaches to the Origin of Cellular Life. *Life* **2015**, *5*, 872–887.
447 doi:10.3390/life5010872.
- 448 24. Damer, B. A Field Trip to the Archaean in Search of Darwin’s Warm Little Pond. *Life* **2016**, *6*, 21.
449 doi:10.3390/life6020021.
- 450 25. Damer, B.; Deamer, D. The Hot Spring Hypothesis for an Origin of Life. *Astrobiology* **2020**, *20*, 429–452.
451 doi:10.1089/ast.2019.2045.
- 452 26. Deamer, D. Where Did Life Begin? Testing Ideas in Prebiotic Analogue Conditions. *Life* **2021**, *11*, 134.
453 doi:10.3390/life11020134.
- 454 27. Trevors, J.T.; Pollack, G.H. Hypothesis: the origin of life in a hydrogel environment. *Progress in Biophysics*
455 *and Molecular Biology* **2005**, *89*, 1–8. doi:10.1016/j.pbiomolbio.2004.07.003.
- 456 28. Allwood, A.C.; Walter, M.R.; Burch, I.W.; Kamber, B.S. 3.43 billion-year-old stromatolite reef from the
457 Pilbara Craton of Western Australia: Ecosystem-scale insights to early life on Earth. *Precambrian Research*
458 **2007**, *158*, 198–227. doi:10.1016/j.precamres.2007.04.013.
- 459 29. Jefferson, K. What drives bacteria to produce a biofilm? *FEMS Microbiology Letters* **2004**, *236*, 163–173.
460 doi:10.1016/j.femsle.2004.06.005.
- 461 30. Bernhardt, H.S. The RNA world hypothesis: the worst theory of the early evolution of life (except for all
462 the others). *Biology Direct* **2012**, *23*.
- 463 31. Schwartz, A.W. Phosphorus in prebiotic chemistry. *Philosophical Transactions of the Royal Society B: Biological*
464 *Sciences* **2006**, *361*, 1743–1749. doi:10.1098/rstb.2006.1901.
- 465 32. Graaf, R.M.d.; Schwartz, A.W. Reduction and Activation of Phosphate on the Primitive Earth. *Origins of*
466 *life and evolution of the biosphere* **2000**, *30*, 405–410. doi:10.1023/a:1006700512902.
- 467 33. Patel, B.H.; Percivalle, C.; Ritson, D.J.; Duffy, C.D.; Sutherland, J.D. Common origins of RNA,
468 protein and lipid precursors in a cyanosulfidic protometabolism. *Nature Chemistry* **2015**, *7*, 301–307.
469 doi:10.1038/nchem.2202.
- 470 34. Maurette, M.; Brack, A.; Kurat, G.; Perreau, M.; Engrand, C. Were micrometeorites a source
471 of prebiotic molecules on the early Earth? *Advances in Space Research* **1995**, *15*, 113–126.
472 doi:10.1016/s0273-1177(99)80071-4.
- 473 35. Rimmer, P.B.; Thompson, S.J.; Xu, J.; Russell, D.A.; Green, N.J.; Ritson, D.J.; Sutherland, J.D.; Quelo, D.P.
474 Timescales for Prebiotic Photochemistry Under Realistic Surface Ultraviolet Conditions. *Astrobiology* **2021**.
475 doi:10.1089/ast.2020.2335.
- 476 36. Cleaves, H.J.; Miller, S.L. Oceanic protection of prebiotic organic compounds from UV radiation. *Proceedings*
477 *of the National Academy of Sciences* **1998**, *95*, 7260–7263.
- 478 37. Fleischmann, E.M. The measurement and penetration of ultraviolet radiation into tropical marine water.
479 *Limnology and Oceanography* **1989**, *34*, 1623–1629. doi:10.4319/lo.1989.34.8.1623.
- 480 38. Pilla, R.M.; Couture, R. Attenuation of photosynthetically active radiation and ultraviolet radiation
481 in response to changing dissolved organic carbon in browning lakes: Modeling and parametrization.
482 *Limnology and Oceanography* **2021**, *66*, 2278–2289. doi:10.1002/lno.11753.
- 483 39. Woese, C.R.; Fox, G.E. The concept of cellular evolution. *Journal of Molecular Evolution* **1977**, *10*, 1–6.
484 doi:10.1007/bf01796132.
- 485 40. Wotton, R.S.; Preston, T.M. Surface Films: Areas of Water Bodies That Are Often Overlooked. *BioScience*
486 **2005**, *55*, 137–145.
- 487 41. Hulshof, J.; Ponnampereuma, C. Prebiotic Condensation Reactions in an aqueous medium: a review of
488 condensing agents. *Origins of life* **1976**, *7*, 197–224.
- 489 42. Ross, D.S.; Deamer, D. Dry/Wet Cycling and the Thermodynamics and Kinetics of Prebiotic Polymer
490 Synthesis. *Life* **2016**, *6*, 28. doi:10.3390/life6030028.
- 491 43. Butcher, S.E.; Pyle, A.M. The Molecular Interactions That Stabilize RNA Tertiary Structure: RNA Motifs,
492 Patterns, and Networks. *Accounts of Chemical Research* **2011**, *44*, 1302–1311. doi:10.1021/ar200098t.
- 493 44. Jaeger, J.A.; Turner, D.H.; Zuker, M. Improved predictions of secondary structures for RNA. *Proceedings of*
494 *the National Academy of Sciences* **1989**, *86*, 7706–7710. doi:10.1073/pnas.86.20.7706.
- 495 45. Freier, S.M.; Kierzek, R.; Jaeger, J.A.; Sugimoto, N.; Caruthers, M.H.; Neilson, T.; Turner, D.H. Improved
496 free-energy parameters for predictions of RNA duplex stability. *Proceedings of the National Academy of*
497 *Sciences* **1986**, *83*, 9373–9377. doi:10.1073/pnas.83.24.9373.

- 498 46. Disalvo, E.; Pinto, O.; Martini, M.; Bouchet, A.; Hollmann, A.; Frías, M. Functional role of water in
499 membranes updated: A tribute to Träuble. *Biochimica et Biophysica Acta (BBA) - Biomembranes* **2015**,
500 *1848*, 1552–1562. doi:10.1016/j.bbamem.2015.03.031.
- 501 47. Disalvo, E.; Martini, M.; Bouchet, A.; Hollmann, A.; Frías, M. Structural and thermodynamic properties of
502 water–membrane interphases: Significance for peptide/membrane interactions. *Advances in Colloid and*
503 *Interface Science* **2014**, *211*, 17–33. doi:10.1016/j.cis.2014.05.002.
- 504 48. Disalvo, E.; Lairion, F.; Martini, F.; Tymczyszyn, E.; Frías, M.; Almaleck, H.; Gordillo, G. Structural and
505 functional properties of hydration and confined water in membrane interfaces. *Biochimica et Biophysica*
506 *Acta (BBA) - Biomembranes* **2008**, *1778*, 2655–2670. doi:10.1016/j.bbamem.2008.08.025.
- 507 49. Damodaran, S. Water activity at interfaces and its role in regulation of interfacial enzymes: a hypothesis.
508 *Colloids and Surfaces B: Biointerfaces* **1998**, *11*, 231–237. doi:10.1016/s0927-7765(98)00040-x.
- 509 50. Rao, C.S.; Damodaran, S. Surface pressure dependence of phospholipase A2 activity in lipid
510 monolayers is linked to interfacial water activity. *Colloids and Surfaces B: Biointerfaces* **2004**, *34*, 197–204.
511 doi:10.1016/j.colsurfb.2004.01.004.
- 512 51. Rao, C.S.; Damodaran, S. Is Interfacial Activation of Lipases in Lipid Monolayers Related to
513 Thermodynamic Activity of Interfacial Water? *Langmuir* **2002**, *18*, 6294–6306. doi:10.1021/la025655n.
- 514 52. Schiraldi, A.; Fessas, D.; Signorelli, M. Water Activity in Biological Systems - A Review. *Polish Journal of*
515 *Food and Nutrition Sciences* **2012**, *62*, 5–13. doi:10.2478/v10222-011-0033-5.
- 516 53. Benkovic, S.J.; Hammes-Schiffer, S. A Perspective on Enzyme Catalysis. *Science* **2003**, *301*, 1196–1202.
517 doi:10.1126/science.1085515.
- 518 54. Boekhoven, J.; Hendriksen, W.E.; Koper, G.J.M.; Eelkema, R.; Esch, J.H.v. Transient assembly of active
519 materials fueled by a chemical reaction. *Science* **2015**, *349*, 1075–1079. doi:10.1126/science.aac6103.
- 520 55. Hanson, R.W. The role of ATP in metabolism. *BIOCHEMICAL EDUCATION* **1989**, *2*.
- 521 56. Nam, I.; Lee, J.K.; Nam, H.G.; Zare, R.N. Abiotic production of sugar phosphates and uridine
522 ribonucleoside in aqueous microdroplets. *Proceedings of the National Academy of Sciences* **2017**,
523 *114*, 12396–12400. doi:10.1073/pnas.1714896114.
- 524 57. Pasek, M.A. Thermodynamics of Prebiotic Phosphorylation. *Chemical Reviews* **2019**.
525 doi:10.1021/acs.chemrev.9b00492.
- 526 58. Yousef, A.; Coy, B.M. Diffusion of surfactants to an interface: Effect of a barrier of oriented water. *Journal*
527 *of Colloid and Interface Science* **1983**, *94*, 497–501. doi:10.1016/0021-9797(83)90289-8.
- 528 59. Cornell, C.E.; Black, R.A.; Xue, M.; Litz, H.E.; Ramsay, A.; Gordon, M.; Mileant, A.; Cohen, Z.R.;
529 Williams, J.A.; Lee, K.K.; Drobny, G.P.; Keller, S.L. Prebiotic amino acids bind to and stabilize
530 prebiotic fatty acid membranes. *Proceedings of the National Academy of Sciences* **2019**, *116*, 17239–17244.
531 doi:10.1073/pnas.1900275116.
- 532 60. Xue, M.; Black, R.A.; Cohen, Z.R.; Roehrich, A.; Drobny, G.P.; Keller, S.L. Binding of Dipeptides to Fatty Acid
533 Membranes Explains Their Colocalization in Protocells but Does Not Select for Them Relative to Unjoined
534 Amino Acids. *The Journal of Physical Chemistry B* **2021**, *125*, 7933–7939. doi:10.1021/acs.jpbc.1c01485.
- 535 61. Powner, M.W.; Gerland, B.; Sutherland, J.D. Synthesis of activated pyrimidine ribonucleotides in
536 prebiotically plausible conditions. *Nature* **2009**, *459*, 239–242. doi:10.1038/nature08013.
- 537 62. Deamer, D. Electrochemical energy for living systems. *Current Opinion in Electrochemistry* **2021**, *29*, 100742.
538 doi:10.1016/j.coelec.2021.100742.
- 539 63. Summers, D.P.; Rodoni, D. Vesicle Encapsulation of a Nonbiological Photochemical System Capable of
540 Reducing NAD⁺ to NADH. *Langmuir* **2015**, *31*, 10633–10637. doi:10.1021/la502003j.
- 541 64. Deamer, D. POLYCYCLIC AROMATIC HYDROCARBONS: PRIMITIVE PIGMENT SYSTEMS IN THE
542 PREBIOTIC ENVIRONMENT. *Adv. Space Res.* **1992**, *12*, 183–189.
- 543 65. Robertson, R.M.; Laib, S.; Smith, D.E. Diffusion of isolated DNA molecules: Dependence on length and
544 topology. *Proceedings of the National Academy of Sciences* **2006**, *103*, 7310–7314. doi:10.1073/pnas.0601903103.