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# Bioverse: Exoplanet Biosignature Patterns Under the UV Threshold Hypothesis for the Origin of Life

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#### ABSTRACT

The origins of life on Earth may provide testable hypotheses about biosignature distributions in exoplanet populations. Motivated by the cyanosulfidic origins-of-life scenario, we investigate the hypothesis that a minimum past near-ultraviolet (NUV) flux is necessary for abiogenesis. Using a semi-analytical Bayesian framework und the Bioverse survey simulator, we estimate the probability of obtaining strong evidence for or against this UV Threshold Hypothesis with potential future biosignature surveys. Our results indicate that a correlation between past NUV flux and current biosignature occurrence is observationally testable, but requires sample sizes of at least 50 planets for meaningful constraints. These constraints are critically sensitive to the intrinsic abiogenesis rate, as well as host star properties, particularly maximum past NUV fluxes. Surveys targeting extreme NUV flux values and planets orbiting M dwarfs enhance the likelihood of obtaining conclusive results, with sample sizes  $\gtrsim 100$  providing  $\gtrsim 80\%$  likelihood of strong evidence if abiogenesis rates are high and required NUV fluxes are moderate. Conversely, the fraction of inhabited planets sharply decreases for required fluxes exceeding a few hundred erg/s/cm<sup>2</sup>.

Our findings demonstrate the potential of upcoming exoplanet surveys to test origins-of-life hypotheses. Beyond specific scenarios, our work underscores the broader value of realistic survey simulations in identifying testable science questions, optimizing mission strategies, and advancing theoretical and experimental studies of abiogenesis. This approach emphasizes the need to contextualize biosignature detections, enhancing their credibility and offering new approaches to explore the emergence of life on Earth and beyond.

# 1. INTRODUCTION

A wide variety of scenarios for the origin of life have 25 been proposed (e.g., Baross & Hoffman 1985; Brasier et al. 2011; Mulkidjanian et al. 2012; Fox & Strasdeit 27 2013; Deamer & Georgiou 2015; Westall et al. 2018). 28 While we may still be far away from conclusively test-29 ing them, new prospects in the search for conditions 30 favorable to life have opened up by thinking of the ori-31 gin of life as a planetary phenomenon and identifying 32 global-scale environmental properties that might sup-33 port pathways to life (Sasselov et al. 2020). In partic-34 ular, specific planetary conditions are needed to create 35 stockpiles of initial compounds for prebiotic chemistry; 36 and planetary processes are required to trigger the pre-37 biotic synthesis. Such planetary conditions can be hy-38 pothesized for exoplanets located in the habitable zone 39 (HZ) of their host star, with persistent liquid water on 40 their surface. For example, the deep-sea hydrothermal 41 (or hydrothermal-sedimentary) context for the origin of 42 life requires a direct contact of an ocean and the plan-43 etary mantle/crust (e.g., Baross & Hoffman 1985). The 44 alternate scenario of a surface locally subject to wet-dry

45 cycles requires a planetary exposure to mid-range UV 46 irradiation, as a source of energy and an agent of se-47 lection in chemical evolution (e.g., Deamer et al. 2019). 48 This "UV Threshold Hypothesis" states that UV light in as a specific wavelength range played a constructive role in 50 getting life started on Earth (Ranjan & Sasselov 2016; 51 Ranjan et al. 2017a; Rimmer et al. 2018; Rapf & Vaida 52 2016), and it could provide a probabilistic approach to 53 the interpretation of possible future biosignature detec-54 tions (e.g., Catling et al. 2018a; Walker et al. 2018). The association of chemical pathways to life and plan-56 etary environmental conditions offers a new opportunity 57 to test alternate scenarios for life emergence based on 58 planetary-level data collected from the upcoming ob-59 servations of populations of exoplanets. Deep-sea hy-60 drothermal scenarios require planetary conditions that 61 may not be met on ocean worlds with large amounts 62 of water, where the water pressure on the ocean floor 63 is high enough to form high-pressure ices (Noack et al. 64 2016; Kite & Ford 2018). In this case, a testable predic-65 tion would be that planets with high-pressure ices do not 66 show biosignatures. Likewise, if UV light is required to 67 get life started, then there is a minimum planetary UV

flux requirement to have an inhabited world. This requirement is set by competitor thermal processes; if the photoreaction does not move forward at a rate faster than the competitor thermal process(es), then the abiogenesis scenario cannot function. On the other hand, abundant UV light vastly in excess of this threshold does not increase the probability of abiogenesis, since once the UV photochemistry is no longer limiting, some other thermal process in the reaction network will be the rate-limiting process instead. Therefore, a putative dependence of life on UV light is best encoded as a step function (e.g., Ranjan et al. 2017a; Rimmer et al. 2018, 2021a).

The goal of this work is to evaluate the potential of potential future exoplanet surveys to test the hypothesis that a minimum past NUV flux is required for abiogenesis. We focus on one version of the UV Threshold Hypothesis, the so-called cyanosulfidic scenario, which has been refined to the point where the required threshold flux has been measured to be  $F_{\rm NUV,min}=(6.8\pm3.6)\times10^{10}\,{\rm photons\,cm^{-2}\,s^{-1}\,nm^{-1}}$  integrated from 200–280 nm at the surface (Rimmer of et al. 2018, 2021b; Rimmer 2023; Ranjan et al. 2023).

We first follow a semi-analytical Bayesian analysis to 92 estimate probabilities of obtaining strong evidence for 93 or against this hypothesis. Under the UV Threshold 94 Hypothesis  $(H_1)$ , the probability of an exoplanet having 95 detectable biosignatures is zero if the near-ultraviolet 96 (near-Ultraviolet (NUV)) irradiation is less than the 97 threshold, and it is equal to the (unknown) probabil-98 ity of abiogenesis,  $f_{\text{life}}$  if NUV exceeds the threshold for a sufficiently long period of time. Under the null hypothesis  $(H_0)$ , that probability simply is  $f_{life}$ , that is, it does not correlate with the UV flux. Figure 1 shows these hypotheses as derived from the predictions of the 104 cyanosulfidic scenario. Given a sample of planets, where 105 for some of them we have convincing biosignature detec-106 tions but remain agnostic on  $f_{\rm life}$ , we ask what evidence for  $H_1$  and  $H_0$  we can expect to obtain.

A real exoplanet survey will be subject to observational biases, sample selection effects, and the underlying demographics of the planet sample. To assess the information gain of a realistic exoplanet survey, we employed Bioverse (Bixel & Apai 2021; Hardegree-Ullman et al. 2023; Schlecker et al. 2024; Hardegree-Ullman et al. 2024), a framework that integrates multiple components including statistically realistic simulations of exoplanet populations, a survey simulation module, and a hypothesis testing module to evaluate the statistical power of different observational strategies.

This paper is organized as follows: In Section 2, we introduce both our semi-analytical approach and Bioverse

121 simulations for testing the UV Threshold Hypothesis. 122 Section 3 presents the results of these experiments for 123 a generic survey as well as for a realistic transit survey. 124 In Section 4, we discuss our findings before concluding 125 with a summary in Section 5.

#### 2. METHODS

# 2.1. Fraction of inhabited planets with detectable biosignatures

Here, we conduct a theoretical experiment on the 130 UV Threshold Hypothesis by relating the occurrence 131 of life on an exo-earth candidate with a minimum past 132 quiescent stellar UV flux, focusing on the prebiotically 133 interesting NUV range from 200–280 nm.

SUKRIT: got a good reference for this?

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Our concrete hypothesis shall be that life only occurs on planets that at some point in their history have received such radiation at a flux exceeding a threshold  $_{138}$   $F_{\rm NUV,min}$ .

## 2.2. Semi-analytical approach

We first assessed the expected probabilities of obtaining true negative or true positive evidence for the UV Threshold Hypothesis  $(H_1)$  above, as well as the probability for misleading or inconclusive evidence, under idealized conditions. This serves as a first-order estimate of the information content of a survey, before we take into account impacts from exoplanet demographics, sample selection, and survey strategy.

Presumably, not all habitable worlds are inhabited and not all inhabited worlds develop detectable biosignatures. The fraction of exo-Earth candidates (EEC) that are both inhabited and harbor detectable biosignatures at the time when we observe them remains speculative; we aggregate them in the unitless parameter  $f_{\rm life}$ . Let us consider the probability to detect a biosignature  $F_{\rm log}$  and let our observable be the inferred past NUV find flux of the planet  $F_{\rm NUV}$ . Under Hypothesis  $H_1$  (Equation 8), there exists a special unknown value of  $F_{\rm NUV}$ , noted  $F_{\rm NUV,min}$  such that

$$P(L|F_{\text{NUV}}, H_1) = f_{\text{life}} \quad \text{if } F_{\text{NUV}} > F_{\text{NUV,min}}$$
 (1)

$$P(L|F_{\text{NUV}}, H_1) = 0$$
 otherwise (2)

where  $f_{\rm life}$  is the unknown probability of abiogenesis. The corresponding null hypothesis is that there exists no such special value of  $F_{
m NUV}$  and that

$$P(L|F_{\text{NUV}}, H_{\text{null}}) = f_{\text{life}}.$$
 (3)

If we now define a sample of size n as  $X = \{F_{\text{NUV},i}, L_i\}_{i \in [1,n]}$  where  $L_i$  is equal to 1 if life is detected and 0 otherwise, we can calculate the evidence

Hypothesis: Life only originates on planets with particular UV irradiance

### Prediction

H1 Correlation between past UV flux and biosignature occurrence

Past UV flux and the occurrence of biosignatures are correlated

H0 No correlation between past UV flux and biosignature occurrence

Figure 1. UV Threshold Hypothesis and null hypothesis derived from the cyanosulfidic scenario.

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for hypothesis  $H_i$  against  $H_j$  through the Bayes factor

$$BF_{H_i,H_j} = \frac{P(X|H_i)}{P(X|H_j)},\tag{4}$$

with  $P(X|H_i)$  and  $P(X|H_i)$  likelihoods of obtaining the sample X under either hypothesis.

If we define  $k = \sum L_i$  and denote Y the random vari- $_{174}$  able that describes it,  $H_{\rm null}$  represents the likelihood that the number of planets with life in the sample follows 176 the binomial distribution

$$P(Y = k|H_{\text{null}}) = \binom{n}{k} f_{\text{life}}^k (1 - f_{\text{life}})^{n-k}.$$
 (5)

Under  $H_1$ , Y also follows a binomial distribution, however it is conditioned by  $n_{\lambda} = Card(\{F_{\text{NUV,i}} \text{ if } F_{\text{NUV,i}} >$ 180  $F_{\text{NUV,min}}\}_{i\in[1,n]}$ ) the number of values of  $F_{\text{NUV}}$  in the 181 experiment that exceed  $F_{\rm NUV,min}$ 

$$P(Y = k|H_1) = \binom{n_{\lambda}}{k} f_{\text{life}}^k (1 - f_{\text{life}})^{n_{\lambda} - k}.$$
 (6)

Hence, 183

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$$BF_{H_1,H_{\text{null}}} = \frac{P(Y=k|H_1)}{P(Y=k|H_{\text{null}})} = \frac{\binom{n_{\lambda}}{k}}{\binom{n}{k}} (1-f_{\text{life}})^{n_{\lambda}-n}, \tag{7a}$$
 184 and

$$BF_{H_{\text{null}},H_1} = \frac{P(Y=k|H_{\text{null}})}{P(Y=k|H_1)} = \frac{\binom{n}{k}}{\binom{n_{\lambda}}{k}} (1-f_{\text{life}})^{n-n_{\lambda}}.$$
 (7b)

Given a sample of planets, where for some of them we 188 have convincing biosignature detections but remaining agnostic on  $f_{life}$ : What evidence for  $H_1$  and  $H_{null}$  can we 190 expect to get? Our Bayes factor (Equation 7a) is deter-191 mined by the unknown variables  $f_{\text{life}}$  and  $F_{\text{NUV,min}}$ , as well as the number of planets with biosignature detections in the sample k. To compute the distribution of evidences, we repeatedly generated samples under  $H_1$  and 195  $H_{\text{null}}$  and computed the Bayes factors  $BF_{H_1,H_{\text{null}}}$  and 196  $BF_{H_{\text{null}},H_1}$ . We then evaluated the fraction of Monte 197 Carlo runs in which certain evidence thresholds (Jeffreys 198 1939) were exceeded.

2.3. Exoplanet survey simulations with Bioverse

To assess the diagnostic power of realistic exoplanet 201 surveys, we employed our survey simulator and hypoth-202 esis testing framework Bioverse (Bixel & Apai 2021). 203 The general approach is as follows:

- 1. Exoplanet population synthesis: We populate the Gaia Catalogue of Nearby Stars (Smart et al. 2021) with synthetic exoplanets whose orbital parameters and planetary properties reflect our current understanding of exoplanet demographics (Bergsten et al. 2022; Hardegree-Ullman et al. 2023). Here, we also inject the demographic trend in question - in this case we assign biosignatures according to  $H_1$ , i.e., to planets in the HZ that have received NUV fluxes above a certain threshold.
- 2. Survey simulation: We simulate the detection and characterization of these exoplanets with a hypothetical survey, taking into account the survey's sensitivity, target selection, and observational biases. To model the sensitivity of the information gain of a proposed mission to sample selection and survey strategy, we conduct survey simulations with Bioverse using different sample sizes and survey strategies.
- 3. Hypothesis testing: We evaluate the likelihood that a given survey would detect a specified demographic trend in the exoplanet population and estimate the precision with which the survey could constrain the parameters of that trend. A common definition of the null hypothesis  $H_0$ , which is also applied here, is that there is no relationship between the independent variable (here: maximum NUV flux) and the dependent variable (here: biosignature occurrence). The alternative hypothesis  $H_1$  proposes a specific relationship between the independent and dependent variables. Bioverse offers either Bayesian model comparison or non-parametric tests to evaluate the evidence for or against the null hypothesis.

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To determine the diagnostic capability of a given survey, Bioverse runs multiple iterations of the simulated
survey and calculates the fraction of realizations that
successfully reject the null hypothesis. We use this metric, known as the statistical power, to quantify the potential information content of the survey, identify critical design trades, and find strategies that maximize the
survey's scientific return.

### 2.3.1. Simulated star and planet sample

We generated two sets of synthetic exoplanet popula-249 tions, one for FGK-type stars and one for M-type stars. 250 The stellar samples are drawn from the Gaia Catalogue of Nearby Stars (Smart et al. 2021) with a maximum 252 Gaia magnitude of 16 and a maximum stellar mass of 253 1.5  $M_{\odot}$ . We included stars out to a maximum distance 254  $d_{\text{max}}$  that depends on the required planet sample size. <sup>255</sup> Planets were generated and assigned to the synthetic 256 stars following the occurrence rates and size/orbit dis-257 tributions of Bergsten et al. (2022). Following Bixel & <sup>258</sup> Apai (2021), we considered only transiting EECs with radii  $0.8 S^{0.25} < R < 1.4$  that are within the HZ (see 260 Section 2.3.2). The lower limit was suggested as a min-261 imum planet size to retain an atmosphere (Zahnle & <sup>262</sup> Catling 2017). To generate planet samples larger than what the stellar catalog in combination with these oc-264 currence rates yields, we scaled up the occurrence rates 265 by a constant factor that yields the desired number of 266 planets. This was in particular necessary for the FGK 267 sample, where the rate of transiting planets that occupy the HZ is low.

For all survey simulations and hypothesis tests, we repeated the above in a Monte Carlo fashion to generate randomized ensembles of synthetic star and planet populations (Bixel & Apai 2021).

# 2.3.2. Habitable zone occupancy and UV flux

To test the UV Threshold Hypothesis, we require that life occurs only on planets with sufficient past UV irration exceeding the origins of life threshold  $F_{\rm NUV,min}$ . Further, we require this flux to have lasted for a minimum duration  $\Delta T_{\rm min}$  to allow for a sufficient "origins timescale" (Rimmer 2023). All common Origin of Life scenarios require water as a solvent; we thus consider only rocky planets that may sustain liquid water on their surface, i.e., that occupy their host star's momentary HZ during the above period, as well as at the epoch of observation. To determine HZ occupancy, we take into account the evolution of the host star's luminosity and HZ boundaries.

The HZ describes a region around a star where a planet with Earth's atmospheric composition and climate feedbacks can maintain liquid water on its sur-

290 face (e.g., Ramirez & Kaltenegger 2017; Ramirez 2018; 291 Mol Lous et al. 2022; Spinelli et al. 2023; Tuchow & 292 Wright 2023). Here, we adopt estimates that define 293 the HZ as the region between the runaway greenhouse 294 transition, where the stellar instellation cannot anymore 295 be balanced through infrared cooling to space (Inger-296 soll 1969), and the maximum greenhouse limit, corre-297 sponding to the maximum distance at which surface 298 temperatures allowing liquid water can be maintained 299 through a  $CO_2$  greenhouse effect (Kasting 1991; Kast-300 ing et al. 1993; Underwood et al. 2003; Kopparapu et al. 301 2013, 2014). We use the parametrization in Kopparapu 302 et al. (2014) to derive luminosity and planetary mass-303 dependent edges of the HZ  $a_{\rm inner}$  and  $a_{\rm outer}$ .

To determine HZ occupancy, we interpolated the stel305 lar luminosity evolution grid of Baraffe et al. (1998) us306 ing a Clough Tocher interpolant (Nielson 1983; Alfeld
307 1984, see left panel of Figure 2) to compute the evolu308 tion of the inner (runaway greenhouse) and outer (max309 imum greenhouse) edges as a function of planet mass
310 and stellar spectral type (Kopparapu et al. 2014). Be311 ing a local interpolation method, Clough Tocher enables
312 rapid processing while producing a smooth interpolating
313 surface that highlights local trends. From this, we get
314 each planet's epochs within and outside the HZ.

For the NUV flux, we use the age- and stellar mass- dependent NUV fluxes in the HZ obtained by Richey-Yowell et al. (2023), which considers GALEX UV data in the wavelength range of 177–283 nm. We linearly interpolate in their measured grid, where we convert spectral type to stellar mass using the midpoints of their mass ranges (0.75  $\rm M_{\odot}$  for K stars, 0.475  $\rm M_{\odot}$  for early-type M stars, and 0.215  $\rm M_{\odot}$  for late-type M stars). Outside the age and stellar mass range covered in Richey-Yowell et al. (2023), we extrapolate using nearest simplex (see right panel of Figure 2).

We then determined which planets were both in the HZ and had NUV fluxes above  $F_{\rm NUV,min}$ . To avoid considering short transitional phases, we require this situation to last for a minimum duration  $\Delta T_{\rm min} \geq 1\,{\rm Myr}$ . We assigned the development of life to a random fraction  $f_{\rm life}$  of all temperate planets fulfilling these requirements. For the probability of a planet having detectable biosignatures,  $P({\rm bio})$ , the UV Threshold Hypothesis then states

$$H_1: P(\text{bio}) = \begin{cases} 0, & F_{\text{NUV}} < F_{\text{NUV}, \text{min}} \\ f_{\text{life}}, & F_{\text{NUV}} \ge F_{\text{NUV}, \text{min}} \text{ and in HZ for } \Delta t \ge 1 \text{ Myr} \end{cases}$$
(8)

336 and the corresponding null hypothesis  $H_{\text{null}}: P(\text{bio}) =$  337  $f_{\text{life}}$ , i.e., no correlation with UV flux.

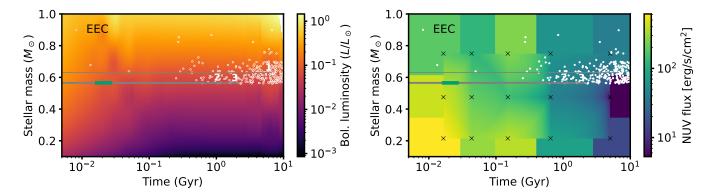


Figure 2. Interpolated stellar luminosity evolution (left) and evolution of the NUV flux in the HZ (right) as a function of host star mass. The scatter plots show age and host star mass of the transiting planets in the synthetic FGK sample; crosses denote the estimated NUV values in Richey-Yowell et al. (2019). A few example tracks for an example threshold flux of  $F_{\text{NUV,min}} = 300.0\,\text{erg}\,\text{s}^{-1}\,\text{cm}^{-2}$  are shown; extended overlap of HZ occupancy and high NUV flux (green sections) fulfills our requirement for abiogenesis. Planet 1 is an EEC that never receives sufficient NUV flux for abiogenesis. Planet 2 and Planet 3 enter the HZ at different times and receive sufficient NUV flux for different durations until their respective host star evolves below the threshold.

# 2.3.3. Transit survey simulations

With the synthetic star and planet samples generated,
we used Bioverse's survey module to simulate noisy measurements of key observables with a transit survey. We
saza assumed a hypothetical mission that can target a large
planet sample with high photometric precision and conduct a biosignature search on these planets (e.g., Apai
et al. 2019, 2022). The simulated survey was designed to
measure planetary instellation (for HZ occupancy) with
measure planetary instellation (for HZ occupancy) with
apprecision of 5% and host star effective temperature
with a precision of 50.0 K. The maximum past NUV
flux a planet received can be determined within a precision of 5%. To marginalize over choices of biosignatures
and their detectability, which are beyond the scope of
this study, we assumed that any inhabited planet would
show a biosignature detectable by the survey.

# 2.3.4. Hypothesis testing

To evaluate the evidence for correctly rejecting the null hypothesis, we employed the Mann-Whitney U test (Mann & Whitney 1947). This is a non-parametric test of the null hypothesis that two independent samples were drawn from a population with the same distribution and is in particular sensitive to one sample being stochastically greater than the other. We used the Mann-Whitney U test to compare the distributions of NUV fluxes of planets with and without biosignatures. The implementation in Bioverse relies on the scipy.stats.mannwhitneyu function (Virtanen et al. 2020) and returns a p-value, for which we set a significance level of  $\alpha = 0.05$  to reject the null hypothesis. For every hypothesis test, we repeated randomized survey realizations to estimate the fraction of successful rejections.

370 tions of the null hypothesis, i.e., the statistical power of 371 the survey.

### 3. RESULTS

## 3.1. Semi-analytical assessment

In Section 2.2, we computed the probability for true positive evidence for  $H_1$  and  $H_{\rm null}$ , respectively (Equations 7a, 7b). Figure 3 shows how these evidences are distributed for sample sizes 10 and 100, and how likely we are to obtain strong evidence ( $BF_{H_i,H_j} > 10$ ). For n=10, strong true evidence for  $H_1$  ( $H_{\rm null}$ ) can be expected in  $\sim 30\,\%$  ( $\sim 40\,\%$ ) of all random experiments. In the majority of cases, the outcome of the survey will be inconclusive. The situation improves with larger samples: for n=100, 80 % of random samples permit conclusive inference (strong true evidence) under either  $H_1$  or  $H_{\rm null}$ .

The expected resulting evidence further depends on the a priori unknown abiogenesis rate  $f_{\rm life}$  and on the NUV flux threshold. Figure 4 illustrates this dependency: For very low values of either parameter, samples drawn under the null or alternative hypotheses are indistinguishable and the Bayesian evidence is always low. Both higher  $f_{\rm life}$  and higher NUV flux thresholds increase the probability of obtaining strong evidence. Larger sample sizes enable this at lower values of these parameters.

So far, we have assumed random, uniform distribusions of  $f_{\rm life}$ ,  $F_{\rm NUV,min}$ , and  $F_{\rm NUV}$ . A high biosignature detection rate  $f_{\rm life}$  increases the evidence (cmp. Equation 7a) but we cannot influence it. The same true for  $F_{\rm NUV,min}$ , where again higher values increase the evidence as the binomial distribution for  $H_1$  gets increasingly skewed and shifted away from the one

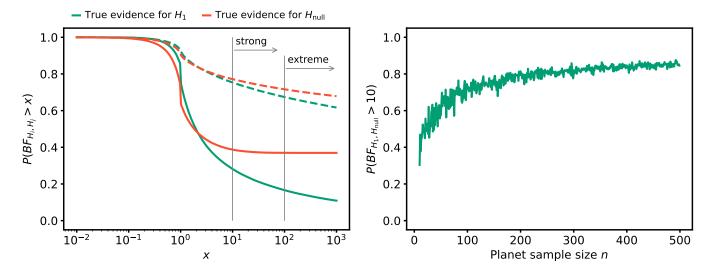
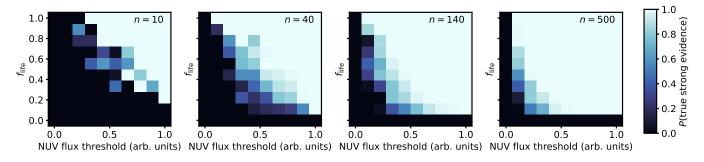


Figure 3. Obtaining true strong evidence with different sample sizes. Left: Probability to reach given evidence levels for  $H_1$  and  $H_{\text{null}}$  under sample sizes n = 10 (solid) and n = 100 (dashed). Vertical lines denote thresholds for "strong" evidence,  $BF_{H_i,H_j} > 10$ , and "extreme" evidence,  $BF_{H_i,H_j} > 100$ . Right: Probability of obtaining true strong evidence for  $H_1$  as a function of sample size n.



**Figure 4.** Probability of obtaining true strong evidence for different abiogenesis rates, NUV flux thresholds, and sample sizes. For each of these parameters, higher values increase the probability of yielding strong evidence.

405 for  $H_{\text{null}}$ . However, one might 'cherry-pick' exoplanets 406 for which a biosignature test is performed based on apriori available contextual information (Catling et al. 2018b) in order to maximize the science yield of investing additional resources. For instance, the distri-410 bution of  $F_{\rm NUV}$  in the planet sample can be influenced 411 by the survey strategy, and a targeted sampling ap-412 proach could favor extreme values. We model this by 413 distributing  $F_{\text{NUV}}$  according to different Beta functions and introduce a selectivity parameter  $s \in ]-1,1[$  such 415 that  $F_{\rm NUV} \sim Beta(1/10^s, 1/10^s)$ . Figure 5 shows how 416 the probability of obtaining true strong evidence for  $H_1$ 417 scales with selectivity s. For large samples, a high selec-418 tivity  $(s \sim 1)$  can increase the probability of obtaining 419 true strong evidence from  $\sim 70\%$  for s=0 (random 420 uniform distribution) to > 90%.

With HZ occupancy as a requirement for abiogene-423 sis, and barring selection biases beyond stellar brightness, the host star distribution of inhabited planets in a 425 simulated transit survey is skewed toward later spectral 426 types. For a fixed planet sample size, the fraction of in-427 habited planets is highest for planets orbiting M dwarfs 428 due to the higher NUV fluxes in the HZ of these stars (see Figures 2, 6). Their NUV fluxes are generally high-430 est at early times  $\lesssim 100 \, \mathrm{Myr}$ . These host stars, in par-431 ticular late subtypes, also provide extended periods of 432 increased NUV emission that overlap with times when 433 some of these planets occupy the HZ (see Figure 2), our 434 requirement for abiogenesis (compare Equation 8). Thus - under the UV Threshold Hypothesis - most inhabited transiting planets in the sample orbit M dwarfs. Here, we are interested in the statistical power of a

Here, we are interested in the statistical power of a transit survey with a plausible sample selection and size.
In the following, we fix the sample size to 250 and consider two different survey strategies targeting FGK and

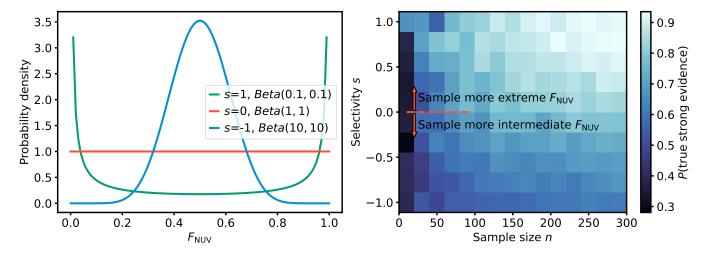


Figure 5. Scaling of the probability of obtaining true strong evidence with sample selectivity. Left: Sampling distribution for different selectivity parameters s. Right: Resulting P(true strong evidence), where  $f_{\text{life}}$  and  $F_{\text{NUV,min}}$  are randomly drawn from a uniform distribution. Sampling more extreme values of  $F_{\text{NUV}}$  is more likely to yield strong evidence.

<sup>441</sup> M dwarfs, respectively. We further investigate the sen-<sup>442</sup> sitivity of the survey to the a priori unknown threshold <sup>443</sup> NUV flux  $F_{\text{NUV,min}}$  and the abiogenesis rate  $f_{\text{life}}$ .

# 3.2.1. Selectivity of simulated transit surveys

In Section 3.1, we demonstrated that the probabilate ity of obtaining true strong evidence for the hypothesis that life only originates on planets with a minimum past NUV flux is sensitive to the distribution of sampled past NUV fluxes, i.e., the selectivity of the survey (compare Figure 5). For both surveys targeting M dwarfs and those targeting FGK dwarfs, the maximum NUV distribution is rather unimodal. Applying the approach from Sect. 3.1 of fitting a Beta function to the distribution, we find rather low selectivities (see Figure 6), which is likely a detriment for statistical hypothesis tests.

# 3.2.2. Expected biosignature pattern

A representative recovery of the injected biosignature 457 pattern is shown in Figure 6. There, we assumed an abiogenesis rate of  $f_{\rm life}=0.8$  and a minimum NUV flux of  $F_{\text{NUV,min}} = 300.0 \,\text{erg}\,\text{s}^{-1}\,\text{cm}^{-2}$ . All injected biosignatures are assumed to be detected without false positive ambiguity, and the maximum NUV flux is estimated 463 from the host star's spectral type and age with an uncer-464 tainty corresponding to the intrinsic scatter in the NUV 465 fluxes in Richey-Yowell et al. (2023). This leads to a 466 distribution of biosignature detections with detections 467 increasingly occurring above a threshold inferred NUV 468 flux. In this example case, the few biosignature detections in the FGK sample lead to a higher evidence than 470 in the M dwarf sample, where the majority of planets 471 are above the threshold NUV flux.

Figure 7 shows the fraction of inhabited planets under the UV Threshold Hypothesis for different threshold 474 NUV fluxes and for a high abiogenesis rate of  $f_{\rm life}=1$ .
475 This fraction decreases sharply with increasing thresh476 old flux, as fewer planets receive sufficient NUV flux for
477 abiogenesis. Another effect responsible for this drop is
478 that some planets receive the required NUV flux only
479 before entering the HZ – this is especially likely for
480 M dwarfs. For the FGK sample, the fraction of inhab481 ited planets drops at lower threshold fluxes than for the
482 M dwarf sample.

# 3.2.3. Statistical power and sensitivity on astrophysical parameters

We now investigate the sensitivity of the achieved statistical power of our default transit survey to the a priori unconstrained threshold NUV flux  $F_{\rm NUV,min}$  and the abiogenesis rate  $f_{\rm life}$ . Figure 8 shows the statistical power as a function of these parameters for a sample size of N=250. Values of  $F_{\rm NUV,min}$  that lie between the exterma of the inferred maximum NUV flux increase the achieved statistical power of the survey, as in this case the dataset under the alternative hypothesis  $H_1$  differs more from the null hypothesis. Furthermore, a higher abiogenesis rate  $f_{\rm life}$  increases the evidence for  $H_1$ .

Notably, the sensitivity of the M dwarf sample extends into the low NUV flux end due to the broader distribution of maximum past NUV fluxes in this sample. Here, the FGK sample is less sensitive.

Update this with final results from stat. power grid

The only parameter space region with statistical power above 90% lies at abiogenesis rates  $f_{\rm life}>0.9$  and threshold NUV fluxes of  $\sim 200$ –400 erg s<sup>-1</sup> cm<sup>-2</sup>.

# 4. DISCUSSION

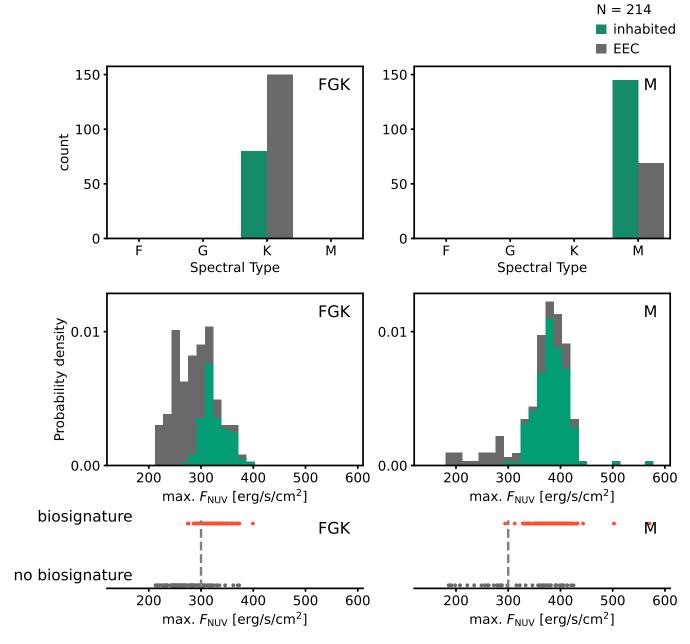


Figure 6. Simulated transit surveys targeting FGK and M stars. Top: Host stars of all transiting EECs and inhabited planets in a simulated transit survey. In the FGK sample, most EECs and all inhabited planets orbit K dwarfs. In an M dwarf sample of the same size, the fraction of inhabited planets is larger. Center: Distribution of inferred maximum past NUV flux in transit surveys targeting EECs around FGK and M stars, respectively. The best-fit beta distributions (gray) correspond to selectivities of  $s_{\rm FGK} = -0.47$  and  $s_{\rm M} = -0.02$ . Green areas show inhabited planets for an abiogenesis rate of  $f_{\rm life} = 0.8$  and a generic threshold NUV flux  $F_{\rm NUV,min} = 300.0\,{\rm erg\,s^{-1}\,cm^{-2}}$ . Bottom: Recovered biosignature detections and non-detections of simulated transit surveys. The dashed line denotes  $F_{\rm NUV,min}$ .

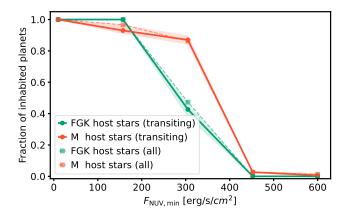


Figure 7. Fraction of inhabited planets for different threshold NUV fluxes under the UV Threshold Hypothesis if the abiogenesis rate is 1. Shaded regions denote 90 percent confidence intervals of randomized sample generations, and dashed lines correspond to samples including non-transiting planets. For all samples, the fraction of inhabited planets drops sharply with increasing threshold NUV flux due to the combined effects of never receiving sufficient NUV flux for abiogenesis or receiving it before entering the HZ.

A key question in the quest to understand the origins of life is which natural processes best explain how living matter spontaneously appears from nonliving matter (e.g., Malaterre et al. 2022). Using astronomical methods, this question will likely not be testable for individual planets but only by the presence of patterns in planetary populations. The cyanosulfidic scenario (Patel et al. 2015), in particular its predicted existence of a minimum NUV flux required for prebiotic chemistry, seems to offer an opportunity to test an origins of life hypothesis with a statistical transit survey sampling planets with varying NUV flux histories. In the following, we discuss the prospects of testing the UV Threshold Hypothesis in light of our results.

# 519 4.1. Sampling strategy for testing a NUV flux threshold

In Sect. 3.1, we show that testing the UV Thresh1010 dollar Hypothesis suffers from 'nuisance' parameters that
1021 render inference through astronomical observations diffi1022 cult. Here, these parameters are the unspecified value of
1024 the NUV threshold hypothesized to exist under  $H_1$ , and
1025 the unknown probability of detectable life emerging on
1026 a habitable planet  $f_{\text{life}}$ . While the inference of a planet's
1027 entire UV flux evolution is difficult (e.g., Richey-Yowell
1028 et al. 2023), the estimated maximum NUV flux that a
1029 planet was exposed to may be used as a proxy, at least if
1030 one is interested in a minimum threshold flux and makes
1031 the assumption that planetary surfaces offer protection
1032 against too high UV flux. Indeed, the distribution of the
1033 number of planets with determined maximum NUV

values  $F_{\text{NUV}}$  depends on both the values of  $F_{\text{NUV},min}$  and  $f_{\text{life}}$  as shown in equation 6.

In our semi-analytical analysis (Section 3.1), we 538 project a possible test performed by a future observer equipped with a sample of exoplanets with derived past maximum NUV exposure for which biosignature detec-541 tion has been attempted. This is necessarily reductive 542 as this observer will have more knowledge about exper-543 imental conditions and will therefore be able to use this information to guide hypothesis testing. For instance, 545 we have made the choice to consider the total number of 546 detected biosignatures as our summary statistic (Equa-547 tion 6), which is not 'sufficient' to infer  $F_{NUV,min}$  and 548  $f_{life}$  separately. However, by conditioning the Bayes factor to these variables (Equation 7a), we calculate the 550 probability distribution of the Bayesian evidence in fa-**551** vor of  $H_1$ . In doing so, we may evaluate how evidence 552 depends on the uncertainty over these unknown parameters in general terms, without assuming which particular test a future observer might choose to actually perform over real data when available. From this, we can see that target selection can strongly affect the conclusiveness of 557 a future test of the UV Threshold Hypothesis.

The particular finding that prioritizing extreme values of past NUV flux can enhance statistical power likely 560 clashes with observational constraints, as the composi-561 tion of the subset of planets that we can observe and 562 for which detection of biosignature can be attempted is 563 not independent from their NUV flux history. Hence, for our future observer, selectivity and sample size may be in conflict. This trade-off can be quantified in terms 566 of expected evidence yield, which we have done in Sec-567 tion 3.1. Our analysis shows that regardless of selec-568 tivity, sample sizes smaller than 50 likely result in in-569 conclusive tests, and that increasing selectivity towards 570 extreme  $F_{\text{NUV}}$  offers limited inference gains compared 571 to the uniform case (s = 0; Figure 5). For larger sam-572 ples, however, a narrow distribution of  $F_{\text{NUV}}$  may pre-573 vent inference entirely. We thus argue that selecting a 574 sample with  $F_{\text{NUV}}$  distributed uniformly or emphasiz-575 ing extreme values should – barren any practical coun-576 terarguments – be considered in any future attempt at 577 testing the UV Threshold Hypothesis. Since the prac-578 tical implementation of an exoplanet survey can stand 579 in the way of such a selection, the following discussion 580 focuses on the results of our transit survey simulations 581 with Bioverse.

# 4.2. How planetary context may constrain the UV Threshold Hypothesis

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It comes to no surprise that the success rate of such a test is sensitive to the sample size of the survey and

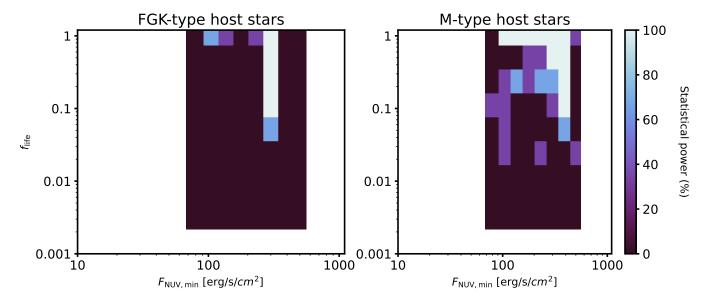


Figure 8. Statistical power as a function of threshold NUV flux and abiogenesis rate. Even for a large sample (here: N = 250), a high statistical power of the transit survey requires high abiogenesis rates  $f_{\text{life}}$ . Intermediate values of  $F_{\text{NUV,min}}$  are more likely to yield strong evidence than extreme values; the sensitivity of the M dwarf sample extends into the low NUV flux end.

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586 to the occurrence of life on temperate exoplanets. As we have shown, the statistical power of this test also 588 depends on the distribution of past NUV fluxes in the sample and on the threshold flux. Optimizing the survey to sample a wide range of NUV flux values, par-591 ticularly at the extremes, can enhance the likelihood of obtaining strong evidence for or against the hypothesis. Intermediate values of the threshold NUV flux are more likely to yield strong evidence than extreme valby ues, as the dataset under the alternative hypothesis  $H_1$ differs more from the null hypothesis in this case while 597 still being sufficiently populated. The threshold flux is, 598 of course, a priori unknown and we cannot influence it. If, however, better theoretical predictions for the re-600 quired NUV flux for abiogenesis become available (Rim-601 mer et al. 2021a), the survey strategy can be further optimized, for instance by targeting planets that are estimated to have received a NUV flux slightly below and above this threshold or by applying a bisection algorithm in a sequential survey (Fields et al. 2023). 605

### 4.3. An M dwarf opportunity

An interesting aspect lies in the distribution of host star properties, as different spectral types probe different ent past NUV flux regimes. FGK stars show a narrow distribution of maximum past NUV fluxes in the HZ, which may limit the diagnostic power of a survey depending on the (unknown) threshold NUV flux. With a pure FGK sample, a survey will be more sensitive to the low NUV flux end of the hypothesis. A lack of biosignatures on these planets would support the UV Threshold Hypothesis. Their presence would have little constrain-

617 ing power; it would either suggest that low NUV fluxes 618 are sufficient for abiogenesis or indicate a different abio-619 genic pathway (e.g., Westall et al. 2018).

On the other hand, M dwarfs show a wider distribu-621 tion of maximum past NUV fluxes in their HZs. While 622 old M dwarfs can be considered low-UV environments, a large fraction of them emit high NUV fluxes into their 624 HZ during their early stages, in particular later sub-625 types (Richey-Yowell et al. 2023). An M dwarf sample 626 will thus help to test the high NUV flux end of the UV 627 Threshold Hypothesis; a higher occurrence of biosigna-628 tures here would support the hypothesis that a higher 629 NUV flux is beneficial or necessary for life. As our sim-630 ulations show, the higher and more variable NUV fluxes 631 in M dwarfs increase the likelihood of obtaining strong 632 evidence for or against the hypothesis. The combination 633 of a lack of UV radiation today, which makes biosigna-634 ture gases more detectable (Segura et al. 2005), and a 635 UV-rich past that may have enabled abiogenesis could 636 make M dwarfs the preferred targets for biosignature 637 searches.

These findings underscore the importance of constraining the UV emission profiles of EEC host stars throughout their evolutionary stages to assess the viability of M-dwarf planets as testbeds for theories on the origins of life (Rimmer et al. 2021a; Ranjan et al. 2023).

### 4.4. Sensitivity to astrophysical parameters

Our Bioverse simulations that take into account exoplanet demographics, the evolution of habitability and NUV fluxes, and observational biases show that not only the likelihood of a conclusive test of the UV Thresh648 old Hypothesis, but also the likelihood of successful biosignature detection itself is extremely sensitive to the 650 threshold NUV flux if the hypothesis is true. Even 651 if all biosignatures can be detected and the nominal 652 abiogenesis rate is very high, say  $f_{\rm life}=1,$  under the 653 condition that prebiotic chemistry requires a minimum 654 NUV flux and liquid water, if the threshold flux turns 655 out to be high the probability of finding life on a ran-656 domly selected planet may be very low. As we showed, 657 for high required fluxes the two requirements of simul-658 taneous HZ occupancy and sufficient NUV flux conspire to diminish the fraction of inhabited planets in 660 the sample. Taking the inferred fluxes from Richey-Yowell et al. (2023) at face value (but taking into ac-662 count intrinsic scatter), a minimum required NUV flux of of  $\gtrsim 400\,\mathrm{erg\,s^{-1}\,cm^{-2}}$  reduces the fraction of inhabited planets to below  $\sim 1\%$ . This not only calls for a large sample size and a targeted sample selection preferring 666 high expected past NUV fluxes, but also highlight the 667 necessity of continued theoretical and experimental research into the role of UV radiation in prebiotic chem-669 istry (Ranjan et al. 2017b; Rimmer et al. 2018, 2021a).

# 4.5. Contextual support for potential biosignature detections

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The predicted interplay of NUV flux and HZ occupancy in enabling abiogenesis via the cyanosulfidic scenario could in principle be used to add or remove credibility from a tentative biosignature detection. For exmaple, with a strong belief that this scenario is the only
viable one for the origins of life, a biosignature detection
on a planet orbiting a strongly UV-radiating star may
add credibility to the detection. Conversely, a biosigmature detection on a planet estimated to have received
very little UV radiation would increase the likelihood of
the detection in the latter case be confirmed, it could be
the detection in the latter case be confirmed, it could be
used to falsify the UV Threshold Hypothesis.

Our simulations find no clear criterion for the credibility of a biosignature detection based on spectral type of
the host star, as both FGK and M dwarf samples show
similar maximum past NUV flux distributions. The
abiogenesis rate in both samples show similar trends
with the threshold NUV flux, and the fraction of inhabited planets drops at similar threshold fluxes (see
Section 3.2). A potentially inhabited planet's host star
spectral type may thus not be a strong indicator for the
credibility of a biosignature detection in the context of
the UV Threshold Hypothesis.

# 4.6. Overall prospects for testing the UV Threshold Hypothesis

Our results show that the UV Threshold Hypothesis is testable with potential future exoplanet surveys, but 700 that the success of such a test depends on the sample 701 size, the distribution of past NUV fluxes, and several 702 unknown astrophysical nuisance parameters. Even un-703 der idealized conditions, obtaining strong evidence for or 704 against the hypothesis likely requires sample sizes on the 705 order of 100 (see Section 3.1). This is true for a future 706 transit survey, the specifics of which we have reflected in 707 our Bioverse simulations (see Section 3.2). However, we 708 have shown that the impacts from the combined require-709 ments of the UV Threshold Hypothesis on the fraction 710 of inhabited planets in a sample are comparable in the 711 non-transiting case. The likely required sample sizes are 712 larger than the projected sample sizes of most planned missions; for example the Habitable Worlds Observatory 714 (HWO) expects to characterize a sample of only  $\sim 25$ 715 Earth analogs (Mamajek & Stapelfeldt 2023; Tuchow 716 et al. 2024).

Meaningful tests of the UV Threshold Hypothesis may thus require a different survey strategy that increases the sample of characterized planets. The Nautilus mission (Apai et al. 2019, 2022) aims to characterize up to  $\sim 1000$  EEC via utilizing an innovative telescope design.

#### DANIEL wants to work on this paragraph

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To prepare for future biosignature surveys, it is important to refine predictions on the role of UV radiation in prebiotic chemistry with both theoretical and experimental work.

## 4.7. Caveats

Our work is based on a number of assumptions and simplifications that may affect the results and conclusions. We discuss some of these caveats here.

#### 4.7.1. Existence of an atmosphere-crust interface

By its nature, cyanosulfidic scenario relies on rock surfaces exposed to the planetary atmosphere. Water worlds that have their entire planeary surface covered by oceans contradict this requireent and do not allow for the wet-dry cycling inherent to this origin of life scenario. The competition of tectonic stress with gravitational crustal spreading (Melosh 2011) sets the maximum possible height of mountains, which in the solar system does not exceed ~20 km. Such mountains will be permanently underwater on water worlds. Another impediment to wet-dry cycles may be tidal locking of the planet as it stalls stellar tide-induced water movement and diurnal irradiation variability. CITE!

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Our assumptions on past UV flux neglect the contribution of stellar flares, which may be hypothesized as an
alternative source of UV light (Buccino et al. 2007; Ranjan et al. 2017a). This concerns mainly ultracool dwarfs,
due to their low quiescent emission and high pre-main
sequence stellar activity (Buccino et al. 2007; West et al.
Joseph Grant Start show inadequate activity levels for a sufficient
start show inadequate activity levels for a sufficient
contribution through flares (Glazier et al. 2020; Ducrot

### 4.7.3. Atmosphere transmission

We do not take into account absorption of UV radiation by the planetary atmosphere. Theoretical work suggests that the atmosphere of prebiotic Earth was largely
transparent at NUV wavelengths with the only known
source of attenuation being Rayleigh scattering (Ranjan & Sasselov 2017; Ranjan et al. 2017a). We thus
approximated surface UV flux using top-of-atmosphere
fluxes. If there are planets in a sample that do not have
a transparent atmosphere at NUV wavelengths and require higher fluxes for abiogenesis, the fraction of inhabited planets in the sample will be lower. However,
these planets will not pollute the below-threshold subsample, as they will not be able to host life under the
UV Threshold Hypothesis.

# 5. CONCLUSIONS

We propose that specific origins-of-life scenarios may leave a detectable imprint on the distribution of biosignatures in exoplanet populations. We have investigated the potential of upcoming exoplanet surveys to test the hypothesis – motivated by the cyanosulfidic origins-of-life scenario – that a minimum past NUV flux is required for abiogenesis. To this end, we first employed a semi-analytical Bayesian analysis to estimate probabilities of obtaining strong evidence for or against this hypothesis. We then used the Bioverse framework to assess the diagnostic power of realistic transit surveys, taking into account exoplanet demographics, time-dependency of habitability and NUV fluxes, observational biases, and target get selection.

Our main findings are:

- 1. The UV Threshold Hypothesis of the cyanosulfidic scenario for the origins of life should lead to a correlation between past NUV flux and current occurrence of biosignatures that may be observationally testable.
- 2. The required sample size for detecting this correlation depends on the abiogenesis rate on temperate exoplanets and the distribution of host star properties in the sample; in particular their maximum

past NUV fluxes. Samples smaller than 50 planets are unlikely to yield conclusive results.

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- 3. Under the UV Threshold Hypothesis, the fraction of inhabited planets in a transit survey is sensitive to the threshold NUV flux and is expected to drop sharply for required fluxes above a few hundred erg s<sup>-1</sup> cm<sup>-2</sup>.
- 4. If the predicted UV correlation exists, obtaining strong evidence for the hypothesis is likely ( $\gtrsim 80\,\%$ ) for sample sizes  $\ge 100$  if the abiogenesis rate is high ( $\gtrsim 50\,\%$ ) and if no very high NUV fluxes are required. A survey strategy that targets extreme values of inferred past NUV irradiation increases the diagnostic power.
- 5. Samples of planets orbiting M dwarfs overall yield higher chances of successfully testing the UV Threshold Hypothesis. They may also be more likely to yield biosignature detections under this hypothesis.

check if new grid still supports this

Overall, our work demonstrates that future exoplanet surveys have the potential to test the hypothesis that a minimum past NUV flux is required for abiogenesis. More generally, we found that models of the origins of life provide hypotheses that may be testable with these surveys. Conducting realistic survey simulations with representative samples is important to identify testable science questions, support trade studies, help define science cases for future missions, and guide further theoretical and experimental work on the origins of life. Our work highlights the importance of understanding the context in which a biosignature detection is made, which can not only help to assess the credibility of the detection but also to test competing hypotheses on the origins of life on Earth and beyond.

The authors thank Kevin Heng, Dominik Hintz, and 833 Rhys Seeburger for insightful discussions. This ma-834 terial is based upon work supported by the Na-835 tional Aeronautics and Space Administration under 836 Agreement No. 80NSSC21K0593 for the program "Alien Earths". The results reported herein bene-838 fited from collaborations and/or information exchange within NASA's Nexus for Exoplanet System Science 840 (NExSS) research coordination network sponsored by 841 NASA's Science Mission Directorate. This work has 842 made use of data from the European Space Agency 843 (ESA) mission Gaia (https://www.cosmos.esa.int/gaia), 844 processed by the Gaia Data Processing and Analysis 845 Consortium (DPAC, https://www.cosmos.esa.int/web/ 846 gaia/dpac/consortium). Funding for the DPAC has been provided by national institutions, in particular the institutions participating in the Gaia Multilateral Agreement.

### AUTHOR CONTRIBUTIONS

M.S., D.A., and S.R. conceived the project, planned its implementation, and interpreted the results. M.S. de-

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veloped the planetary evolution component to Bioverse, 854 carried out the hypothesis tests and statistical analyses, and wrote the manuscript. D.A. leads the "Alien 856 Earths" program through which this project is funded, 857 helped to guide the strategy of the project, and provided 858 text contributions. A.A. carried out the semi-analytical s59 computations regarding the correlation of past UV flux and biosignature occurrence. S.R. advised on planetary 861 NUV flux evolution and the cyanosulfidic scenario of the 862 origins of life. R.F. wrote the initial draft of the Introand advised on the evolutionary biology aspects 864 of the project. K.H.-U. contributed to the Bioverse soft-865 ware development and simulations. T.L. supported the selection of testable hypotheses and provided text con-867 tributions to the initial draft. S.M. advised on the scope s68 of the project and supported the selection of testable 869 hypotheses. All authors provided comments and sug-870 gestions on the manuscript.

# REPRODUCIBILITY

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