

# Microbes as Agents of Change: Historical, Cultural, and Astrobiological Perspectives

Tran Quoc Hoang<sup>a,1,\*</sup>, Nguyen Hoang Tuong Vy<sup>a</sup>, Nguyen Vy Van<sup>a</sup>, Nguyen Le Ngoc Vy<sup>a</sup>, Nguyen Hoang Quynh Anh<sup>a</sup>

<sup>a</sup>*International University - VNU-HCM, School of Biotechnology, Quarter 33, Linh Xuan Ward, Ho Chi Minh City, Vietnam, 71309<sub>b</sub>,*

---

## Abstract

This is the abstract. Lorem ipsum dolor sit amet, consectetur adipiscing elit. Vestibulum augue turpis, dictum non malesuada a, volutpat eget velit. Nam placerat turpis purus, eu tristique ex tincidunt et. Mauris sed augue eget turpis ultrices tincidunt. Sed et mi in leo porta egestas. Aliquam non laoreet velit. Nunc quis ex vitae eros aliquet auctor nec ac libero. Duis laoreet sapien eu mi luctus, in bibendum leo molestie. Sed hendrerit diam diam, ac dapibus nisl volutpat vitae. Aliquam bibendum varius libero, eu efficitur justo rutrum at. Sed at tempus elit.

*Keywords:* keyword1, keyword2

---

**Submission Note:** Paper prepared for the Microbiology (BT321IU) course, International University - VNU-HCM.

## 1. Introduction

Microbes are among the most abundant and influential life forms on Earth, contributing substantially to planetary biomass and shaping ecosystems at every level despite their microscopic size. Throughout history, microorganisms have reshaped human societies, influenced cultural practices, and catalyzed scientific breakthroughs that redefine modern life. From pandemics that altered civilizations to extremophiles that challenge our understanding of life's limits, microbes exhibit a level of ubiquity and power that is both surprising and foundational. Microbiology, the scientific discipline devoted to the study of microorganisms, reveals the extraordinary breadth of microbial influence across both natural and human systems. As such, microbes are not simply sources of disease, but powerful drivers of historical change, cultural production, and emerging scientific frontiers such as astrobiology. Their largely unseen impact spans multiple disciplines and eras, highlighting the deep interconnectedness between microbial life and human advancement.

This paper is organized into four principal sections to examine these multidimensional contributions of microorganisms. The first explores microbes that changed history by analyzing the social and medical consequences of pandemics such as smallpox and the Black Death. The

second focuses on pioneering microbiologists whose discoveries transformed science and medicine, including Pasteur, Koch, and Fleming. The third investigates microbial influence in art and culture, with emphasis on bacterial art and natural dye production. The final section extends the discussion beyond Earth, examining extremophiles, microbial survival under space-like conditions, and their implications for astrobiology.

Rather than treating microbes as passive background conditions to which humans merely respond, this paper argues that microorganisms function as active agents of change—entities whose biological properties constrain, redirect, and at times override human intentions across history, culture, and science. By shaping demographic collapse, enabling new artistic media, and redefining planetary habitability, microbes do not simply influence human systems; they participate in them. Viewing microbes as agents rather than environmental noise clarifies why they repeatedly reorganize social structures, cultural meaning, and scientific definitions of life itself.

## 2. Microbes that Changed History

This section examines the all-encompassing influence of pathogenic microbes on human civilization, focusing on two specific agents that altered demographic, economic, and social structures globally. Rather than acting as passive causes of disease, these microorganisms functioned as active historical forces, reshaping political systems, economic structures, and patterns of human belief on a global scale.

### 2.1. The Black Death and Yersinia pestis

The Black Death, a devastating pandemic caused by the bacterium *Y. pestis* that peaked between 1346 and 1353, serves as a primary example of microbial impact on human

---

\*Corresponding author

Email addresses: [tranquochoang@protonmail.com](mailto:tranquochoang@protonmail.com) (Tran Quoc Hoang), [BTBTIU24133@student.hcmiu.edu.vn](mailto:BTBTIU24133@student.hcmiu.edu.vn) (Nguyen Hoang Tuong Vy), [BTBTIU24126@student.hcmiu.edu.vn](mailto:BTBTIU24126@student.hcmiu.edu.vn) (Nguyen Vy Van), [BTBTIU24131@student.hcmiu.edu.vn](mailto:BTBTIU24131@student.hcmiu.edu.vn) (Nguyen Le Ngoc Vy), [BTBTIU24007@student.hcmiu.edu.vn](mailto:BTBTIU24007@student.hcmiu.edu.vn) (Nguyen Hoang Quynh Anh)

<sup>1</sup>Student ID: BTBTWE24036

history. The disease is believed to have traveled along the Silk Road from Central Asia to Europe, facilitated by the expansion of trade routes and the movement of armies [1, 2]. Upon reaching Europe, the plague caused a demographic collapse of unprecedented scale, claiming an estimated 50% of the population in affected regions of Europe and Asia [3, 4].

The massive loss of life triggered profound societal and economic shifts. The sudden scarcity of labor empowered the surviving peasantry, which led to the decline of the feudal system as workers demanded higher wages and better conditions [5, 6]. The psychological trauma of the pandemic also reshaped religious and social attitudes, manifesting in extreme movements such as the flagellants and the persecution of minority groups [7, 8]. In this sense, *Yersinia pestis* acted not merely as a pathogen, but as a catalyst for systemic change, accelerating social transformations that might otherwise have taken centuries.

The transmission cycle of *Y. pestis* historically involved the black rat (*Rattus rattus*) and the rat flea (*Xenopsylla cheopis*) [9]. After feeding on an infected host, *Y. pestis* blocks the flea's gut, causing regurgitation of the bacteria into new hosts during later bites [10]. While bubonic plague was the most common form, the pneumonic form allowed for direct person-to-person transmission—thereby accelerating the spread [11, 12]. Fig. 1 provides a visualized diagram of the transmission pathway. The biological characteristics of *Y. pestis*—meaning its zoonotic reservoir, vector-based transmission, and capacity for airborne spread—directly determined the speed, scale, and severity of its historical impact.

## 2.2. Smallpox and Global Eradication

Shifting focus to another microbial agent that fundamentally altered human history: Smallpox. Caused by the *Variola major* virus and carrying a fatality rate of  $\approx 30\%$ , the disease played a decisive role in the colonization of the Americas. Following the arrival of Europeans, smallpox decimated indigenous populations, such as the Aztecs and Incas, who lacked prior immunity [14]. This “virgin soil” epidemic killed tens of millions—perhaps up to 95% of the indigenous population of the Americas—far exceeding the deaths caused by warfare and facilitating the collapse of indigenous empires [15, 16].

The fight against smallpox marked the birth of immunology. In 1796, Edward Jenner observed that milkmaids who contracted cowpox (a mild disease) were immune to smallpox. He tested this hypothesis by inoculating a young boy with material from a cowpox lesion and subsequently exposing him to smallpox; the boy remained healthy [17, 18]. Jenner’s work established the principle of vaccination, a term derived from *vacca* (Latin for cow), replacing the riskier practice of variolatio, an earlier technique that intentionally induced a mild smallpox infection to confer immunity [19].

Smallpox holds the distinction of being the first and only human infectious disease to be eradicated. Following a massive global campaign by the World Health Organization (WHO) involving vaccination, surveillance, and quarantine, the disease was declared eradicated in 1980 [20, 21]. As such, the disease marks a turning point in human–microbe relations, where scientific understanding allowed humans strategically counter microbial influence.

Together, the Black Death and smallpox demonstrate that microbial impact is not limited to mortality statistics but extends to structural reorganization of societies. In both cases, the biological features of the pathogen—mode of transmission, immunity dynamics, and host susceptibility—directly shaped political power, labor systems, and colonial outcomes. These outcomes were not accidental side effects of disease but emergent properties of microbial life interacting with human networks. In this sense, microbes acted less like external shocks and more like historical actors whose constraints forced new social equilibria.

## 3. Famous Microbiologists and Groundbreaking Discoveries

This section examines how key microbiologists transformed microbes from invisible forces of fate into identifiable, controllable agents—thereby reshaping history, medicine, and society.

### 3.1. Louis Pasteur: A Core Founder of Modern Microbiology

#### 3.1.1. Germ Theory

Louis Pasteur, a French chemist, fundamentally changed the understanding of biology by disproving the theory of spontaneous generation. Through his elegant experiments using swan-necked flasks, he demonstrated that microorganisms in the air, not a “vital force,” were the cause of contamination in sterile broth [22]; see Fig. 8]. Pasteur’s work laid the foundation for the germ theory of disease, which establishes that specific invisible microbes are responsible for specific illnesses [23].

#### 3.1.2. Pasteurization

In addition to establishing the germ theory, Pasteur applied his knowledge of fermentation to solve problems in the French wine and silk industries. He discovered that heating liquids to a specific temperature (60–90°C) killed the bacteria responsible for spoilage without destroying the product. This process, known as pasteurization, revolutionized food safety and is still widely used today for milk and other beverages [24].

#### 3.1.3. Vaccine Development

Pasteur also pioneered the development of laboratory-attenuated vaccines. He created vaccines for chicken

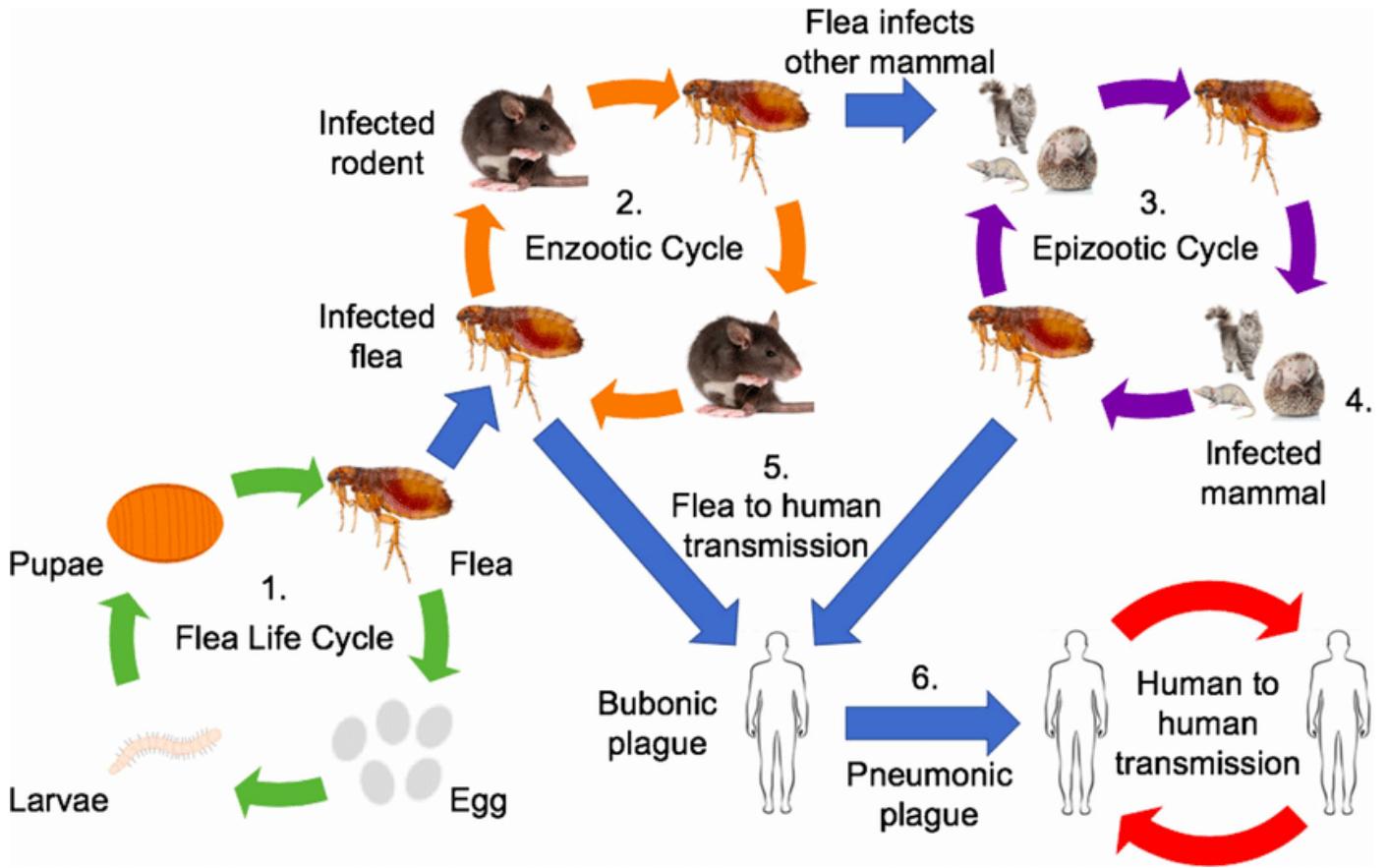


Fig. 1: Mechanisms of Plague (*Y. pestis*) Transmission and Disease Progression. The cycle begins with (1) the Flea Life Cycle: Eggs, laid in moist soil/burrows, hatch into larvae that feed on rodent feces and develop into pupae and adult fleas. Infected adult fleas parasitize rodents. (2) The Enzootic Cycle maintains *Y. pestis* among rodent hosts and reservoirs, primarily vectored by fleas (*X. cheopis*, *X. brasiliensis*, and *S. fonsquerniei*). (3, 4) Epizootic Spills occur when fleas infect non-host mammals, which often lack resistance, rapidly succumb to the disease, and spread the plague geographically. (5) Human Zoonotic Transmission: Humans are typically infected via the bite of an infected flea from a rodent or secondary mammal host. (6) Human-to-Human Spread: Bubonic plague may progress to a highly contagious pneumonic plague via lung infection, enabling *Y. pestis* transmission through infectious respiratory droplets generated by coughing. Reproduced from Mackay-Alderson J, Quastel M, Wilson E, Bellamy D [13].

cholera and anthrax by using weakened strains of the bacteria [25]. His most famous medical achievement was the development of the rabies vaccine in 1885, which he successfully used to save the life of Joseph Meister, a boy bitten by a rabid dog [26, 27].

### 3.2. Alexander Fleming: The Serendipitous Discovery

#### 3.2.1. Discovery of Penicillin

In 1928, Scottish bacteriologist Alexander Fleming made a serendipitous discovery that launched the antibiotic era. Upon returning from a holiday, he noticed that a mold, *Penicillium notatum*, had contaminated a Petri dish of *Staphylococcus* bacteria. Crucially, he observed that the bacteria surrounding the mold had been destroyed [28].

#### 3.2.2. Impact on Medicine

Fleming identified the mold's active substance as penicillin. Although mass production was later achieved by Howard Florey and Ernst Chain during World War II, Fleming's discovery provided the first effective treatment

for bacterial infections that had previously been fatal [29, 30]. Penicillin was hailed as a "miracle drug," saving millions of lives from wound infections and diseases like syphilis and pneumonia, though Fleming himself warned early on about the potential for antibiotic resistance [31].

### 3.3. Robert Koch and Koch's Postulates

#### 3.3.1. Methodological Rigor

Robert Koch, a German physician, is often regarded as one of the founders of medical microbiology alongside Pasteur. He introduced rigorous scientific methods to the field, most notably "Koch's Postulates." This set of four criteria is used to establish a causal relationship between a specific microbe and a specific disease: (1) the microbe must be present in every case of the disease; (2) it must be isolated and grown in pure culture; (3) inoculation of the culture into a healthy host must reproduce the disease; and (4) the microbe must be recovered again from the experimentally infected host [32, 33].

### 3.3.2. Key Discoveries

Using these methods, Koch identified the specific causative agents of several deadly diseases, including *Bacillus anthracis* (anthrax), *Vibrio cholerae* (cholera), and *Mycobacterium tuberculosis* (tuberculosis) [34, 35, 36]. His work provided the definitive proof for the germ theory of disease and established the standards for diagnostic microbiology [37]. Though foundational, Koch's Postulates are limited when applied to viruses, unculturable organisms, and polymicrobial diseases.

Pasteur, Koch, and Fleming are often presented as isolated geniuses, but their collective impact lies in how they transformed microbes from invisible fate into governable entities. Germ theory reframed illness as a biological interaction, antibiotics redefined survivability, and laboratory standards stabilized microbial knowledge across institutions. The rise of antibiotic resistance illustrates that microbial evolution continues to negotiate, resist, and sometimes overturn human control, reinforcing the argument that microbes remain active participants rather than solved problems.

## 4. Microbes in Art and Culture

This section explores the scientific mechanisms underlying microbial involvement in aesthetic and cultural domains, focusing on bacterial pigment production as a medium for art and the role of fungi and yeasts in creating sustainable dyes for textiles.

### 4.1. Bacterial Art (Agar Art/Bio-Painting)

Bacterial art is a form of art resulting from the special combination of microbiology with the power of visual expression by utilizing the Petri dish as a living canvas. This form of art relies entirely on the underlying biological mechanisms by which certain bacterial species synthesize brightly colored pigments [38].

The medium consists of nutrient-rich agar plates, where different bacterial strains are inoculated and cultured to grow into intricate images. Vibrant colors are generated by chromogenic bacterial secondary metabolites produced through distinct metabolic and biosynthetic pathways, which makes it easier for scientists and artists to isolate and identify the strains that produce the compounds in sufficient amount [39].

Two bacterial species employed in bio-painting are widely studied for their distinctive pigment production. The first is *Serratia marcescens*, a Gram-negative bacterium that is also a producer of prodigiosin. Prodigiosin is a red pigment belonging to the prodiginine group, characterized by a tripyrrole chemical structure, and has been successfully used in textile coloring processes [40]. The second example is *Chromobacterium violaceum*. This strain produces violacein, a distinctive bisindole-violet or blue pigment [41].

Violacein and prodigiosin are well-known hydrophobic bacterial chromogenic pigments, and these two are responsible for the purple and red color phenotype of the bacterial strain. Fig. 7 shows the chemical structure and the colored phenotypes of the bacterial strains that produce these compounds.

By integrating microbiological techniques with the principles of aesthetic design, bacterial art blurs the traditional boundaries between science and art. Fig. 2 shows the representation of various colors producing microorganisms on a Petri plate. Fig. 3 shows agar art from living microbes. Those are examples of colorful bacteria applied to agar in an artistic manner (A-C) ASM Agar Art Contest winners.



Fig. 2: Representation of various colors producing microorganisms on a Petri plate. Reproduced from Tuli HS, Chaudhary P, Beniwal V, Sharma AK [40].

Fig. 9 shows The 2016 American Society for Microbiology Agar Art Contest winner, The First Race, illustrated fertilization using four pigment-producing bacterial species grown on selective agar. Created by Md Zohorul Islam (University of Copenhagen), the artwork used *Staphylococcus aureus* (red), *S. xylosus* (green), *S. hyicus* (white), and *Corynebacterium glutamicum* (yellow), with additional hues produced by mixing these microbes.

### 4.2. Fungi in Pigmentation and Textiles

The rapid development of greener approaches in industrial processes, such as textile dyeing, has been driven by the environmental concerns regarding synthetic dyes [43]. In this context, fungi have become a promising sources of natural pigments.

Fungi have shaped the history of natural dyes for centuries, with uses like ancient lichen dyes extracted from *Roccella* and *Ochrolechia* species for purples and violets. There were many traditional dyes made from lichens in the Scottish Highlands, including red dyes from the cudbear lichen (*Lecanora tartarea*), the common orange lichen (*Xanthoria parietina*), and several species of leafy *Parmelia*

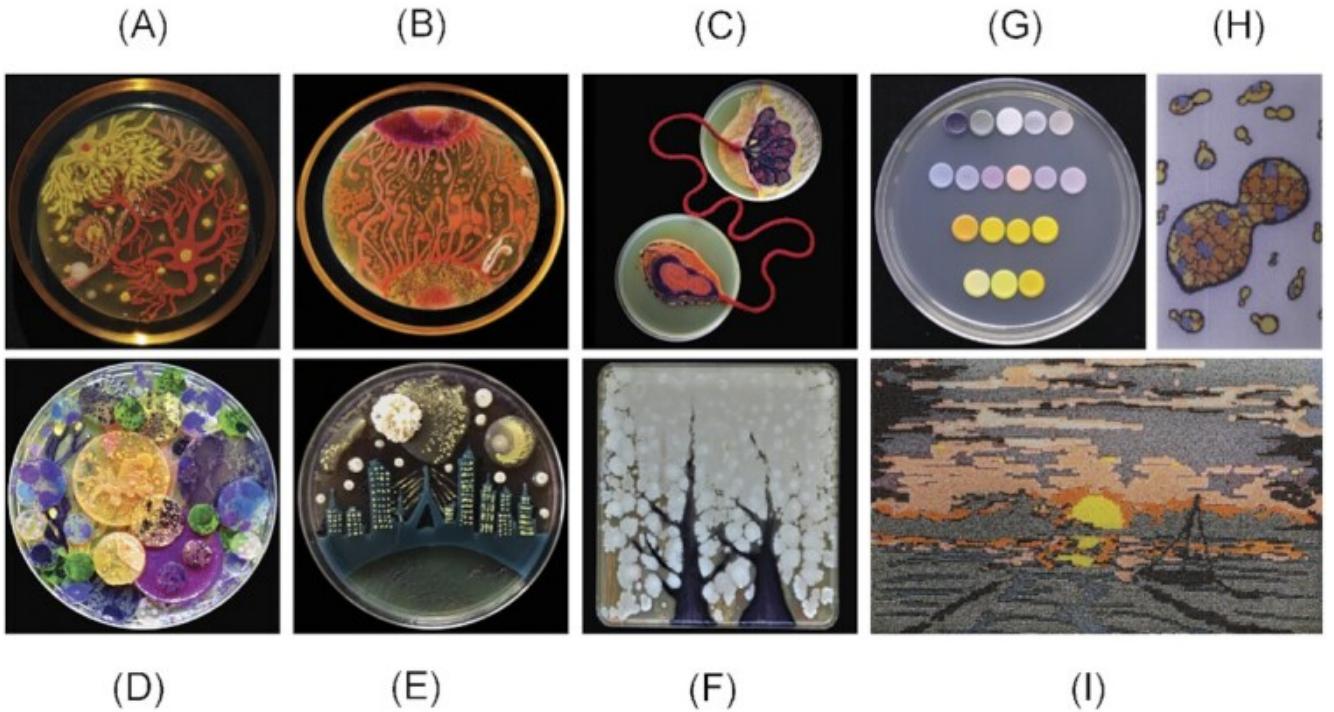


Fig. 3: Examples of Agar Art from Living Microbes. Panels (A-C) display winning entries from the ASM Agar Art Contest. (A) "Cell to cell" (2015) using yellow *Nesterenkonia*, orange *Deinococcus*, and *Sphingomonas*. (B) "Neurons" (2015) using red *Serratia*, yellow *Nesterenkonia*, orange *Deinococcus*, and *Sphingomonas*. (C) "Sustenance" (2018): (Top) pink colonies, orange *Nesterenkonia*, and *Deinococcus radiodurans*. (Bottom) Dark-violet recombinant *E. coli* (violacein pathway), red *Serratia marcescens*, and white *Bacillus*. (D) "Remainders": microbial paintings on food-colored agar plugs. (E) "Boston skyline" (2019): sculpted and collaged agar and bacteria. (F) "Bacillus surprise": dark-violet recombinant *E. coli* and white *Bacillus*. (G) A palette of colored recombinant yeast (*Saccharomyces cerevisiae*) expressing chromogenic proteins (black/grey/purple from violacein, blue from anemone gene, pink from RFP, and various yellows/oranges from beta-carotene). (H) "Puzzle Pieces" (2017) and (I) "Sunset at the End" (2016): patterns printed using engineered *S. cerevisiae*. Reproduced from Frankel E, Temple J, Dikener E, Berkmen M [42].

lichens. During the 15th and 17th centuries, purple lichen dyes continued to play an important role in the European dyeing industry [44]. And today, the traditional practices have evolved into modern bio-dyes via fermentation to serve the demand of today's sustainable textile production [45].

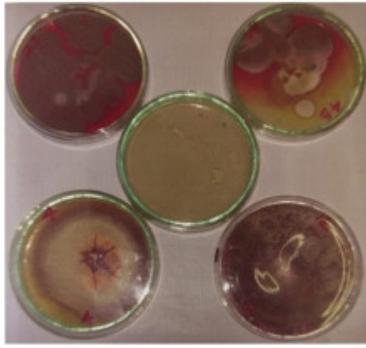
From an industrial perspective, most fungal pigments are water-soluble, making production and extraction relatively straightforward. They can be easily cultivated and scaled up in industrial fermentation systems and can be harvested without the need for organic solvents, enhancing their sustainability and environmental compatibility (see Fig. 4). For reference, Table 2 shows a list of fungal pigments produced on an industrial scale.

Fungal pigments cover a broad color spectrum, such as yellow, orange, red, purple, blue, brown, and black. Key producers are *Monascus* spp. (red, yellow, orange), *Penicillium* spp. (red, yellow, brown), *Aspergillus* spp. (yellow, brown), *Fusarium* spp. (pink, violet, red), and *Blakeslea trispora* (-carotene) [44]. Numerous fungal genera have demonstrated efficient pigment production [47]. For instance, *Penicillium brevicompactum* can produce a mixture of yellow, orange, and red pigments [48] while *Penicillium*

*purpurogenum* generates red pigments with strong dyeing performance on fabrics such as wool, silk and polyester. Fig. 5 shows the effect of red pigment produced by *Penicillium purpurogenum* application on different fabrics (wool, silk, polyester, viscose). Furthermore, different fungal species with their active pigment for application in the textile industry are shown in Fig. 10. Fungal pigments have demonstrated successful dyeing performance on a wide range of textile materials, including cotton, linen, silk, and several synthetic fibers [49, 50]. Table 1 shows the fungal pigments and their application in the textile industry.

## 5. Bacteria in Space (Astromicrobiology)

Astromicrobiology investigates how microorganisms such as bacteria and archaea survive, adapt, and behave under the extreme physical and chemical conditions found beyond Earth. These are known as extremophiles and include organisms adapted to extremes of temperature, pressure, radiation, salinity, and pH, inhabiting environments such as hydrothermal vents, permafrost, hypersaline lakes, and acidic waters [51]. Their survival depends on their metabolisms and cellular structures evolved specifically for these extremes. Accurately simulating extraterrestrial ul-



**A) Growth of pigmented fungi in agar medium**



**B) Maintenance of pigmented fungi**

1. Control
2. *Penicillium purpurogenum*
3. *Paecilomyces farinosus*
4. *Emericella nidulans*
5. *Fusarium moniliforme*
6. *Monascus purpureus*



**C) Scale-up of pigment producing fungi in fermenter**

Fig. 4: Scale-up process for pigment-producing fungi from laboratory culture to industrial fermentation. The process moves from initial isolation and growth in a Petri dish to larger-scale production. (A) Growth and visualization of pigmented fungi on solid agar medium (Petri dish). (B) Maintenance and preservation of the fungal strain for long-term use. (C) Scale-up of the culture in a liquid medium within a fermenter, which is a critical step for sustainable and high-volume industrial pigment production. Reproduced from Venil CK, Velmurugan P, Dufossé L, Renuka Devi P, Veera Ravi A [46].



Fig. 5: Effect of red pigment produced by *Penicillium purpurogenum* application on different fabrics (wool, silk, polyester, viscose). Reproduced from Elkhatib W, Elnahas M, Daba G [50].

traviolet (UV) radiation, which is significantly more intense and has a wider wavelength range than that which reaches Earth's surface, is one of the main challenges in astromicrobiology [52]. The solar UV spectrum cannot be

completely replicated in space by even the most powerful artificial UV generators on Earth. Nevertheless, simulation studies are useful in spite of this drawback. As an illustration of how extremophiles might take advantage of the chemical environment of other planets, the acidophilic bacterium *Acidithiobacillus ferrooxidans* was found to grow successfully in simulated Martian regolith, even without additional nutrients. In true space conditions, however, the most informative results come from orbital experiments such as the ADAPT study, which exposed genetically UV-resistant *Bacillus subtilis* spores to space radiation in Low Earth Orbit (LEO) [53]. Fig. 11 provides an overview of the interconnected medical and microbiological challenges associated with spaceflight and long-term habitation of extraterrestrial environments, illustrating how microbial survival, human health, and planetary conditions are tightly linked. As such, this section argues that microbial survival in space fundamentally re-

Table 1: Fungal pigments and their application in the textile industry

Fungi	Pigment	Color	Fabrics
<i>Alternaria alternata</i>	Anthraquinones	Reddish-brown	Cotton
<i>Penicillium Oxalium</i>	Anthraquinones	Arpink red	Wool
<i>Chlorociboria aeruginosa</i>	Quinones	Green	Bleached cotton, spun polyacrylic, spun polyamide, spun polyester, worsted wool
<i>Scytalidium cuboideum</i>		Red	
<i>Scytalidium ganodermophthorum</i>		Yellow	
<i>Aspergillus</i> sp.	Quinones	Brown cotton	Silk, silk cotton
<i>Acrostalagmus</i> (NRC 90)	Quinones	Brown	Wool
<i>Alternaria alternata</i> (NRC17)	Quinones	Reddish-brown	Wool
<i>Alternaria</i> sp. (NRC 97)	Quinones	Brown	Wool
<i>Aspergillus niger</i> (NRC 95)	Quinones	Brown	Wool
<i>Bisporomyces</i> sp. (NRC 63)	Quinones	Deep brown	Wool
<i>Penicillium murcianum</i>	Cartenoids	Yellow	Wool
<i>Talaromyces australis</i>	2,4-Di-tert-butylphenol	Red	Cotton fabric
<i>Phoma harbarum</i>	Magenta pigment	Magenta	Nylon
<i>Talaromyces verruculosus</i>	Polyketide	Red	Cotton fabric
<i>Monascus purpureus</i>	Rubropunctamine	Red	Wool

Note. Fungal pigments and their application in the textile industry. Adapted from Elkhateeb W, Elnahas M, Daba G [50].

shapes how we define habitability, contamination, and the origin of life.

### 5.1. Extremophiles Defined

Extremophiles are organisms, mostly bacteria and archaea, that not only survive but flourish in environments previously thought to be incompatible with life. This applies to a wide range of settings, including high pressure, temperature, pH, radiation, salinity, and nutrient availability, since they exist in severe habitats [54]. Extremophiles offer crucial insights about the boundaries of life on Earth and assist scientists in determining the potential habitability of alien settings since they live in such harsh circumstances.

Extreme settings can be either naturally occurring or man-made; these microbes are just the result of evolution and the best adapted to their surroundings. Thousands of new species are found and named by scientists each year. Microorganisms have played a significant role in this massive expansion of species discoveries in recent years. Other terms that are used to categorize particular kinds of extremophiles include: Low pH acidophiles, high pH alkaliphiles, anaerobic extremophiles (anti-oxygen), cryophiles (thrive in cold temperatures), piezophiles (high pressure), psychrophiles (thrive in low temperatures), thermophiles (live in temperatures above 40°C), hyperthermophiles (above 80°C), xerophiles, methanogens, Toxitolerance Deep-sea hydrothermal vents, ice sheets and permafrost, volcanic systems and hot springs, salt pans and hypersaline lakes, and extremely acidic or alkaline waters are just a few examples of the various types of extreme conditions found in nature [55, 56]. The wide range of extreme habitats in which these organisms thrive is summarized in

Fig. 6, which illustrates the diversity and spatial distribution of extreme environments within Earth's crust.

By surviving in such environments, extremophiles reveal the true boundaries of Earth-based life. This helps astrobiologists evaluate the potential habitability of extraterrestrial environments such as Mars, Europa, Enceladus, or exoplanets, and understand the biochemical strategies that life might use elsewhere. Extremophilic habitats may occur naturally (e.g., hydrothermal vents, volcanoes, permafrost) or be produced by human activity (industrial brines, highly acidic mining runoffs). Regardless of origin, extremophiles are the best-adapted products of evolution within these conditions [57].

### 5.2. Survival on the International Space Station (ISS)

In the space experiment “Molecular adaption strategies of microorganisms to different space and planetary UV climate conditions” (ADAPT), bacterial endospores of the highly UV-resistant *Bacillus subtilis* strain MW01 were exposed for 559 days on the ESA EXPOSE-E facility mounted outside the ISS. These facilities allow biological samples to be exposed directly to Space vacuum, Full-spectrum solar UV radiation (> 110 nm), Cosmic ionizing radiation, Rapid temperature cycling, Simulated Martian surface conditions [58]. The experimental configuration of the EXPOSE-E facility and the placement of the ADAPT samples on the exterior of the ISS are shown in Fig. 12.

One of the main goals of Mars exploration missions is to find organic compounds on the planet's surface. Therefore, comprehending the preservation of organic matter in the Martian environment is a crucial step in interpreting interpreting data to be obtained in the future. The organic substances (glycine, serine, phthalic acid, phthalic acid in the

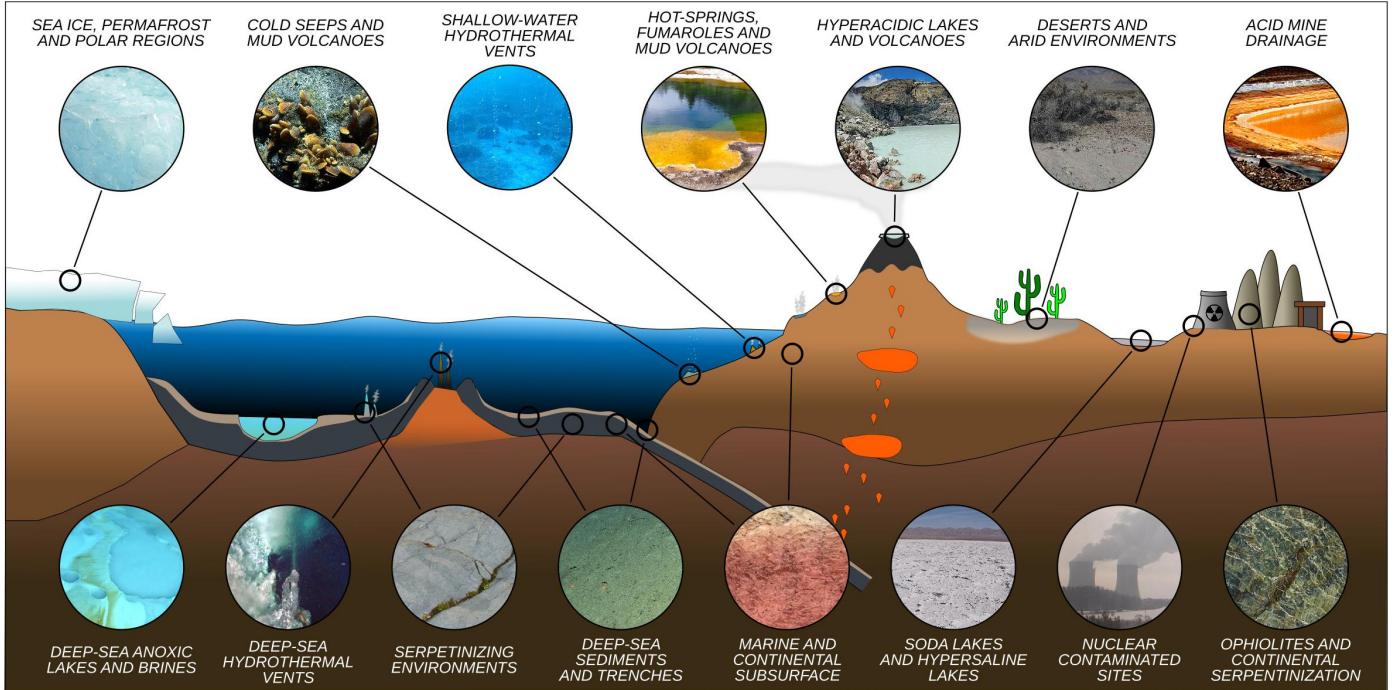


Fig. 6: Representative idealized cross section of Earth’s crust showing the diversity of extreme environments and their approximate location. Reproduced from Merino et al. [55].

presence of a mineral phase, and mellitic acid) completely degraded after a 1.5-year exposure to Mars-like surface UV radiation conditions in space. Under Martian surface conditions, their half-lives ranged from 50 to 150 hours. Amino acids and a dipeptide in pure form and embedded in meteorite powder were exposed to space conditions for 18 months in order to study the chemical behavior of organic molecules in the space environment. The samples were then brought back to Earth and examined in a lab for reactions brought on by solar UV and cosmic radiation [59]. The findings demonstrate that the chemical makeup of the exposed molecules and the UV light’s wavelengths affect resistance to irradiation. Aspartic acid, aminobutyric acid, and the dipeptide were the most changed substances. Alanine, valine, glycine, and aminoisobutyric acid were the most resistant. The findings also show that meteorite powder has a protective effect, highlighting the significance of exogenous contributions to the prebiotic organics inventory on early Earth.

### 5.2.1. Findings

The findings attest to some microorganisms’ extraordinary resistance. In LEO, spores of the bacterium *Bacillus subtilis*, including the UV-resistant strain from the ADAPT study, endured an astonishing 1.5 years. Despite the catastrophic mix of vacuum, desiccation, and strong sun radiation, this survival took place. Furthermore, certain eukaryotic life forms, such as the green alga *Stichococcus* sp. and the rock-dwelling Antarctic fungi *Cryomyces antarcticus*, also endured the exposure, particularly when they had minimal shielding. This suggests that protection

afforded by even thin layers of soil or rock is sufficient to shield life from the worst effects of space, providing experimental support for the theory of lithopanspermia. In stark contrast, the basic organic molecules necessary for life, like amino acids glycine and serine, were found to be highly vulnerable. They rapidly degraded under UV radiation comparable to the Martian surface, showing half-lives of just 50 to 150 hours. This finding is essential for Mars exploration, suggesting that the absence of organic signatures detected on the planet’s surface may simply be the result of rapid photodegradation rather than an indication that life never existed there [59].

### 5.2.2. Forward Contamination

Experiments like EXPOSE have proved that some terrestrial bacteria are resilient in space, which raises serious worries about forward contamination. This is the inadvertent transfer of germs from Earth to other celestial bodies on spacecraft. Highly resilient species, like *Bacillus subtilis* spores or cryptoendolithic fungi (microbes that live in rocks), have demonstrated their ability to withstand the severe temperatures, radiation, and vacuum of space. They may be able to travel to and infect delicate locations like Mars, Europa, or Enceladus if they can withstand extended exposure in LEO [60]. If extraterrestrial ecosystems exist, such contamination could interfere with native biochemical processes or introduce false positives for life, which could jeopardize future astrobiological research. Because even a tiny number of hitchhiking microbes could alter the chemical environment, this possibility is particularly significant for missions intended to

find biosignatures or identify subsurface habitability [61]. To protect scientific integrity and the biological isolation of other worlds, international space organizations adhere to stringent COSPAR Planetary Protection requirements, which require rigorous sterilization procedures, cleanroom assembly, and microbial load reduction.

### 5.2.3. Panspermia

The scientific viability of panspermia, particularly lithopanspermia, the hypothesis that life could be carried between planets via rocks ejected by impacts is greatly strengthened by the demonstrated resistance of microbes in the ISS exposure tests [62]. Although the EXPOSE experiments lasted only 1.5 years, far shorter than the millions of years required for natural interplanetary travel, the ability of bacterial spores, lichens, and rock-inhabiting fungi to survive vacuum, cosmic radiation, and severe temperature cycles provides the first controlled evidence that certain microorganisms can endure the essential stages of space transfer [63]. If shielded inside rock pores or mineral matrices, these organisms could theoretically remain protected from destructive ultraviolet radiation and cosmic particles, increasing their likelihood of surviving long-duration transit. These findings do not prove that panspermia has occurred, but they meaningfully strengthen its feasibility by demonstrating that the biological component of interplanetary transport is physically possible. As a result, the concept of life dispersal across planetary systems, once purely speculative, now rests on experimental data showing that at least some Earth organisms possess the durability required to persist beyond their home planet.

## 6. Conclusion

The exploration of microbial life across history, culture, and science illuminates the multifaceted and often underestimated role of microorganisms in shaping human civilization and the cosmos. Microbes are not merely biological entities but historical agents, driving significant societal shifts—from the plagues that altered demographics and political power structures to the development of early fermented foods that sustained populations and helped launch biotechnology. Their influence extends far beyond the laboratory; microbes have inspired artists, fueled creative expression, and challenged our understanding of life through astrobiological studies aboard the ISS. Whether thriving in the scorching heat of hydrothermal vents, frozen deep within polar ice, or surviving direct exposure to cosmic radiation, microbes continually expand the boundaries of where life can exist and how it can adapt. Beyond the laboratory, microbes have also found a place in the creative sphere, serving as a unique living medium for bio-art and creating vibrant, growing installations that challenge traditional artistic boundaries and encourage reflection on life’s smallest forms. Finally, the field of astrobiology confirms that microbes are cosmic pioneers, surviving in extreme terrestrial environments that mimic con-

ditions on other planets. They are fundamental to the search for extraterrestrial life and will be vital for supporting long-duration human space missions.

Together, these examples demonstrate that microbes are powerful agents of change—shaping ecosystems, societies, culture, and our understanding of life beyond Earth. Far from passive or peripheral, microorganisms drive ecological processes, inspire artistic expression, and define the biological limits of habitability on Earth and elsewhere.

## 7. Abbreviations

Abbreviation	Full Term
ADAPT	Molecular adaption strategies of microorganisms to different space and planetary UV climate conditions
COSPAR	Committee on Space Research
EXPOSE-E	European Space Agency Exposure Facility E
ISS	International Space Station
LEO	Low Earth Orbit

Here is a clean, publication-ready *Author Contributions* section in the same style and structure as your previous one, with your own contribution intentionally omitted as requested:

---

## 8. Author Contributions

- Tran Quoc Hoang: Conceived the study and developed the overall methodology (Conceptualization, Methodology). Managed the project and compiled the manuscript in Quarto (Project Administration, Software). Performed literature synthesis and wrote *Section 2 “Microbes that Changed History”* and *Section 3 “Famous Microbiologists and Groundbreaking Discoveries”* (Investigation, Writing – Original Draft). Contributed to reviewing and editing the manuscript (Writing – Review & Editing).
- Nguyen Hoang Tuong Vy: Wrote *Section 5 “Bacteria in Space (Astromicrobiology)”* and the *Conclusion* (Writing – Original Draft). Contributed to reviewing and editing the manuscript (Writing – Review & Editing).
- Nguyen Vy Van: Wrote *Section 4 “Microbes in Art and Culture”* and the *Introduction* (Writing – Original Draft). Conducted related background investigation (Investigation). Contributed to reviewing and editing the manuscript (Writing – Review & Editing).
- Nguyen Le Ngoc Vy: Co-authored the *Abstract* and summarized the manuscript to prepare presentation slides (Data Curation, Visualization). Contributed

to reviewing and editing the manuscript (Writing – Review & Editing).

- Nguyen Hoang Quynh Anh: Co-authored the *Abstract* and summarized the manuscript to prepare presentation slides (Data Curation, Visualization). Contributed to reviewing and editing the manuscript (Writing – Review & Editing).

## **9. Declaration of Competing Interest**

The authors declare no competing interests.

## **10. Data/Code Availability**

The Quarto project underlying this paper—including the manuscript, figures, and bibliography—is available at <https://github.com/ht2905/historical-microbes>.

## **11. Appendix A: Supplementary Figures**

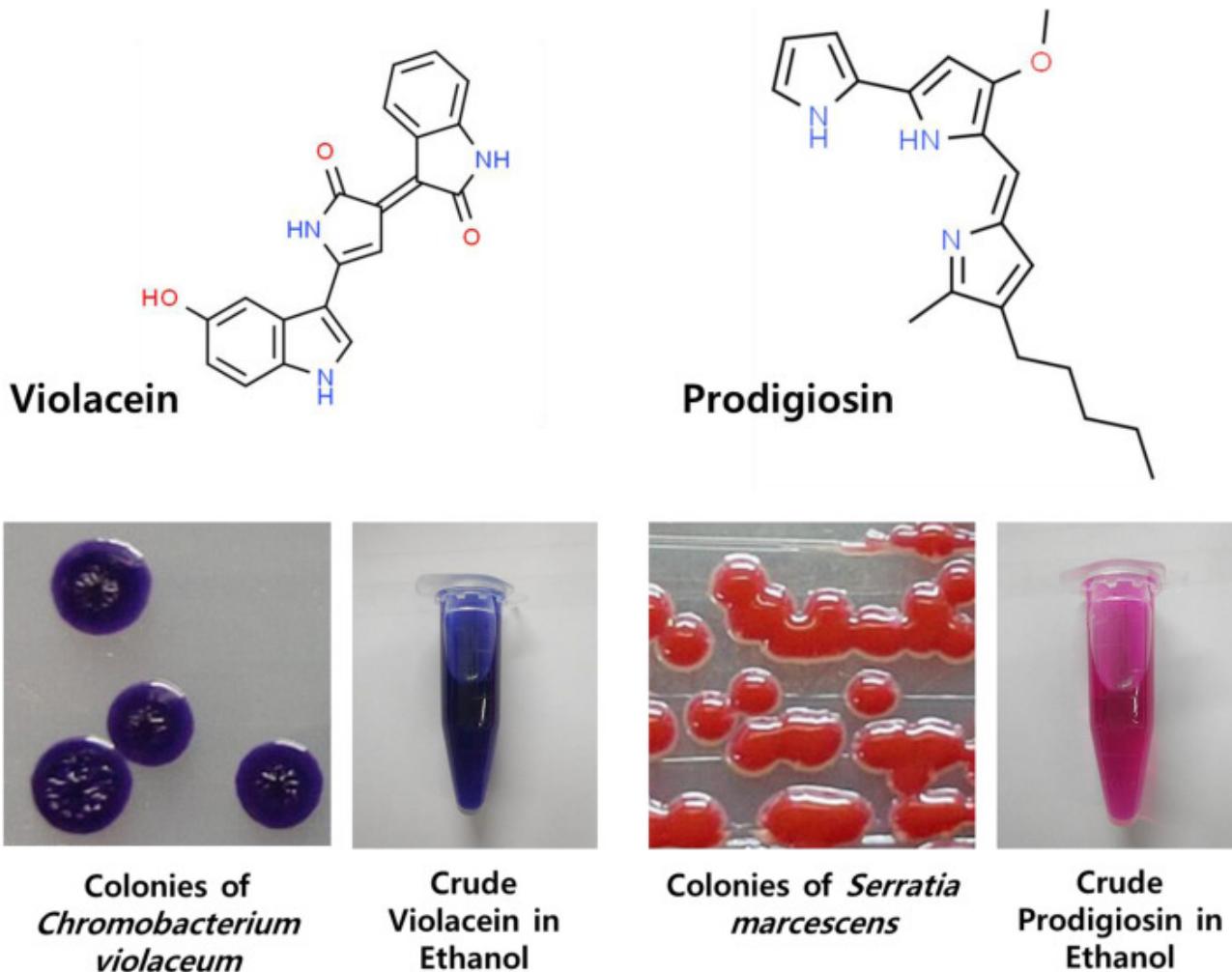


Fig. 7: Chemical structures of the bacterial pigments violacein (produced by *Chromobacterium violaceum*) and prodigiosin (produced by *Serratia marcescens*), shown alongside the respective colored phenotypes of the producing bacterial strains. Reproduced from Choi SY, Lim S, Yoon K, Lee JI, Mitchell RJ [39].

Table 2: List of fungal pigments produced on an industrial scale

Fungal species	Pigments
<i>Monascus</i> species	Ankaflavin (yellow), monascorubramine (red), rubropunctatin (orange)
<i>Ophiocordyceps unilateralis</i>	Erythrostominone (red), 3,5,8-TMON (red)
<i>Blasckeslea tripora</i>	$\beta$ -Carotene (yellow-orange), lycopene (red)
<i>Ashbya gossypii</i>	Riboflavin (yellow)
<i>Penicillium oxalicum</i>	Anthraquinone derivative (red), anthraquinones (red and other hues) Arpink red <sup>TM</sup> , secalonic acid D (yellow)

Note. List of fungal pigments produced on an industrial scale, synthesized from Caro Y, Venkatachalam M, Lebeau J, Fouillaud M, Dufosse L [66].

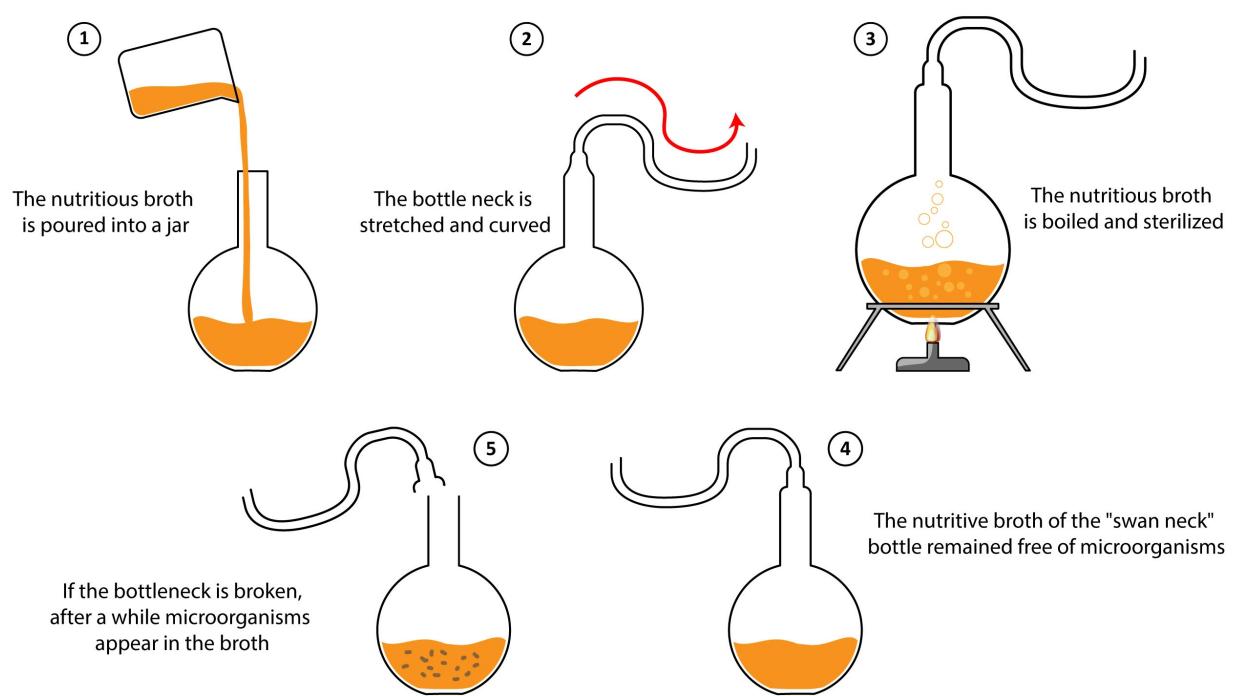


Fig. 8: Pasteur's swan-necked flask experiment illustrating how airborne microorganisms, rather than a “vital force,” contaminate sterile broth. Reproduced from Shutterstock [64].

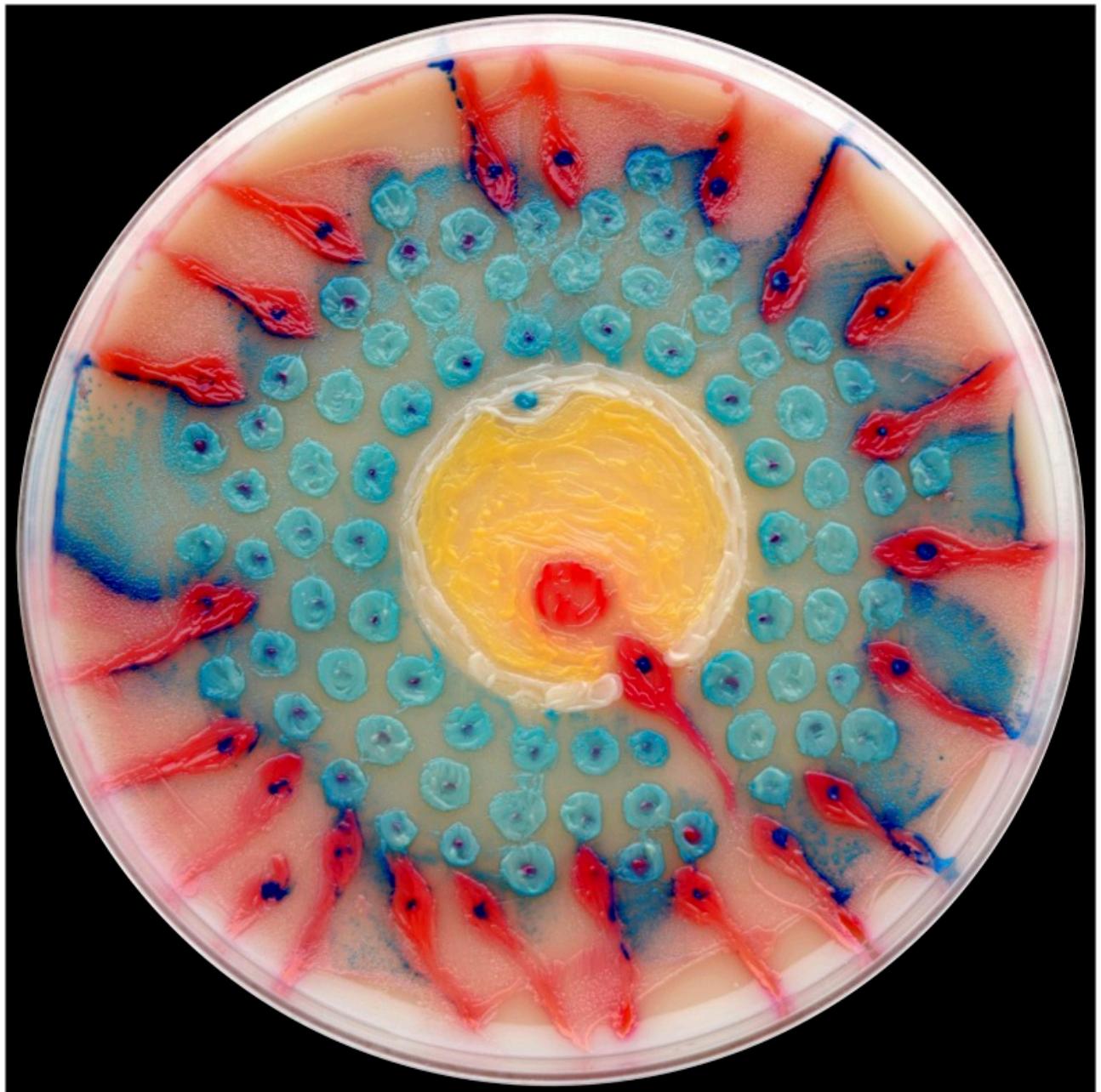
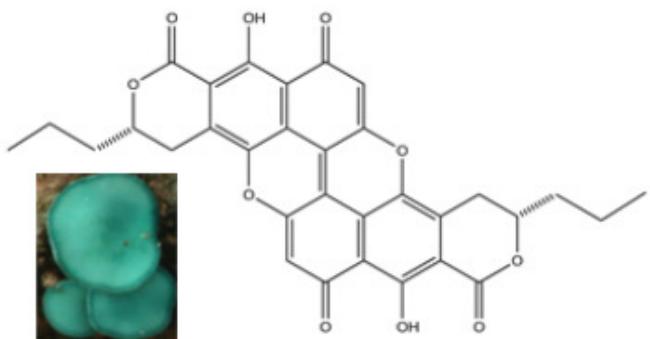
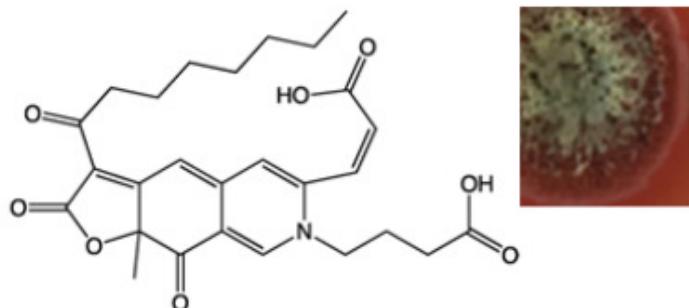
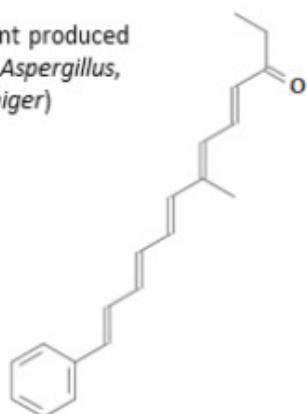


Fig. 9: The 2016 winner of the American Society of Microbiology's Agar Contest, titled The First Race, depicted fertilization. Graduate student Md Zohorul Islam of the University of Copenhagen "painted" with four bacteria on a selective agar canvas. The red was *Staphylococcus aureus*, a pathogen in both humans and animals. *Staphylococcus xylosus*, a commensal organism in human skin, generated green. The white was *Staphylococcus hyicus*, an animal pathogen responsible for grassy pig disease. And yellow came from *Corynebacterium glutamicum*, a bacterium used to produce amino acids, such as L-glutamate and L-lysine. Other colors came from mixing two or more of these microbes. Artwork by Md Zohorul Islam (University of Copenhagen, Copenhagen). Image courtesy of American Society for Microbiology. Reproduced from Madhusoodanan Y [65].

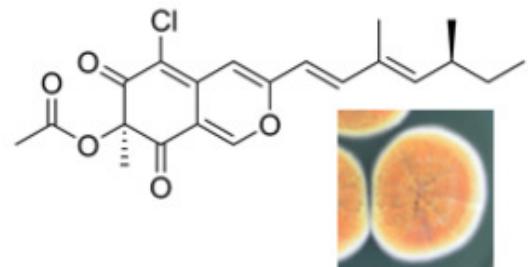


Xylindein (quinone pigment, dimeric naphthoquinone derivative, produced by fungi from genus *Chlorociboria*)

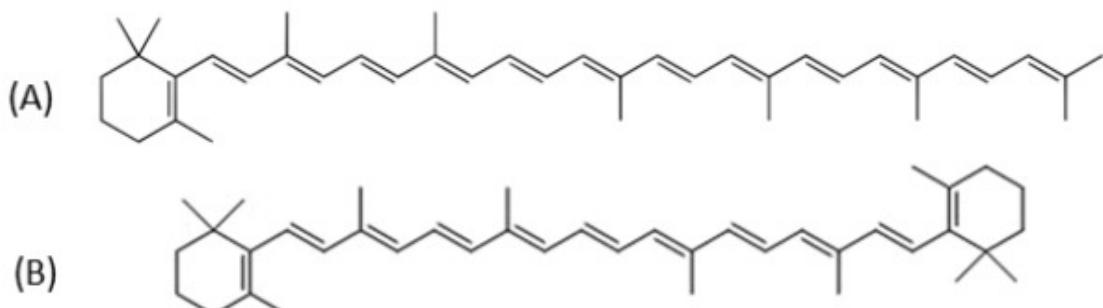
Asperyllone (pigment produced by fungi from genus *Aspergillus*, e.g. *A. awamori*, *A. niger*)



6-[(Z)-2-Carboxyvinyl]-N-gamma Aminobutyric Acid-PP-V (azaphilone pigment, produced by the fungus *Talaromyces albobiverticillius*)



Sclerotiorin (orange azaphilone pigment produced by fungi such as *Penicillium sclerotiorum*)



Torulene (A) and β-carotene (B) (carotenoid pigments, produced by the yeast *Sporidiobolus pararoseus*)



Fig. 10: Chemical structures of fungal pigments with potential coloring properties that could be used in textile dyeing. Reproduced from Venil CK, Velmurugan P, Dufossé L, Renuka Devi P, Veera Ravi A [46].

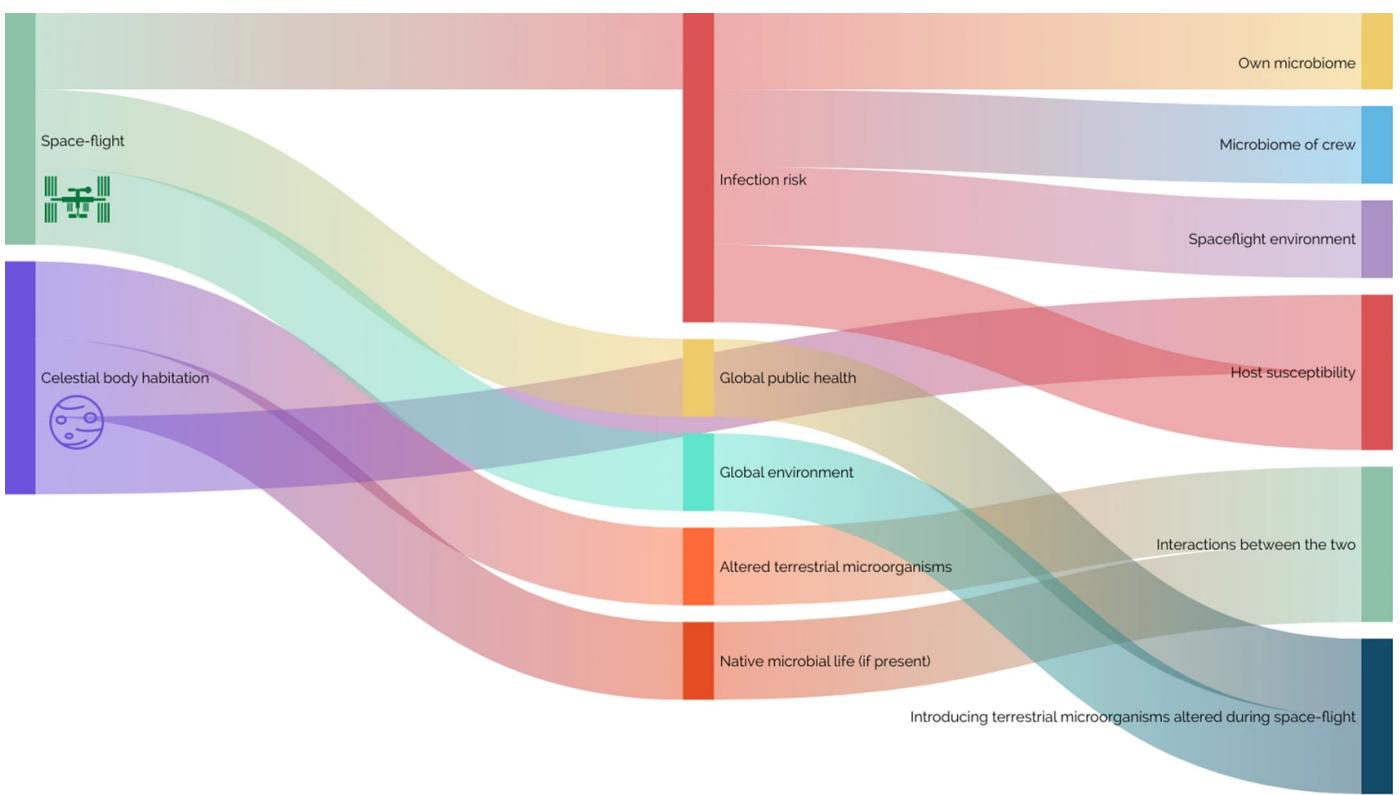


Fig. 11: A sankey diagram visualising the key medical astro-microbiology considerations relating to spaceflight and celestial body habitation. Reproduced from McDonagh et al. [67].

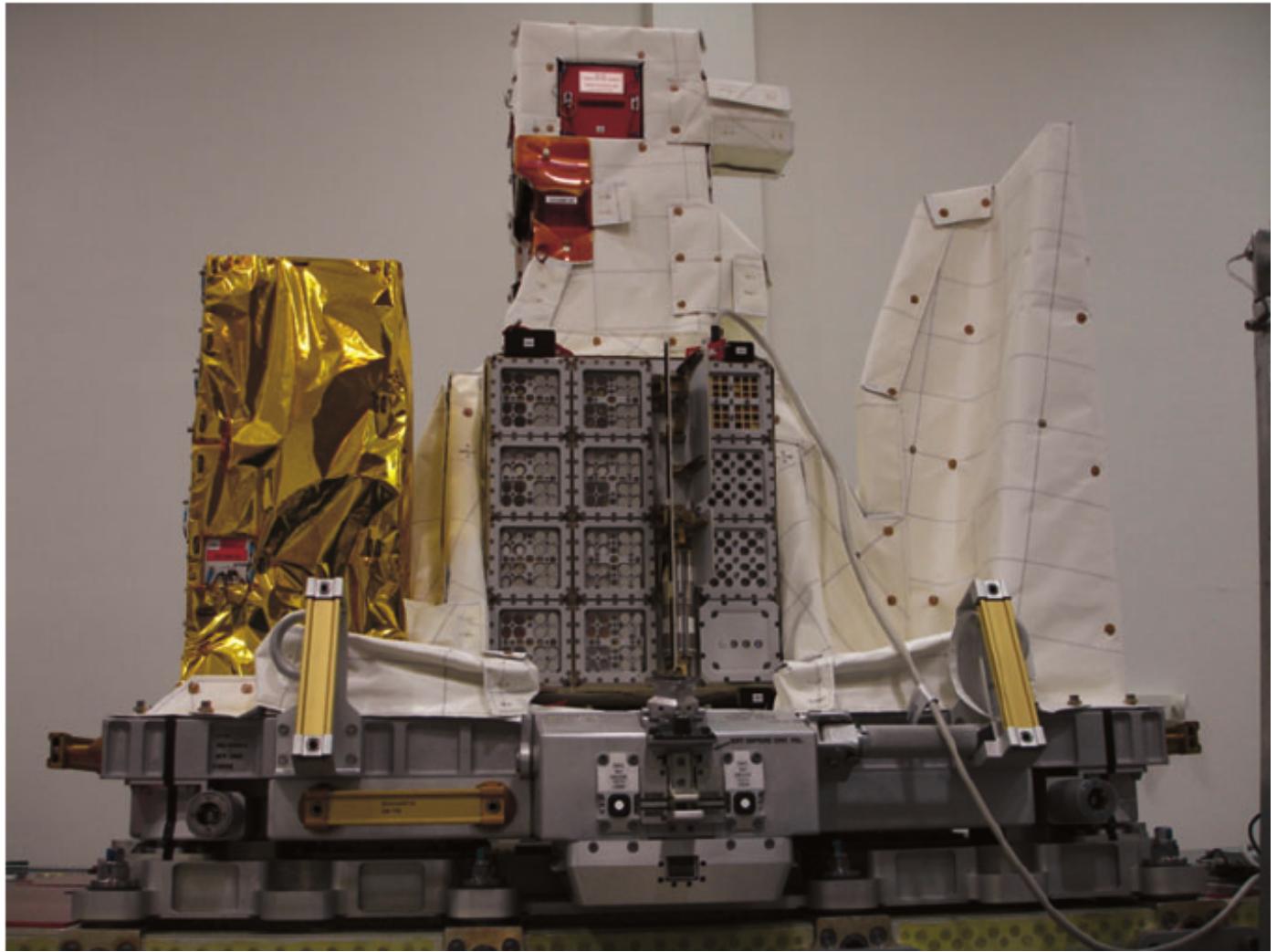


Fig. 12: EXPOSE-E payload, fully integrated and accommodated onto EuTEF at Kennedy Space Center, USA. Arranged vertically from left to right are tray 1 and tray 2—experiments in the four compartments of both trays from bottom compartment to top are ADAPT, PROTECT,  $\frac{1}{2}$  ADAPT, and  $\frac{1}{2}$  PROTECT sharing the third compartment and LIFE in the top compartment; on the right, separated from tray 2 by the three open lids and their motor drives, is tray 3 with R3DE in the bottom compartment, two compartments with PROCESS, and the top compartment with SEEDS. On the right half of each compartment of trays 1 and 2 the mirroring effect of the 0.1% ND filters can be seen. On the left side of EXPOSE-E, the experiment MEDET (wrapped in golden multilayer insulation) is located. Reproduced from Rabbow E et al. [58].

## References

- [1] Black Death | Plague, Death Toll, Definition, Cause, Symptoms, Effects, & Facts | Britannica, <https://www.britannica.com/event/Black-Death> (Nov. 2025).
- [2] M. A. Spyrou, L. Musralina, G. A. Gnechi Ruscone, A. Kocher, P.-G. Borbone, V. I. Khartanovich, A. Buzhilova, L. Djansugurova, K. I. Bos, D. Kühnert, W. Haak, P. Slavin, J. Krause, The source of the Black Death in fourteenth-century central Eurasia, *Nature* 606 (7915) (2022) 718–724. [doi:10.1038/s41586-022-04800-3](https://doi.org/10.1038/s41586-022-04800-3).
- [3] The Black Death and European Expansion, <https://www.history.ox.ac.uk/black-death-and-european-expansion>.
- [4] J. Aberth, From the Brink of the Apocalypse: Confronting Famine, War, Plague and Death in the Later Middle Ages, 2nd Edition, Routledge, London, 2013. [doi:10.4324/9781315881133](https://doi.org/10.4324/9781315881133).
- [5] Decameron Web | Plague, <https://webhelper.brown.edu/decameron/plague/effects/social.php>.
- [6] Labor Shortages Alter Europe's Social Structure | Research Starters | EBSCO Research, <https://www.ebsco.com/research-starters/history/labor-shortages-alter-europe-s-social-structure>.
- [7] J. J. Mark, Religious Responses to the Black Death, World History Encyclopedia (Apr. 2020).
- [8] M. H. Zentner, The Black Death and Its Impact on the Church and Popular Religion.
- [9] R. J. Eisen, A. P. Wilder, S. W. Bearden, J. A. Montenieri, K. L. Gage, Early-Phase Transmission of *Yersinia pestis* by Unblocked *Xenopsylla cheopis* (Siphonaptera: Pulicidae) Is as Efficient as Transmission by Blocked Fleas, *J Med Entomol* 44 (4) (2007) 678–682. [doi:10.1093/jmedent/44.4.678](https://doi.org/10.1093/jmedent/44.4.678).
- [10] B. J. Hinnebusch, D. M. Bland, C. F. Bosio, C. O. Jarrett, Comparative Ability of *Oropsylla montana* and *Xenopsylla cheopis* Fleas to Transmit *Yersinia pestis* by Two Different Mechanisms, *PLOS Neglected Tropical Diseases* 11 (1) (2017) e0005276. [doi:10.1371/journal.pntd.0005276](https://doi.org/10.1371/journal.pntd.0005276).
- [11] D. J. D. Earn, J. Ma, H. Poinar, J. Dushoff, B. M. Bolker, Acceleration of plague outbreaks in the second pandemic, *Proceedings of the National Academy of Sciences* 117 (44) (2020) 27703–27711. [doi:10.1073/pnas.2004904117](https://doi.org/10.1073/pnas.2004904117).
- [12] Plague, <https://www.who.int/news-room/fact-sheets/detail/plague>.
- [13] J. Mackay-Alderson, M. Quastel, E. Wilson, D. Belamby, Factors influencing the re-emergence of plague in Madagascar, *Emerging Topics in Life Sciences* 4 (2020) 423–433. [doi:10.1042/ETLS20200334](https://doi.org/10.1042/ETLS20200334).
- [14] K. B. Patterson, T. Runge, Smallpox and the Native American, *The American Journal of the Medical Sciences* 323 (4) (2002) 216–222. [doi:10.1097/00000441-200204000-00009](https://doi.org/10.1097/00000441-200204000-00009).
- [15] J. S. Marr, J. T. Cathey, New Hypothesis for Cause of Epidemic among Native Americans, *New England, 1616–1619, Emerg Infect Dis* 16 (2) (2010) 281–286. [doi:10.3201/edi1602.090276](https://doi.org/10.3201/edi1602.090276).
- [16] G. Legnaghi de Almeida, C. Santos, The microscopic settler: Smallpox dissemination in the context of european New World explorationEl colono microscópico: Diseminación de la viruela en el contexto de la exploração europea del Nuevo Mundo, *Esboços: histórias em contextos globais* 32 (2025) 1–22. [doi:10.5007/2175-7976.2025.e101292](https://doi.org/10.5007/2175-7976.2025.e101292).
- [17] S. Riedel, Smallpox and biological warfare: A disease revisited, *Proc (Bayl Univ Med Cent)* 18 (1) (2005) 13–20. [doi:10.1080/08998280.2005.11928026](https://doi.org/10.1080/08998280.2005.11928026).
- [18] N. Barquet, P. Domingo, Smallpox: The triumph over the most terrible of the ministers of death, *Ann Intern Med* 127 (8 Pt 1) (1997) 635–642. [doi:10.7326/0003-4819-127-8\\_part\\_1-199710150-00010](https://doi.org/10.7326/0003-4819-127-8_part_1-199710150-00010).
- [19] P. Vaishali, N. Boopathy, Edward Jenner: The Pioneer of Vaccination and His Enduring Legacy in Modern Medicine, *Cureus* 16 (Sep. 2024). [doi:10.7759/cureus.68805](https://doi.org/10.7759/cureus.68805).
- [20] Smallpox, <https://www.who.int/health-topics/smallpox>.
- [21] K. A. Simonsen, J. Snowden, Smallpox, in: StatPearls, StatPearls Publishing, Treasure Island (FL), 2025.
- [22] Louis Pasteur - Microbiology, Germ Theory, Pasteurization | Britannica, <https://www.britannica.com/biography/Louis-Pasteur/Spontaneous-generation> (Nov. 2025).
- [23] Louis Pasteur and germ theory - Medicine in 18th- and 19th-century Britain, c.1700-c.1900 - Edexcel - GCSE History Revision - Edexcel, <https://www.bbc.co.uk/bitesize/guides/zxbqjsg/revision/2>.
- [24] Pasteurization | Definition, Process, Inventor, & Facts | Britannica, <https://www.britannica.com/technology/pasteurization> (Nov. 2025).
- [25] K. A. Smith, Louis Pasteur, the Father of Immunology?, *Front Immunol* 3 (2012) 68. [doi:10.3389/fimmu.2012.00068](https://doi.org/10.3389/fimmu.2012.00068).

- [26] B. .-. A. du texte Repiquet, Méthode de M. Pasteur pour prévenir la rage après morsure, par R. Repiquet,..., 1891.
- [27] R. Rappuoli, 1885, the first rabies vaccination in humans, *Proceedings of the National Academy of Sciences* 111 (34) (2014) 12273–12273. [doi:10.1073/pnas.1414226111](https://doi.org/10.1073/pnas.1414226111).
- [28] R. Gaynes, The Discovery of Penicillin—New Insights After More Than 75 Years of Clinical Use, *Emerg Infect Dis* 23 (5) (2017) 849–853. [doi:10.3201/eid2305.161556](https://doi.org/10.3201/eid2305.161556).
- [29] The discovery of antibiotics.
- [30] S. Y. Tan, Y. Tatsumura, Alexander Fleming (1881–1955): Discoverer of penicillin, *Singapore Med J* 56 (7) (2015) 366–367. [doi:10.11622/smedj.2015105](https://doi.org/10.11622/smedj.2015105).
- [31] Sir Alexander Fleming – Nobel Lecture, <https://www.nobelprize.org/prizes/medicine/1945/fleming/lecture/>.
- [32] J. A. Segre, What does it take to satisfy Koch’s postulates two centuries later? Microbial genomics and Propionibacteria acnes, *J Invest Dermatol* 133 (9) (2013) 2141–2142. [doi:10.1038/jid.2013.260](https://doi.org/10.1038/jid.2013.260).
- [33] L. Walker, H. LeVine, M. Jucker, Koch’s Postulates and Infectious Proteins, *Acta Neuropathol* 112 (1) (2006) 1–4. [doi:10.1007/s00401-006-0072-x](https://doi.org/10.1007/s00401-006-0072-x).
- [34] R. Koch, Die Ätiologie der Milzbrand-Krankheit, begründet auf die Entwicklungsgeschichte des Bacillus Anthracis (Mar. 2010). [doi:10.25646/5064](https://doi.org/10.25646/5064).
- [35] Ueber den augenblicklichen Stand der bakteriologischen Choleradiagnose, <https://zenodo.org/records/2077681>.
- [36] G. K. Gesundheitsamt, Mittheilungen aus dem Kaiserlichen Gesundheitsamte: Bd. Die Aetiologie der Tuberkulose, Norddeutsche Buchdruckerei und Verlaganstalt ; zu beziehen durch Louis Gerschel Verlagsbuchhandlung (G. Goßmann), 1884.
- [37] J. S. Tyagi, The timeless legacy of Robert Koch: Founder of medical microbiology, *Reson* 11 (9) (2006) 20–28. [doi:10.1007/BF02834330](https://doi.org/10.1007/BF02834330).
- [38] L. K. Charkoudian, J. T. Fitzgerald, C. Khosla, A. Champlin, In Living Color: Bacterial Pigments as an Untapped Resource in the Classroom and Beyond, *PLoS Biol* 8 (10) (2010) e1000510. [doi:10.1371/journal.pbio.1000510](https://doi.org/10.1371/journal.pbio.1000510).
- [39] S. Y. Choi, S. Lim, K.-h. Yoon, J. I. Lee, R. J. Mitchell, Biotechnological Activities and Applications of Bacterial Pigments Violacein and Prodigiosin, *J Biol Eng* 15 (2021) 10. [doi:10.1186/s13036-021-00262-9](https://doi.org/10.1186/s13036-021-00262-9).
- [40] H. S. Tuli, P. Chaudhary, V. Beniwal, A. K. Sharma, Microbial pigments as natural color sources: Current trends and future perspectives, *Journal of Food Science and Technology* 52 (8) (2014) 4669. [doi:10.1007/s13197-014-1601-6](https://doi.org/10.1007/s13197-014-1601-6).
- [41] T. Keerthana, S. S. Subhashini, Microbial Pigments: Types, Production Methods, and Diverse Applications in Various Industries: A Review, *IJSAT* 16 (1) (2025) 1593. [doi:10.71097/IJSAT.v16.i1.1593](https://doi.org/10.71097/IJSAT.v16.i1.1593).
- [42] E. Frankel, J. Temple, E. Dikener, M. Berkmen, Bridging the gap with bacterial art, *FEMS Microbiol Lett* 370 (2023) fnad025. [doi:10.1093/femsle/fnad025](https://doi.org/10.1093/femsle/fnad025).
- [43] M. Gomes, G. Soares, A Review of Bacterial Pigments: Harnessing Nature’s Colors for Functional Materials and Dyeing Processes, *Textile & Leather Review* 6 (2023) 167–190. [doi:10.31881/TLR.2023.016](https://doi.org/10.31881/TLR.2023.016).
- [44] L. J. Rather, S. Jameel, S. A. Ganie, K. A. Bhat, Lichen Derived Natural Colorants: History, Extraction, and Applications, in: *Handbook of Renewable Materials for Coloration and Finishing*, John Wiley & Sons, Ltd, 2018, Ch. 6, pp. 103–114. [doi:10.1002/9781119407850.ch6](https://doi.org/10.1002/9781119407850.ch6).
- [45] M. E. Pailliè-Jiménez, P. Stincone, A. Brandelli, Natural Pigments of Microbial Origin, *Front. Sustain. Food Syst.* 4 (Sep. 2020). [doi:10.3389/fsufs.2020.590439](https://doi.org/10.3389/fsufs.2020.590439).
- [46] C. K. Venil, P. Velmurugan, L. Dufossé, P. Renuka Devi, A. Veera Ravi, Fungal Pigments: Potential Coloring Compounds for Wide Ranging Applications in Textile Dyeing, *J Fungi (Basel)* 6 (2) (2020) 68. [doi:10.3390/jof6020068](https://doi.org/10.3390/jof6020068).
- [47] L. Lin, J. Xu, Production of Fungal Pigments: Molecular Processes and Their Applications, *J Fungi (Basel)* 9 (1) (2022) 44. [doi:10.3390/jof9010044](https://doi.org/10.3390/jof9010044).
- [48] B. Basto, N. R. da Silva, J. A. Teixeira, S. C. Silvério, B. Basto, N. R. da Silva, J. A. Teixeira, S. C. Silvério, Production of Natural Pigments by *Penicillium brevicompactum* Using Agro-Industrial Byproducts, *Fermentation* 8 (10) (Oct. 2022). [doi:10.3390/fermentation8100536](https://doi.org/10.3390/fermentation8100536).
- [49] A. Kramar, M. M. Kostic, A. Kramar, M. M. Kostic, Bacterial Secondary Metabolites as Biopigments for Textile Dyeing, *Textiles* 2 (2) (2022) 252–264. [doi:10.3390/textiles2020013](https://doi.org/10.3390/textiles2020013).
- [50] W. Elkhateeb, M. O. Elnahas, G. Daba, Wide Range Applications of Fungal Pigments in Textile Dyeing, in: T. Satyanarayana, S. K. Deshmukh (Eds.), *Fungi and*

- Fungal Products in Human Welfare and Biotechnology, Springer Nature, Singapore, 2023, pp. 289–304. [doi:10.1007/978-981-19-8853-0\\_10](https://doi.org/10.1007/978-981-19-8853-0_10).
- [51] M. Yokoyama, What is Astro Microbiology?, <https://www.news-medical.net/life-sciences/What-is-Astro-Microbiology.aspx> (Oct. 2018).
- [52] G. Horneck, D. M. Klaus, R. L. Mancinelli, Space Microbiology, *Microbiology and Molecular Biology Reviews* 74 (1) (2010) 121–156. [doi:10.1128/mmbr.00016-09](https://doi.org/10.1128/mmbr.00016-09).
- [53] R. Moeller, P. Setlow, G. Reitz, W. L. Nicholson, Roles of small, acid-soluble spore proteins and core water content in survival of *Bacillus subtilis* spores exposed to environmental solar UV radiation, *Appl Environ Microbiol* 75 (16) (2009) 5202–5208. [doi:10.1128/AEM.00789-09](https://doi.org/10.1128/AEM.00789-09).
- [54] G. Gallo, M. Aulitto, Advances in Extremophile Research: Biotechnological Applications through Isolation and Identification Techniques, *Life (Basel)* 14 (9) (2024) 1205. [doi:10.3390/life14091205](https://doi.org/10.3390/life14091205).
- [55] N. Merino, H. S. Aronson, D. P. Bojanova, J. Feyhl-Buska, M. L. Wong, S. Zhang, D. Giovannelli, Living at the Extremes: Extremophiles and the Limits of Life in a Planetary Context, *Front. Microbiol.* 10 (Apr. 2019). [doi:10.3389/fmicb.2019.00780](https://doi.org/10.3389/fmicb.2019.00780).
- [56] B. N. Rekadwad, W.-J. Li, J. M. Gonzalez, R. Pun-chappady Devasya, A. Ananthapadmanabha Bhagwath, R. Urana, K. Parwez, Extremophiles: The species that evolve and survive under hostile conditions, *3 Biotech* 13 (9) (2023) 316. [doi:10.1007/s13205-023-03733-6](https://doi.org/10.1007/s13205-023-03733-6).
- [57] Microbiology | Definition, History, & Microorganisms | Britannica, <https://www.britannica.com/science/microbiology> (Dec. 2025).
- [58] E. Rabbow, P. Rettberg, S. Barczyk, M. Bohmeier, A. Parpart, C. Panitz, G. Horneck, R. von Heise-Rotenburg, T. Hoppenbrouwers, R. Willnecker, P. Baglioni, R. Demets, J. Dettmann, G. Reitz, EXPOSE-E: An ESA Astrobiology Mission 1.5 Years in Space, *Astrobiology* 12 (5) (2012) 374–386. [doi:10.1089/ast.2011.0760](https://doi.org/10.1089/ast.2011.0760).
- [59] M. Wassmann, R. Moeller, E. Rabbow, C. Panitz, G. Horneck, G. Reitz, T. Douki, J. Cadet, H. Stan-Lotter, C. S. Cockell, P. Rettberg, Survival of Spores of the UV-Resistant *Bacillus subtilis* Strain MW01 After Exposure to Low-Earth Orbit and Simulated Martian Conditions: Data from the Space Experiment ADAPT on EXPOSE-E, *Astrobiology* 12 (5) (2012) 498–507. [doi:10.1089/ast.2011.0772](https://doi.org/10.1089/ast.2011.0772).
- [60] C. Pacelli, A. Cassaro, L. Aureli, R. Moeller, A. Fujimori, S. Onofri, The Responses of the Black Fungus *Cryomyces Antarcticus* to High Doses of Accelerated Helium Ions Radiation within Martian Regolith Simulants and Their Relevance for Mars, *Life (Basel)* 10 (8) (2020) 130. [doi:10.3390/life10080130](https://doi.org/10.3390/life10080130).
- [61] S. Onofri, J.-P. de Vera, L. Zucconi, L. Selbmann, G. Scalzi, K. J. Venkateswaran, E. Rabbow, R. de la Torre, G. Horneck, Survival of Antarctic Cryptodolitic Fungi in Simulated Martian Conditions On Board the International Space Station, *Astrobiology* 15 (12) (2015) 1052–1059. [doi:10.1089/ast.2015.1324](https://doi.org/10.1089/ast.2015.1324).
- [62] Panspermia | Research Starters | EBSCO Research, <https://www.ebsco.com>.
- [63] P. S. Wesson, Panspermia, Past and Present: Astrophysical and Biophysical Conditions for the Dissemination of Life in Space, *Space Sci Rev* 156 (1-4) (2010) 239–252. [arXiv:1011.0101](https://arxiv.org/abs/1011.0101), [doi:10.1007/s11214-010-9671-x](https://doi.org/10.1007/s11214-010-9671-x).
- [64] Pasteur-Theorie für Lebensbeginn: Stock-Vektorgrafik (Lizenzfrei) 1182778669, <https://www.shutterstock.com/de/image-vector/pasteur-theory-life-beginning-1182778669>.
- [65] J. Madhusoodanan, Science and Culture: Petri palettes create microbial masterpieces, *Proc Natl Acad Sci U S A* 113 (40) (2016) 11056–11058. [doi:10.1073/pnas.1610867113](https://doi.org/10.1073/pnas.1610867113).
- [66] Y. Caro, M. Venkatachalam, J. Lebeau, M. Fouillaud, L. Dufossé, Pigments and Colorants from Filamentous Fungi, in: *Fungal Metabolites*, Springer, Cham, 2015, pp. 1–70. [doi:10.1007/978-3-319-19456-1\\_26-1](https://doi.org/10.1007/978-3-319-19456-1_26-1).
- [67] F. McDonagh, M. Cormican, D. Morris, L. Burke, N. K. Singh, K. Venkateswaran, G. Miliotis, Medical Astro-Microbiology: Current Role and Future Challenges, *J Indian Inst Sci* 103 (3) (2023) 771–796. [doi:10.1007/s41745-023-00360-1](https://doi.org/10.1007/s41745-023-00360-1).