OPINION ARTICLE



REVISED The evolution of universal adaptations of life is driven

by universal properties of matter: energy, entropy, and

interaction [version 3; peer review: 3 approved]

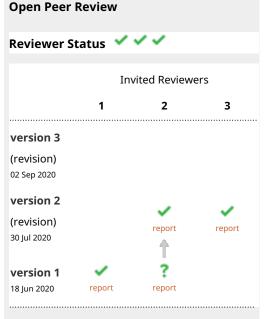
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Abstract

The evolution of multicellular eukaryotes expresses two sorts of adaptations: local adaptations like fur or feathers, which characterize species in particular environments, and universal adaptations like microbiomes or sexual reproduction, which characterize most multicellulars in any environment. We reason that the mechanisms driving the universal adaptations of multicellulars should themselves be universal, and propose a mechanism based on properties of matter and systems: energy, entropy, and interaction. Energy from the sun, earth and beyond creates new arrangements and interactions. Metabolic networks channel some of this energy to form cooperating, interactive arrangements. *Entropy*, used here as a term for all forces that dismantle ordered structures (rather than as a physical quantity), acts as a selective force. Entropy selects for arrangements that resist it long enough to replicate, and dismantles those that do not. Interactions, energy-charged and dynamic, restrain entropy and enable survival and propagation of integrated living systems. This fosters *survival-of-the-fitted* – those entities that resist entropic destruction - and not only of the fittest - the entities with the greatest reproductive success. The "unit" of evolution is not a discrete entity, such as a gene, individual, or species; what evolves are collections of related interactions at multiple scales. Survival-of-the-fitted explains universal adaptations, including resident microbiomes, sexual reproduction, continuous diversification, programmed turnover, seemingly wasteful phenotypes, altruism, co-evolving environmental niches, and advancing complexity. Indeed survival-of-the-fittest may be a particular case of the survival-of-the-fitted mechanism, promoting local adaptations that express reproductive advantages in addition to resisting entropy. Survival-of-the-fitted accounts for phenomena that have been attributed to neutral evolution: in the face of entropy, there is no neutrality; all variations are challenged by



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Any reports and responses or comments on the article can be found at the end of the article.

ubiquitous energy and entropy, retaining those that are "fit enough". We propose experiments to test predictions of the survival-of-thefitted theory, and discuss implications for the wellbeing of humans and the biosphere.

Keywords

Evolution, Energy, Entropy, Interaction, Cooperation, Niche, Fitness, Fittedness

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REVISED Amendments from Version 2

Summary of changes V2-to-V3

These amendments address comments in Susan Stepney's review. See the review and the authors' response for more details.

1. In the introduction, we stated that we plan to address the mechanism of unicellular evolution in a future paper.

2. In the section "Entropy is life's nemesis and evolution's facilitator", we edited our statement of the second law of thermodynamics, and added published recitations of two other brief statements of the law.

3. In the above section, we added to the list of papers exploring the connections between evolution and entropy citations of "Investigations" by Kauffman and "Into the cool: Energy flow, thermodynamics, and life" by Schneider and Sagan.

4. In the section "Interactions", we reordered and rephrased the sentences to clarify and add context to Feynman's quote that "everything is made of atoms".

5. In the section "programmed renewal," we cited the work of Maturana and Varela on autopoiesis and the definition of living systems.

6. In the section "Metabolism protects against entropic dissolution," we added citations to Kauffman's discussion of the constrained release of energy.

7. In the above section, we clarified that bonding releases entropy into the environment as heat.

8. In the section "Humans are responsible for widespread niche deconstruction", in the discussion of biomass comparisons, we explicitly included humans.

9. In the section "Analysis and modeling," we cited Kauffman's modeling approaches appearing in Investigations and in Answering Schroedinger's "What is Life" and compared them with our approach.

10. In the section "Predictions" we cited Lenski's long running evolution experiment.

11. We corrected several typos.

Any further responses from the reviewers can be found at the end of the article

Introduction

Evolution may be defined as the transitions over time of living systems, which include cells, organisms, species and ecosystems, along with their component genes and other molecules. Evolution can take place when several conditions are fulfilled: *variations* among the evolving systems; *interactions* with the environment, both the organic and the inorganic; *destruction* of some variants and continuing *propagation* of others; and mechanisms that can *select* between the two fates. Any mechanisms proposed to drive the processes of selection and heritability would also have to explain, or at least not contradict, the reality of presently living systems; observed life constitutes a reality test for theories of evolution.

The generally accepted mechanism of evolution is *Natural Selection* resulting from *survival-of-the-fittest*¹. The neo-Darwinian theory of evolution has expanded Darwin's concept of Natural

Selection to include modern genetics. Natural Selection proposes that variant individuals must struggle for resources in a necessarily limited environment; the individual who manifests superior adaptive fitness to its environment wins out in the struggle to survive and procreate. The winner's offspring inherit the winner's genes and so enrich the species with increased frequencies of the fittest genes. Fitness, or "being fittest" is equated with reproductive success². Iteration of this process over generations eventually leads to an optimally adapted genotype for the species in the given environment^{3,4}. Thus, species would be expected to evolve an optimum uniformity in a stable environment. Survival-of-the-fittest selection in a new environment generates a new species and restarts the optimization process.

However, the rationale behind the theory of Natural Selection has been called into question by the recent discovery that all multicellular organisms—plants, invertebrates and vertebrates—are holobionts composed of gene products of the eukaryote organism in cooperative interaction with great numbers of resident microbiome prokaryote genes and organisms⁵. In a word, there are no individual organisms to engage in classical Survival-of-the-Fittest struggle^{6–8}. All multi-cellular organisms are groups.

Indeed, microbiome organisms and their genes are acquired somatically after birth and are independent of host eukaryote genome reproduction to a significant degree. Consequently, the inheritance of eukaryote genes alone cannot endow holobiont offspring with the phenotypic fitness of a progenitor holobiont. For example, a mouse that has no microbiome, a germ-free mouse, develops an abnormal gut, a crippled immune system and other abnormalities⁹. If there be no genetically independent, procreating individuals, individual eukaryote reproductive success alone cannot be responsible for genetic progress.

Recent publications have discussed the inheritance of the microbiome¹⁰ and controversy surrounds the definition of the holobiont: for example, whether the holobiont is an integrated individual or a community^{6,11}. These matters raise questions critical to our understanding of the holobiont's role in the process of evolution. The issue has yet to be resolved.

Another recent controversy surrounding the mechanism of evolution relates to the concept of niche construction – the co-evolution of the environmental niche with its resident species¹².

The theory of *Extended Evolutionary Synthesis* (EES)¹³⁻¹⁵ reconsiders evolution in the light of niche construction, the holobiont and other challenges facing classical thinking. The issue is subject to debate¹⁶.

We shall not attempt to analyze or resolve these controversies; rather we draw a distinction between universal adaptations expressed by essentially all multicellular organisms and local adaptations expressed by particular species in a defined environment. This paper deals with universal adaptations of multicellulars observed in most all environments including cooperation, sexual reproduction, continuous diversification, programmed turnover and others, analyzed below. We reason that universal adaptations must have been driven and selected for survival by ubiquitous physical features of matter: *Energy*, *Entropy* and *Interaction*.

We shall first describe energy and entropy and then proceed to interaction and how their integration can account for universal adaptations. Along the way, we contrast our proposed material mechanism with the classical mechanism of survival-of-the-fitted. We do not deal directly with the evolution of unicellular eukaryotes or prokaryotes; we plan to address the mechanism of unicellular evolution in a future paper.

This paper has evolved from preliminary thoughts about the adaptive immune system¹⁷, the role of information in art, science and evolution¹⁸ and the evolutionary relationship between information and entropy¹⁹.

Energy and entropy Energy is an imperative of life

Darwin did not refer to energy as a formative factor in evolution, and neo-Darwinian gene-centered discourse has paid little attention to either energy or entropy. In contrast, we shall show that energy and entropy (and networks of cooperative interaction) are more consequential to the evolution of universals than is the selfish Darwinian struggle.

Energy is defined by physicists in various ways using different mathematical formulations, each formulation suitable for a different situation²⁰. Nevertheless, we can discuss the evolutionary role of energy without having to get into the mathematical details. Dictionaries define energy as the impetus behind all motion and all activity and as the capacity to do work²¹. Capacity to do work relates to energy in an organized or ordered form. Living systems, as we discuss below, exploit and are also damaged by energy, both ordered and disordered. The world is permeated with energy in many forms: irradiation from the

sun, the earth and outer space; the force of gravity; impacts with asteroids; chemical interactions; electromagnetism; the weather—wind, rain, snow, hailstones, cold, heat; lightning; tides, rivers, floods, and droughts; fires; seismic movements and eruptions; to these inorganic elements we can add energy originating from living systems and from human technology.

Entropy is life's nemesis and evolution's facilitator

The word entropy is formally defined as a measure of the number of possible microstates of a system²². The term is informally used to refer to disorder *per se*, to a measurement of the amount of disorder, and to the tendency, processes or forces that spontaneously effect deterioration, destruction, or dissolution²³. To avoid ambiguity²⁴, we use the term to include all the circumstances and events that can lead to the destruction or dissolution of any specific arrangement.

The concept of entropy derives from the second law of thermodynamics, which states that that disorder in an isolated system will continuously increase spontaneously until random homogeneity is reached; more specifically, quoting from Wehrl²², "the entropy of a closed system never decreases; it can only remain constant or increase"; or, stated as the maximum entropy principle, "the entropy of a closed system in equilibrium always takes the maximal possible value". Entropy, as we said above, is formally defined as a statistical property of any system that contains many component parts or states - so called microstates: the greater the number of parts or micro-states, the less likely will be any particular, given arrangement; randomness will prevail (Figure 1). Moreover, the internal energy inherent in all substances²⁵ - seen, for example in Brownian motion - propels them, sooner or later, to fall apart-unless, as we discuss below, cooperative interactions and other processes hold that arrangement together or renew it.

Living systems depend on the specific arrangements and interactions of molecules, cells, organs, individuals, species, and ecosystems²⁶. Consequently, life has had to adapt to entropy,

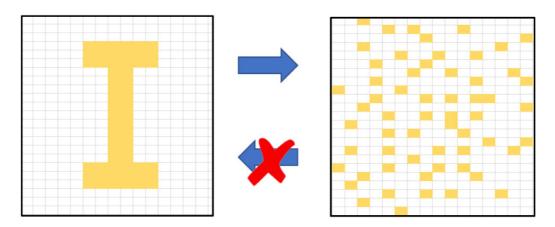


Figure 1. Entropy; the spontaneous dissolution of order into randomness. In this simplified conceptual representation, the border rectangles mark a closed system. The "I" structure in the left frame will spontaneously dissolve into a random disorder of its component pixels, shown in the right frame; the spontaneous process is irreversible. The challenge facing life is to resist entropic dissolution of essential structure. The figure was inspired by P. W. Atkins²⁷.

the metaphoric destroyer of specific arrangements. Ludwig Boltzmann, one of the founding fathers of thermodynamics, already made the point in 1875²⁸:

• The general struggle for existence of animate beings is not a struggle for raw materials – these, for organisms, are air, water and soil, all abundantly available – nor for energy which exists in plenty in any body in the form of heat, but a struggle for [negative] entropy, which becomes available through the transition of energy from the hot sun to the cold earth.

"Negative entropy" refers to "order".

In his book *What is Life*, the physicist Erwin Schrödinger concluded that entropy—the decay of orderly arrangements into atomic chaos—is the ultimate challenge facing living systems²⁹. Schrödinger did not attempt to relate evolution to entropy, and accepted Darwinian Natural Selection and survival-of-the-fittest as a fact (page 11). He made it clear, however, that living organisms have to pay for their order by exporting into the environment a due measure of entropy in the form of heat.

The mystery of life, says Schrödinger, is that "existing order displays the power of maintaining itself and of producing orderly events." Schrödinger offers no mechanism that might maintain and produce order by restraining entropy, other than to state that "we, no doubt, draw on experience concerning social organization and other events which involve the activity of organisms. And so it might seem that something like a vicious circle is implied." In other words, Schrödinger speculates that the orderly arrangements of life emerge from the order inherent in life's ongoing activities—in a "vicious circle"; cooperative, "social" interactions somehow restrain the effects of entropy.

Despite the physical definition of entropy as a statistical measure of microstates, physicists Boltzmann and Schrödinger both relate to entropy as a concrete threat to life.

It is clear that entropy is life's nemesis, but how can entropy also function as evolution's facilitator? Above, we pointed out that evolution requires *variation*, *interaction*, *selection*, *destruction and propagation*. Obviously, entropy plays important roles in variation and destruction, but, as we shall see, entropy will also turn out to be a major factor in selection.

Over the years, many theoretical papers have been written to reconcile the evolution of life with entropy and the second law of thermodynamics (for example, see 30-34); but, as far as we know, none of these authors have challenged the paradigmatic neo-Darwinian mechanism of survival-of-the-fittest, and none of them have invoked entropy as a selective factor in evolution, as we do here.

Prevalent adaptations have evolved to mitigate the effects of entropy

Adaptation is often described as the aim of evolution³⁵: adaptation designates a state in which an organism or species fits the

constraints and demands of its environment; to adapt is to solve a vital survival problem imposed by the environment. This concept of adaptation, however, is paradoxical; for example, wings are local adaptations for flying and fins for swimming. Local adaptations are limited to a species in a certain environment; wings and fins only solve fitness problems if one is already a bird or a fish; an elephant would only have more problems surviving with wings or fins.

Levins and Lewontin state that; *It is difficult to think of any physical force or universal physical law that represents a fixed problem to which all organisms must find a direct solution*³⁶. If you think about it, however, there is one universal physical law to which adaptation is required *a priori* for living – all creatures in any environment have to deal with entropy. Therefore, adaptations that restrain entropy are ubiquitous in living systems; extending Schrödinger, we propose that cooperative, "social" interaction is one such adaptation.

Interactions

Interactions fashion reality

The physicist Richard Feynman referring to the atomic and subatomic scales, claimed that "all mass is interaction"³⁷ (Prologue); similarly, Carlo Rovelli stated that all reality is interaction³⁸. Feynman suggested a sentence that bears the most information about our world for posterity: "all things are made of atoms—little particles that move around in perpetual motion, attracting each other when they are a little distance apart, but repelling upon being squeezed into one another"²⁵. Since most matter in the biosphere is made of atoms, everything made of such matter is made of moving, interacting atoms. Here we apply dynamic interaction as a building block at higher, biological scales represented by molecules, cells, organisms, species and ecosystems.

Cooperative interactions are pervasive and central to life

We define an interaction as a relationship between two or more entities involving a transfer or exchange of matter, information and/or energy. Interactions include both struggle and cooperation: in a struggle, the participants each strive to win and dominate the others – who become the losers. In a cooperative interaction, there are no losers; the participants each gain some benefit, or at the least suffer no loss.

Living systems interact with other living systems, but they also interact with non-living matter like water, minerals and air. Interactions maintain life, and are critical to the evolution of life. Darwin proposed that evolution is driven by interactions expressed as individual struggles for survival. In a limited environment, one's gain must be offset by others' losses. Such a zero-sum game logically generates selfishness^{39–41}. When, however, we look at the dominant characteristics of living systems, what we see is not only selfishness but also interactions that generate cooperation and symbiosis.

Symbiotic cooperation—living and working together—seems to have been decisive in the evolution of complex life: the primordial development of a progenitor eukaryote cell, estimated to have occurred about two billion years ago, is now thought to have resulted from the symbiotic amalgamation of two or more prokaryote (bacteria or archaea) cell types to eventually evolve a new living entity – a eukaryote cell^{42,43}. Prokaryote life was then, and still is consummately fit; the foray of life into a complicated eukaryotic endosymbiosis is a fortuitous meander of nature that has eventually led to us complex humans and to our effects on the biosphere.

The evolution of single-celled eukaryotes into multicellular organisms is also a cooperative enterprise. The phenotype of any multi-cellular individual is the expression of at least two internal cooperative processes: First, there is the developmental and physiological cooperation of the offspring's eukaryote genes and cells, which initially emerged from the endosymbiosis of parental germ cells. One's eukaryote cells share a common DNA genome, but differential epigenetic expressions of this shared genome are key to interactions resulting in growth, development, differentiation, immunity, repair, reproduction, physiology, behavior, metabolism, species and ecosystems. Indeed, all processes of multi-cellular life involve cooperative interactions: all depend on a variety of cooperative interactions between molecules, cells and organisms. Even the development of a mutated, lethal tumor can progress only through cooperative interactions of the tumor cells with a microenvironment of non-tumor cells and tissues, including angiogenesis and immune system complicity⁴⁴.

Cooperation includes the microbiome

The second level of eukaryote cooperative interaction is with a resident microbiome, which may include bacteria, archaea, yeasts, algae and viruses⁴⁵. Research is yet in an early stage, but the microbiome is clearly an essential component of all multi-cellular invertebrate, vertebrate and plant organisms, which, as we mentioned, are holobionts. The gut microbiome in humans and other organisms performs a variety of significant functions: metabolism of otherwise indigestible foodstuffs; production of vitamins and other essential molecules; detoxification of harmful substances; neutralizing or blocking pathogens; assistance in developing the immune system; warming the body; influencing brain development and behavior; and other benefits8. The human species may actually benefit from the ridding of burdensome, aged humans by the timely evolution of "pathogenic" microbiome bacteria⁴⁶. In general, microbiome-host metabolic interactions are critical in human health and disease; an abnormal microbiome (dysbiosis) is a hazard47. The microbiome dynamically changes with age, diet, gender, and other factors; for example, a change in diet can greatly expand or reduce particular components of a holobiont's microbiome and, consequently, affect the holobiont's health, adaptation to environmental changes and survival.

How the essential microbiome and the host organism genome are transmitted faithfully across generations is still a matter for debate¹⁰. As part of this discussion, Stencel and Wloch-Salamon propose that the holobiont is not inherited as a single entity, but is a cooperating group¹¹. Hence, the discovery of the holobiont obliges evolutionary theory to reconsider group selection – a controversial idea that is incompatible with the concept of

evolution driven by the heritable fitness of a single-genome individual 48 .

Social bonding is encoded in species genomes

Biology has recently discovered that all species of vertebrates bear genes encoding molecules like oxytocin that function to enhance cooperation, social bonding, and love, and activate brain reward centers while reducing conflict⁴⁹; evolution for many millions of years seems to have generated mechanisms fostering cooperation and mutual benefit. Indeed, oxytocinlike genes are borne by many invertebrates⁵⁰; and the products of microbiome bacteria stimulate oxytocin production in their cooperating host⁵¹.

Social bonding and empathy would also appear to be programmed in the systems of mirror neurons discovered in the brains of humans and other primates⁵²; these neuronal centers are active in imitation and empathy for others⁵³. Mirror neurons probably evolved even before the primates⁵⁴.

Of course, mutual identification and cohesion within a group may also give rise to prejudices and to episodic conflicts with others outside the group⁵⁵. In any case, it is clear that evolution has generated species outfitted with innate aptitudes for bonding, cooperation, empathy and social interaction. Even the evolution of the dog from the wolf appears to have involved mutual oxytocin-related bonding with humans⁵⁶.

Horizontal gene transfer is a form of cooperation

The transfer of genetic material between existing species can be seen as a further example of non-Darwinian cooperation; a species receives genes from organisms of other species that have evolved in a different environment. Horizontal gene transfer occurs regularly among prokaryotes, but also has occurred in eukaryote evolution. A notable example is the development of the placenta, a formative adaptation in mammalian evolution; placenta function involves the syncytin gene, believed to have been transferred horizontally from a virus⁵⁷. The survival-of-the-fittest theory, which is based on the evolution of genes internal to a single species, cannot easily account for the roles in evolution of horizontal gene transfer^{58,59}.

Cooperation retards entropy

Figure 2 illustrates that cooperative interactions between the components of an arrangement can delay the entropic destruction of the arrangement.

Darwin was aware of the intrinsic conflict between selfish struggle and cooperation¹ (Chapter 7); later evolutionary thinkers have attempted to explain cooperation as a result of an underlying selfish drive by theories of kinship selection or reciprocation agreements (see summary in 60), including the application of mathematical game-theory explanations^{61–63}.

Maynard Smith and Szathmary have tabulated and studied the concept of Major Transitions in Evolution^{64,65}. Like us, they note the importance in evolution of cooperation and sexual

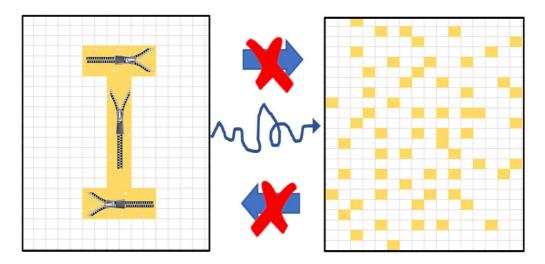


Figure 2. Cooperation retards entropy. Compare this conceptual figure to Figure 1. Here, the pixels that form the "I" structure on the left are held together by cooperative interactions, pictured as zippers that link the pixels. These interactions slow down the deterioration of order and structure, depicted by the long and tortuous arrow from the left to the right frame, while both short arrows are blocked.

reproduction. However, in contrast to our entropic selection theory, Maynard Smith and Szathmary attempt to explain these and all other features of life as resulting from Darwinian competition and survival of the fittest.

Wolf, Katznelson and Koonin have identified competing interaction and "frustrated states" as a formative element in evolution and study how competition between interactions on different scales generates cooperation⁶⁶.

Darwinian game theory explanations for the evolution of cooperation have been widely accepted, but we believe that cooperation as a universal adaptation is more easily understandable once we hedge selfish struggle and include entropy as a selective agent in the process of evolution. An empirical example of the ability of cooperation to restrain entropic degradation is provided by the DNA molecule.

Cooperative interactions prolong the survival of DNA

DNA is well known as the repository of genetic information shared by all forms of eukaryote life. Compare the fragility of single-stranded DNA (ssDNA) with the unparalleled stability of double-stranded DNA (dsDNA)^{67,68}. The paired strands of dsDNA, bound together, are relatively resistant to random damage because each strand is in constant interaction with its sister strand; the two strands can be perceived to be in continuous cooperation; essentially all potentially reactive, non-covalent bonding energy is engaged in cooperative interactions between the strands. In marked contrast to the stable dsDNA dimer, ssDNA and single stranded RNA (ssRNA) are highly susceptible to destruction; reactive bonding energy in the single-strand configuration is not quenched by physiological interactions and is available for haphazard interactions with illicit target

molecules, leading to degradation (Figure 3). Cooperative interaction explains why stable dsDNA can be recovered from mummies or woolly mammoths thousands of years old⁶⁹, while unstable ssDNA and ssRNA survive for only days, at best. dsDNA illustrates how an exchange of energy between the interacting strands retards their entropic dissolution. dsDNA models the universal power of interaction to forestall entropy. We shall cite a higher scale molecular example in the section *Metabolism* below, when we discuss how dynamic metabolic interactions establish networks that restrain entropy.

Interaction also maintains life at the macroscopic scale

We have all observed the role of cooperative interactions in maintaining life at the macroscopic scale: physical isolation or enforced bed rest of an elderly person in a hospital can lead to sudden collapse and death⁷⁰. Why did hospitalization kill the grandmother with the broken hip? She was receiving all the material resources needed for physical survival-food, fluids, oxygen, nursing care; what did she lack? It was not only the concentration of lethal bacteria and other such medical hazards in hospitals that killed the patient; deprivation of her customary daily activities and interactions with her familiar physical and human environments made her more susceptible to undesirable interactions⁷¹. There are many observations that social interaction itself is critical. Retirement or social isolation accelerates mental and physical decline72; newborns isolated from maternal interactions can sicken and die73. In contrast, relationships and responses to challenge delay entropic decline and rejuvenate and maintain body and soul. Life thrives on interaction⁷⁴. As they say; use it or lose it.

The function of interaction in maintaining life may help us understand why some species of birds, and other organisms, have

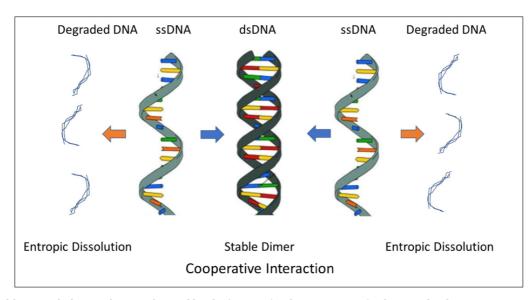


Figure 3. Double-stranded DNA (dsDNA), formed by the interaction between two single strands of DNA (ssDNA), resists entropy and thus enables long term survival of the composite genetic information. The colored bars protruding horizontally from the DNA strands represent non-covalent bonding sites. Interactions between the bonding sites in dsDNA protect the joint molecule from illicit, potentially damaging interactions. DNA in the single stranded form—ssDNA—is not protected from non-physiological interactions and is thereby readily degraded. Image of ssDNA and dsDNA reproduced from 75 under fair use licensing for nonprofit and educational purposes.

evolved to "waste" resources in elaborate courtship displays and bowers and to indiscriminately feed parasitic chicks (which may be obscenely large) of other bird species⁷⁶. Such behaviors clearly contradict individual Darwinian fitness. We propose, however, that seemingly wasteful activities and interactions in themselves maintain cooperative fittedness; as long as a holobiont organism or group has enough matter and energy to act, evolution will not select against a fitted action that is merely inefficient or blatantly wasteful⁷⁷. Interaction itself restrains entropy. Indeed, as we discuss below, living systems shield themselves from entropy, not by saving energy, but by deploying networks of energy to maintain interactions. Cooperation is its own reward.

Prokaryotes too have evolved cooperative interactions

Microbiologists, until recently, have studied prokaryotes isolated in "pure" cultures in the laboratory; in nature, however, prokaryotes, like eukaryotes, live in complex communities of cooperating organisms in microbiomes, in biofilms, in soil, in water, and in other collective environments⁷⁸. Mutual group behaviors are adjusted by the exchange of molecular signals and genetic elements between prokaryotes and, in microbiomes, also between prokaryote and eukaryote cells⁷⁹. Cooperative interactions support the prokaryote world too.

Metabolism

Metabolism is essential to life

Metabolism refers to chemical interactions that produce and transform energy. The term is usually applied to the breakdown or synthesis of molecules that release, consume, or store energy. Life has evolved metabolic arrangements that organize energy: photosynthesis and other contrivances trap disorganized light, heat and chemical energy released by our exploding sun⁸⁰ and our quaking earth⁸¹; living systems, through metabolic networks, transform these potentially destructive forces into organized molecular and cellular structures, processes and networks of interactions that build and maintain life. Living systems also exploit energy and matter, both ordered and disordered, obtained from other living systems – creating food chains and ecosystems⁸². The trapping of available energy, both disordered and ordered, and its conversion into networks of orderly arrangements and interactions, maintain life and its renewal in the face of entropy.

The subject of metabolism was thought to have been solved decades ago with the discoveries of the functions of vitamins and the major cycles, like the Krebs cycle, that control energy transformations. But metabolism has returned lately as a major subject of research. Metabolic pathways can differ in the substances used as fuels, the molecules consumed or synthesized, the activated enzymes, the involvement of aerobic or anaerobic pathways, the efficiency of the processes, and more. The details are beyond our present scope, but dynamic shifts in metabolic processes have been discovered not only to fuel life, but to actually signal and regulate key cellular activities and interactions. Metabolism regulates stem cell differentiation⁸³, immune cell functions⁸⁴, microbiome interactions⁸⁵, brain cell signaling⁸⁶, and the growth and development of cancer cells⁸⁷, among other functions. It is no wonder that oxytocin, the hormone of cooperation and bonding, also affects metabolism⁴⁹.

Metabolism protects against entropic dissolution

The flows of energy through the networks of a living system create a unified structure that resists entropic dissolution. Building a house of cards exemplifies the power of interactive energy: a single card will not stand on a tabletop for more than an instant; two cards interacting as a triangle can stand appreciably longer-the electrostatic interactions between the cards and their balanced support keep them up, and you can progress to build a stable house of cards. That is, until you open the window and the winds of entropy blow it down. If, however, you want your house of cards to survive casual entropic winds, you only have to add a drop of glue to the interfaces between the cards. The glue augments the electrostatic bonding interactions between the cards and the house of cards will hold up, at least until the dog demolishes it. Thus do ongoing metabolic interactions (largely electrostatic) between living systems create a jointly unified structure that can restrain destructive effects of entropy.

Moreover, metabolic networks deal with energy in a step-by-step fashion—one reaction at a time; Stuart Kauffman has pointed out that such a "constrained release of energy" operates to actually reduce the entropy within the metabolic network while it increases the amount of effective work performed by the system^{33,88}. Metabolism, besides energizing the cooperative interactions of life, acts by itself as an entropy-restraining adaptation.

An ongoing interaction between two agents reduces the numbers of random microstates — and thus of the entropy— otherwise present in each of the agents when they are not interacting. For example, as stated in 89:

• "...A spark applied to the mixture [of hydrogen and oxygen] initiates a chemical reaction in which hydrogen and oxygen combine to form water. If the temperature of the system is held constant, the entropy of the system decreases because 3 moles of two differing reactants have been combined to form 2 moles of a single product. The gas now consists of a uniform set of indistinguishable molecules".

Bonding between reacting entities reduces their combined microstates and thus their total amount of entropy; the new combined entity releases entropy into the surroundings as heat.

The reduction of internal entropy achieved by integrated energy networks accounts for the evolution and persistence of interactions that would appear to detract from zero-sum Darwinian fitness. In the eyes of Natural Selection, any arrangement that efficiently saves energy should win in a struggle with arrangements that wastefully squander resources. Nevertheless, the biosphere abounds with inefficient interactions⁷⁷: people waste time and money on trinkets; birds, humans and other suitors invest energy on displays and rituals; butterflies, fish, birds and mammals undertake long migrations filled with peril; many species survive with energy-draining parasitisms. Only some of these "wasteful" arrangements can be explained by variants of a

handicap principle⁹⁰, an argument that has its critics (see summary in 91).

Clearly, one can imagine mutations that could do away with such extravagances. But living systems regularly defy the logic of exclusively Darwinian bookkeeping. Cooperative interactions, like those that connect the two strands of dsDNA or a glued house of cards, can form a hedge against entropy, irrespective of efficiency or wastefulness. In other words, interaction itself resists entropy, as suggested by Schrödinger, and so Fittedness can account for arrangements and interactions that make little sense to survival-of-the-fittest thinking. Metabolism fuels the "preservative social" activities of life envisioned by Schrödinger to restrain entropy²⁹.

Programmed renewal

Lifespans, rates of birth, death and reproduction are programmed in species

Survival requires rebirth. The inevitable dissolution of structure implies that all livings beings are destined, in time, to fall apart and die (Figure 1). Life has had no alternative but to effect its own renewal. In fact, we might define a living entity as one that is able to exploit energy, by way of metabolism, to renew itself⁹².

Darwin founded Natural Selection on the assumption that living organisms strive to survive and reproduce as long and as much as they can¹. The fittest individuals, according to classical survival-of-the-fittest thinking, should have the most offspring. After a century and a half of refinements, qualifications, and other added nuances, the underlying principle of presentday evolutionary theory is that "*Natural selection is all about variation in reproductive success*" and "*Natural selection is not 'survival of the fittest,' but rather 'reproduction of the fittest.*². The fittest should live the longest, provided that longevity does not compromise reproduction^{2,93}.

But when we take a look at life as lived, we see that lifespans and birthrates are essentially programmed by an organism's species more than by an organism's success in struggles with competing organisms^{5,94}. Each species has evolved its genetic endowment to encode a relatively standard lifespan for its member organisms – whether for days, seasons, or years. The apportionment of survival time, barring accidents, is fairly fixed⁹³. Classical explanations based on Natural Selection attribute fixed lifespans to a delicately balanced optimization of factors, such as age of sexual maturity, numbers of offspring, and reproductive cycle timing. But superior or successful individuals do not necessarily live the longest or beget the most offspring.

Indeed, a mindset focusing on biological survival may have delayed the recognition of programmed death⁹⁵ until apoptosis^{96–99} and autophagy¹⁰⁰ became important subjects of research.

Similar to lifespan, the time of birth and replication is closely linked to the style of life evolved by the species as a whole; the time of birth of a new organism, how long the organism lives and its numbers of offspring are associated with the way the species has evolved to make its living and manage its reproductive energy. Humans and elephants, for example, would never survive if they bore litters like rabbits; the rabbit species would never thrive if rabbits lived as long as elephants or people. Quite simply, birthrates, aging rates and spans of life are intrinsic to the genetic endowment of the species.

The human genetic disease progeria makes the point - persons bearing mutations in the LMNA gene manifest accelerated aging and usually die of "old age" in their teenage years¹⁰¹. Standard life times are clearly demonstrated in the turnover rates of the cells that compose our bodies - different cell types manifest half-lives in tune with their healthy functions¹⁰². A cell that competitively out-survives its fellows can grow into a tumor and kill the organism¹⁰³; unbridled survival is destructive. Death, like reproduction, cannot be left to the whims of entropy - entropy is disorganized and death by entropy, although inevitable, is too disorganized to be adaptive. A lifespan is an adaptation⁹⁴, not a prize for successful competition. We propose that programmed birth and death are advantageous to a species because they preempt entropic disorder by imposing order on the inevitable turnover of individuals. Death, the ultimate disorder, becomes ordered.

Diversification, poly-determination, attractors and saltation

Diversification, rather than uniform optimization, characterizes living systems

Replicates of any single material entity are necessarily diverseno two genetically identical cells, for example, can occupy the same place at the same time; each cell expresses its own history and resides, as it were, in a separate environment. Beyond this fundamental, physical diversity, living systems have evolved to express higher orders of biological diversity. The concept of survival-of-the-fittest would lead one to expect that evolution should produce a single, optimal plan, approaching "perfection", in the words of Darwin¹. But evolution does not produce a single optimum; evolution is marked by continuous diversification-living systems, including prokaryotes, eukaryotes, multi-cellular organisms, species and ecosystems, are composed of component parts and processes that are intrinsically variable and diverse. Diversification of genomes is evident in the tens of millions of single-nucleotide polymorphisms (SNPs) and other genetic variations found in healthy people¹⁰⁴.

Tumors express the advantages, at least for the tumor, of diversification: a tumor that includes a great diversity of clones manifests relatively more aggressiveness, greater resistance to therapy, and enhanced fittedness when metastasizing to new organs¹⁰⁵. Lethal tumors have no evolutionary future, neither for themselves nor for their hosts, but they do illustrate a short-term advantage of diversification, however selfish and ultimately self-defeating.

Poly-determination extends diversification and retards entropy

Amazingly, living systems can utilize different combinations of their given component parts and processes to achieve a relatively uniform output. No single internal plan monopolizes a particular system. In other words, living systems can be said to be poly-determined—a given arrangement or behavior can be produced by multiple, diverse networks of interaction¹⁰⁶. A goal in football (soccer) is an accessible example of poly-determination: a scored goal is a goal, but no two are ever achieved through precisely identical field play. The poly-determination of sports is obviously an invention of human game designers; in contrast, the poly-determination of living systems has been selected by the nature of evolution.

Note that poly-determination differs from redundancy; a redundant structure or process is perceived as being a replicate of the one ideal solution; poly-determination, in contrast, is intrinsic to the multiple ways the system works to generate a given output.

A telling example of poly-determined diversification can be seen in the formation of the human species; in how many different ways has evolution been able to devise a human? We can safely assume that no two humans have ever housed identical genomes, brains, immune systems and microbiomes—monozygotic twins may emerge from a single fertilized egg, but their brains and immune systems quickly diverge and their microbiomes and epigenetic landscapes also differ. Since billions of individual holobionts have populated the human species since its inception, we can conclude that evolution has been able to concoct diverse instantiations of the human species using billions of different component recipes.

Poly-determination is also evident in the microbiome-host relationship: diverse combinations of different gut bacteria may perform similar metabolic or other functions in the host gut; in other words, a particular function may be realized by diverse combinations of microbes; as Doolittle and Booth have said: "the song is more important than the singer"¹⁰⁷.

Diversification is another name for *individuality*—no two are exactly the same; hence, evolution, as it were, has fostered individuality. Even artificially inbred mice caged together and fed identical diets manifest individual differences¹⁰⁸. Genetically identical round worms, too, manifest individuality¹⁰⁹.

Diversification, poly-determination and individuality are consequential to disease and therapy: the basic biological differences between people frustrate the traditional assumptions of Western Medicine that a given disease will appear essentially the same in different people and that an effective treatment will work in essentially all patients; actually, different subjects diagnosed with diabetes, dementia, cancer or other conditions, or people infected with a given virus or bacterial pathogen each express different clinical manifestations and will respond differently to particular treatments. Personalized medicine is a necessary consequence of entropy-selected diversification.

Biologic diversification enables living systems to keep ahead of entropy—the failure or loss of one arrangement or interaction can be compensated by alternatives. And, diversity is not a onetime thing; as we shall discuss shortly, ongoing diversification is one of the outcomes of sexual reproduction.

A species acts as an attractor stabilized by polydetermination

Above, we noted that a stable species, like the human species, is poly-determined: the species emerges from variable dynamic interactions between its varying component organisms; different human organisms with their different microbiomes and different states of structure and activity interact to constitute a single species. From this perspective, a species can be likened metaphorically to an *attractor*¹¹⁰; an attractor is mathematically defined as a set of number values towards which a dynamical system will tend to evolve. Similarly, despite their individual differences, the collective of organisms belonging to a given species will tend, on the whole, to exhibit a characteristic species profile; humans will be humans irrespective of their age, gender, ethnicity, and other individual differences; dogs will be dogs despite their various strains; different peach trees will bear peaches, and so forth. A poly-determined species remains stable despite the differences between its component organisms. Diversification and poly-determination endow a species with robust behavior and continuity that actually exploits entropic variation.

Saltation emerges from diversification and polydetermination

SNPs and other diversifications intrinsic to healthy genomes suggest that single random mutations are not likely to change gene functioning; this resistance to phenotypic change indicates that one or a few mutations are not likely to affect an organism functionally. Indeed, healthy tissues tolerate large numbers of mutations¹⁰⁴; the development of a tumor, as we pointed out above, arises from critical "driver mutations"¹¹¹. Since healthy genomes are intrinsically diverse, a meaningful evolutionary change in a given species is also likely to withstand small genetic changes or to a critical "driver" mutation affecting a key phenotypic character.

According to this reasoning, evolution need not advance only in small steps, as taught by Darwin, but may also progress in large jumps, or saltations¹¹². Darwin favored evolution by small increments for fear of encouraging creationist and intelligent design thinking. The roles of diversification and poly-determination in taming entropy, however, provide a physical rationale for saltatory evolution. Survival-of-the-fitted explains saltations as arising from the accumulation of multiple invisible changes that 'finally' exceed a certain quantum threshold; indeed, such changes can involve interactions with other organisms, materials or processes leading to entirely new entities. Phenotypical saltation poses no threat to evolution theories based on small changes.

Sexual reproduction and diversification Sexual reproduction maintains the continuous diversification of species

Sexual reproduction, a defining characteristic of multi-cellular species, entails the random mixing in the offspring of the gene alleles derived from different parental organisms. Gene mixing by sexual reproduction guarantees that no multi-cellular organism will sexually transmit its exact DNA genotype to the next generation. No matter how great the Darwinian fitness of one parent, the baby will never inherit that fit genotype alone – a new baby always inherits a random mix of half the alleles of each of its parents. Sexual reproduction ensures that genetic fitness is not transmissible intact from one generation to the next. By constantly reshuffling genomes, sexual reproduction frustrates the advancement of Darwinian perfection, as assumed to result from the reproductive success of the fittest individuals¹.

In his book¹¹³, Graham Bell asserts that "*Sex is the queen of problems in evolutionary biology*". The problem is still open; it is clearly expressed by Burke and Bonduriansky¹¹⁴:

• "Why sexual reproduction is so widespread despite its substantial costs is one of the most important unsolved problems in evolutionary biology. Because sex is associated with numerous short-term costs that asexual organisms mostly avoid, theory predicts that asexual or parthenogenetic lineages should outcompete and outnumber sexual lineages, all else being equal. However, paradoxically, sex is the dominant mode of reproduction in many lineages of complex eukaryotes."

The community has not yet agreed on a mechanism consistent with survival-of-the-fittest struggle and the debate continues - see 115,116. However, sex, from the perspective of fittedness, is not a problem for evolution, but a solution.

Sexual reproduction eschews the optimum, while it creates continuous innovation – we dare say that the optimal response to entropy is to avoid a single optimum. Darwin himself grappled with the problem of explaining sex¹¹⁷. In any case, diversification is programmatically linked to the renewal of members of a species by sexual reproduction. An individual's offspring automatically diversify the species in each generation. Quite simply, living systems are not constrained by a single "optimum" species genotype; they are programmed to avoid a single optimum; diverse arrangements retard destruction by entropy. Continuous diversification is obviously beneficial, but it is not an obvious outcome of a survival-of-the-fittest mechanism.

Of course, we must not ignore the function of sex in bonding, cooperation, family, and pleasure⁴⁹—evolution has programmed vertebrates like us to enjoy interactive life. Repeated sexual interactions thwart entropy even when they don't necessarily generate renewal.

Survival-of-the-fitted can account for same-sex sexual behavior

Sexual encounters between organisms of the same sex, widely observed in over 1,500 species of organisms, contradicts a basic assumption of Natural Selection¹¹⁵: homosexual interactions produce no offspring; hence there is no survival-of-the-fittest mechanism for transmitting their fitness to future generations of the species. Survival-of-the-fitted, however, can explain the existence of homosexual behavior as a social interaction that enhances cooperative bonding, contributes to the

environmental niche of the evolving group and restrains entropy. Thus it is not surprising that a potential for homosexuality may indeed be genetically maintained within a breeding population¹¹⁸.

The theory of neutral evolution

The theory of neutral evolution aims to explain the development within a species of genetically encoded traits that do not seem to make significant contributions to survival and reproduction; neutral evolutionary characters somehow are free of selection by a survival-of-the-fittest mechanism. Most neutral evolution has been described at the molecular level and includes molecular polymorphisms, variations in protein or nucleic acid sequences between organisms^{119–121}.

Kimura proposed that the mechanism of neutral evolution was based on random mutations that did not affect protein functions; contending selectionists argued that some sort of selection must have taken place¹²². Most evolutionary biologists today would agree with Kimura that neutral evolution results from random genetic mutation and random gene drift¹²¹.

The theory of neutral evolution marks a major distinction between Darwin's theory of Natural Selection and our theory of entropic selection: clearly, there exist genetic traits that manifest no obvious impact on individual survival and reproductive success. However, we reason that all manifestations of evolution must have undergone selection by restraint of entropy. The existence of universal entropy obliges all outcomes of evolution to undergo selection—there can be no neutrality; all persisting living systems and their component parts must withstand entropic dissolution. Evolution cannot be neutral in the face of entropy; all existing variants must, in some way, help maintain stability, or at the very least not totally sabotage stability.

Species and ecosystems

A species is a collective of individual organisms defined by the potential for mutual sexual reproduction

A universal outcome of evolution is the organization of organisms into higher scale species and ecosystems. Species and ecosystems are like corporations – a corporation is defined as an organization whose continuous existence is independent of the turnover of its individual members¹²³. Member organisms (and their molecules) come and go, but the species as a whole persists. Each organism is a transient subunit in the corporate body of a species and an ecosystem.

The organization of multi-cellular organisms into species is so plainly obvious that one might hesitate to enquire in public about its evolutionary origins. Darwin's book *The Origin of Species by Natural Selection*¹, despite its title, did not explore the origin of species as a universal adaptation but rather the origin of the local evolution of one species into another species. The existence of species was apparently seen as an unquestionable given – just as emperors are assumed to be wearing clothes, individuals are assumed to come dressed in species. Let's penetrate the obvious to ask what might be the evolutionary advantage of the higher scale organization of individual creatures into many different species. The word species originates from the Latin term designating entities that look alike (from *specio* – I see). Biologists, in former times, did not know about DNA. We now use the term species in eukaryote biology to designate a collective of creatures that can breed mutually by sexual reproduction thanks to their similar DNA¹²⁴. Not all members of a species need to actually reproduce and they may reproduce only with certain other members of the species, and only at particular times; but they could do so at least potentially. The term species is also applied to prokaryotes, but only some prokaryotes reproduce sexually and there is no functional definition of prokaryote species; for practical purposes, similarity of 16S rDNA sequences is arbitrarily used to define bacterial species¹²⁵. We shall not discuss here the effect of entropic selection on the evolution of prokaryote species.

The linking of species to sexual reproduction in eukaryotes is functional; a species thwarts entropy by the continuous rebirth of its diverse members despite their inevitable death; a species, by way of sexual reproduction, also reshuffles its renewal continuously. Furthermore, sexual reproduction, by species-specific exclusivity, constitutes a barrier against invasion of the species by members of a different species; thus sexual reproduction creates a boundary, as it were, that encloses the species.

Another defining function of a species is the ability of its member organisms to construct and thrive in a shared environmental niche; the members of a species, in principle, can each perform the interactions that characterize the species – the collective of members interacts to exert an amplified effect on their shared environment. In addition, malfunctioning members of the species are replaceable.

Moreover, members of a species can specialize in particular functions: For example, male and female organisms perform different roles in social activities, group defense, habitat construction, and so forth; alpha males, for example, maintain a pack's genetic vigor; experienced adults can educate the newborns; musically gifted persons help maintain social cohesion. So the packaging of organisms into species enables renewing, collective existence and functional similarity along with environmental continuity and individual specialization. A species can be understood as a stable attractor of diverse, continuously turning over organisms. Different species cannot reproduce across their borders, but they may link together to form ecosystems.

Ecosystems transform entropy into order

An ecosystem is a corporation of multiple species and environments that efficiently channels flows of energy through a biosphere network¹²⁶. In an ecosystem, the byproducts of one species can serve as a resource for another species linked in the network. The fecal and urinary excretions of herbivores fertilize the earth and so enable the existence of all the creatures that make their living from soil or from plants; weak and dying prey animals sustain their predators, who, in turn, maintain the health of the prey species and of the environment; dead animals feed many other species along with myriads of prokaryotes who renew the flows of vital chemicals throughout the biosphere. Indeed, the dead body of one whale on the floor of the deep ocean – whale fall – can form the nucleus of an ecosystem that sustains many species including giant isopods, lobsters, shrimp, prawns, hagfish, crabs and others for decades if not for a century¹²⁷. See the whale as a conveyance for harvesting the ocean surface for sun-dependent forms of life and bringing, in the molecules of its dead body, the energy-rich output of the sun to the darkness of the ocean depths. Ecosystems transform sunlight and other disordered energy sources into the staff of life of the biosphere as a whole. Recall that atmospheric oxygen, essential to many forms of life, is the toxic waste of photosynthesis¹²⁸.

Maintenance and repair

Subsystems of maintenance and repair have evolved in response to entropy

Individual living cells and multi-cellular organisms, in responding to entropy, have evolved subsystems that function to repair damage inflicted by accidents, noxious parasites, injury or just plain wear-and-tear. Much of the genetic endowment of individuals and species has evolved to encode molecules and processes directed to repair. Heat shock proteins and other stress responses have been conserved since the onset of prokaryote life¹²⁹; DNA repair processes keep genomes healthy¹³⁰; immune systems of various types have evolved in all species of organisms, including prokaryotes¹³¹. Life has been busy repairing itself since its beginning. We need not get into the details of these and other subsystems of maintenance; their very existence highlights the daily struggle of life with entropy.

Environments and co-evolution An environmental niche emerges from a network of biological and material interactions

Any discussion of evolution must include environmental niches. The *niche* is that segment of the environment within which a species lives and interacts; the niche includes the material, biological and behavioral factors that house the species^{132,133}. The word niche derives from the Latin *nidus*, a nest.

Classical Darwinian theory has tended to present environments as fixed arenas that provide only limited amounts of the energy, materials and space needed for survival; as a consequence, organisms are obliged to struggle amongst themselves for a livelihood¹. Survival-of-the-fittest, as we said, is assumed to reward the individual winners with a larger piece of the fixed environmental pie and with a reproductive advantage that disseminates their more fit genes. Darwin's "tangled bank" of life arises, he concludes, "from the war of nature, from famine and death." However, we see a different picture when we view the environmental niche from the perspective of fittedness: a life-sustaining environment extends beyond serving as a mere source of energy, matter and living space; an enabling environment includes interacting networks of organisms and ecosystems – effectively, the essential environment is itself a tangled web-of-life¹³³.

In contrast to Darwinian reasoning, survival does not depend merely on the ability of a cell, organism, or species to grab a maximal amount of physical energy, matter and space. Survival is not mere sustainability; survival requires that a living entity be integrated within biological and material networks that have evolved to convert entropic disorganization into organized construction. Failure to fit, or the loss of its niche, dooms the cell, organism or species to unrestrained dissolution by unrelenting entropy. Fittedness means fitting in.

Living systems co-evolve with their niches

In contrast to classical survival-of-the-fittest thinking, successful organisms and species find or actually construct their own tailored niches. Consider the human species: the human and the chimpanzee diverged from a common great-ape ancestor about 6 or 7 million years ago¹³⁴. But we evolved without competing with great apes for the same environmental niche. An objective observer would conclude that we humans arose as failed apes. We simply left the apes to their natural environments and invented our own niche. The species of plants and animals that we have domesticated (wheat, maize, rice, fruits, vegetables, chickens, ducks, horses, camels, sheep, goats, and the rest) owe their success to having "found" us to serve as their environments. Deer, barn owls, raccoons, rodents, falcons, pigeons, bats, cockroaches and other feral creatures have constructed their unique niches in our belfries, homes, gardens, farms, suburbs, cities, golf courses, and more; various types of bacteria have adopted our hospitals and our bodies.

We humans evolved human culture as our niche; but niche building is not exclusive to us^{135,136}: beavers are famous for damming lakes to build their own pools; soils are created by combined activities of prokaryotes of various types, plant roots, ants, worms, moles and other creatures. Predator and prey mutually signal each other and organize the kill to maintain both species^{137,138}. Even bacteria build niches in particular organs in our bodies by their metabolic products and their adjustment of oxygen and acid concentrations^{139,140}. Environment-building has been a feature of life on earth from earliest evolutionary times. Levins and Lewontin put it this way:

It is impossible to avoid the conclusion that organisms construct every aspect of their environment themselves. They are not the passive objects of external forces, but the creators and modulators of these forces. The metaphor of adaptation must therefore be replaced by one of construction...³⁴

To live is to find or make a place in the web-of-life

If, indeed, every species fashions its own environment, we may conclude that no two species ever occupy exactly the same environment. Different species may share a physical space in land, sea or air, but each species interacts in its own unique way with that space and with its other tenants. Life converts generic "space" into a tailored "place", a private address, as it were (see discussion in 141). In other words, the *attractor* that stabilizes a living system is not the species alone, but the species together with its co-evolving niche.

The co-evolution of holobionts and tailored niches ensures microbiome transmission

The evolution of any species can take place only if its fittedness, selected by retarding entropy, can be passed on to future generations of offspring. The problem is that the microbiome component of a holobiont is acquired independently of host genetic reproduction; how then can the fittedness of a holobiont group be transmitted somatically to holobiont offspring? The answer, we reason, is by way of niche construction and co-evolution.

The niche of each species has co-evolved with the species in many ways, some of which enable faithful transmission of the collective of microbiome elements to the next generation—not genetically, but by a combination of somatic conditions. In plants, the seeds of a species germinate in particular types of soil suited to the species—suitable soils contain microbiomes needed by the species, within the soil itself or present in adult plants growing close to the seedlings; organisms living in water are bathed by flowing concentrations of suitable microbiomes; eggs and chicks are nested in close microbial contact with their mother birds; newborns emerging from birth canals acquire mother's microbiomes in the course of being born⁸.

The environmental foods that have co-evolved with resident organisms can provide needed microorganisms: humans, for example, are attracted to fermented food stuffs rich in suitable organisms such as lactobacilli, which also help preserve the food; animals transmit microorganisms when they groom and fondle each other¹⁴²; courting rituals and sexual interactions can transmit microbiomes; some species eat feces; attractant odor organs are usually located at oral and anal orifices; attractive odors of men and woman are produced though fermentation of secretions by skin bacteria. The reader may supply additional examples of environments and behaviors that transmit microorganisms among holobionts belonging to one or more species.

Of course, changes in the environment of a species can lead to the loss of an essential element of a species microbiome: Martin Blaser has pointed out that the overuse of antibiotics has contributed to serious pandemics of chronic diseases (diabetes, obesity, heart disease, liver disease, and others) by eliminating from the human environment critical microbiome organisms¹⁴³. Our discovery of antibiotics and their extensive over-use has changed for the worse the environment of the human species, and that of other species too. We are paying a high price, and, if we don't act to characterize and preserve essential microbes, the price will only rise.

Niches are embedded

Species construct collective niches that emerge from the individual niches constructed by the organisms that compose the species. Human culture exemplifies embedded niches: each of us (our microbiomes included) constructs our own dynamically evolving niche as we progress through life, from birth, development, education, occupation, family, and so forth. Each personal niche is fashioned with regard to collective cultural niches that precede, accompany and supersede us—language, systems of belief, values, polity, personality types, science, technologies, diet, and so forth. Niches are multi-scale fractals: niches within niches within niches. Life cannot rest at one scale; entropy must be dealt with at all habitable scales: molecular, genetic, metabolic, cellular, organismal, environmental, etc.

Physical clocks organize biological clocks to help thwart entropy

The orderly dynamics of the physical environment impose entropy-restraining order on life. Many of the most formative arrangements and processes of living systems are organized by biological clocks - including metabolic reactions, cycles of nutrients, courting and fertility, conception, growth, development, migrations, aging, regeneration and repair, illness and death. Life is organized by recurrent time and the clocks of life are tuned to the cycles of nature manifest in the repetitive movements of the earth, moon and sun, which are expressed through cycles of precipitation, tides, seasons, temperatures, weather, dark and light, and even the frequencies of light. Living systems are also tuned to long-term manifestations of physical reality such as the movements of land masses, volcanic activities, and magnetic pole migrations. The details are far beyond the scope of this paper; we only need note that living systems dance to organized cadences of physical reality. The interactions of life have thus evolved to restrain entropic disorder by their linkage to the enduring physical clocks and order of material nature.

Invaders and extinctions

Species can go extinct by niche deconstruction

Evolution, we said at the outset, requires selection of some variations for propagation and others for destruction. At the level of the organism, propagation results from fitting into a network of supporting interactions; the destruction of the unfitted is carried out by the inexorable process of unrestrained entropy. At the level of the species, propagation is effected by construction of a co-evolving environmental niche supported by a flow of ecosystem energy. What causes the extinction of a species? Species can go extinct by the loss of their niche. For example, the dinosaurs underwent extinction when their niches were deconstructed in the wake of the earth's collision with an extraterrestrial body¹⁴⁴. The mammals have replaced the dinosaurs, not by struggling with them, but by constructing new niches with a variety of more effective technologies. The evolution of the organs of speech by H. sapiens may have enabled modern humans to replace the Neanderthals by constructing a more advanced social communications niche145. Likewise, Internet shopping is replacing big-box retail stores¹⁴⁶ because Internet corporations like Amazon have evolved to exploit advanced Internet communications technology (a new mutation, as it were) to construct a new niche environment; the environments of many big-box stores are now drying up following "collision" with the Internet. Niches are found (or made) and niches are lost (or deconstructed). The loss of a niche means the loss of its species; the loss of a person's home niche can generate a homeless person whose niche becomes a street corner.

Invaders can destroy the niches of other species

An invader species is one that enters and thrives in a locale in which it had not previously evolved; often the success of an invading species leads to the decline or loss of other species at one time "native" to that locale. In Israel we have witnessed the spread of Grey Crows from the countryside into human neighborhoods, associated with the disappearance of the local populations of many birds including Bee-eaters; similarly, Myna birds, escaping from the cages of bird fanciers, have proliferated at the expense of house Sparrows, that themselves were invaders from England accompanying British soldiers. Perhaps the most successful invader of once natural habitats have been modern humans, who came out of Africa many tens of thousands, perhaps hundreds of thousands of years ago to take over the planet. Casting the observed events as a struggle between species followed by victory and conquest by one of them does not negate the broader view of this reality as a deconstruction of native networks of interactions in the wake of the construction of new niche interactions. Networks of interaction do not necessarily "compete" with each other, and many such networks in fact continue to coexist.

Obviously, niche deconstruction is not the only way to extinction; a species can be eradicated by a viral or bacterial pandemic, or by the dissemination of some toxic substance, or by predators, or by atomic reactions, and so forth. Entropic mishaps have no limitations.

Humans are responsible for widespread niche deconstruction

Fittedness shoulders human culture with responsibility – we humans, who uniquely exercise conscious choice, are accountable, at least to ourselves, for our role in the state of the biosphere. Even when the human species numbered only some few millions of hunters and gatherers scattered about the earth, we appear to have eradicated many other species using only fire and weapons of stone and wood¹⁴⁷. Now we number in the billions, and are armed with ever-advancing technologies. Without intending to deconstruct the "natural" environment, our domestication of a small number of species has markedly affected the biosphere: the biomass of chickens by now far outweighs the combined biomass of all wild birds and the biomass of cattle outweighs by many fold the biomass of all other mammals combined, including us¹⁴⁸. The states of the earth and the biosphere are in our hands.

According to classical Darwinian thinking, to be fit is to win in the struggle to exploit an ever larger slice of a fixed environmental pie. The environment was viewed as a battle ground on which we, and the rest of the earth's creatures, must fight to survive. Unfortunately, liability for undue exploitation is not inherent in the concept of Natural Selection. One may wonder whether survival-of-the-fittest thinking may have played a role in the ideology of unrestrained exploitation of the earth for economic and political "gain". Our niche encompasses the world; unbridled "gain" will deconstruct us all.

The emergence of complexity

Complexity is easier to recognize than it is to define. But however one defines complexity, it is evident that a biosphere that includes multi-cellular organisms is more complex than a biosphere composed of prokaryotes only^{64–66}. Indeed, increasing complexity seems to evolve relentlessly; see for example the quote "*Everybody seems to know that complexity increases in evolution*" stated by McShea¹⁴⁹. The general increase in complexity is balanced by evidence for particular reductions in complexity, as exemplified in genome reductions discussed by Wolf and Koonin¹⁵⁰; indeed, salmon parasites show an entire loss of the mitochondrial genome and associated nuclear genes required for its transcription¹⁵¹.

Complex species repeatedly have undergone mass extinctions¹⁵². Yet evolution, as it were, picks up the simple leftovers and starts anew accumulating complexity. Why is this so? Complex systems are not more fit in Darwinian terms than are simpler systems; simple systems seem in fact to be much fitter; bacteria have survived all the extinctions suffered by the more complex organisms and will be more likely than us to survive global warming or nuclear catastrophe¹⁵³.

Entropic selection is compatible with the idea that complexity itself fuels more complexity¹⁵⁴. Prevailing energy and entropy, along with biological diversification, continuously drive the emergence of new mutations, combinations, interactions and functional variations. A given molecular entity will tend to increase its range of interactions over evolutionary time; the oxytocin molecule, for example, has evolved to perform a range of functions in bonding, sexual and social behavior, metabolism, inflammation and the control of stress and aggression49. In each case the oxytocin molecule has added new molecular partners and so, over time, it has participated in new and more complex network interactions. Heat shock stress proteins first evolved as chaperones in primitive bacteria; they have gone on to serve as chaperones in eukaryote cells and in multi-cellular organisms, and have evolved into important signal molecules for the mammalian immune system¹⁵⁵.

Evolution seems to have created increasing complexity by adding innovations to its existing molecules, cells, organisms and species. Certain aspects of this general process have been studied under the concepts of preadaptation or exaptation¹⁵⁶. Round worms express some 20,000 genes¹⁵⁷ compared to a similar number of genes expressed in humans¹⁵⁸. These numbers are estimates subject to revision. Nevertheless, it is clear that evolution has proceeded by establishing increasingly more varied interactions and functions for a given number of genes already present in its primitive toolbox; chimpanzees and humans share more than 95% of their genes¹³⁴—the greater complexity of humans, expressed in language and global culture, emerges from more complex uses for common genes. Advancing complexity seems to have emerged both from novel regulatory DNA sequences and even from gene loss and not necessarily from entirely new genes^{159,160}.

Autocatalysis of complexity is also expressed in the increased likelihood that a random variation or mutation in a gene, molecule, cell, or organism will by chance find a fit in a large landscape of different interacting entities. A complex infrastructure will tend to provide opportunities to increase its complexity; consider the ever-increasing complexity of the Internet's evolution before our eyes. Indeed, the jump of viruses like Ebola, influenza and corona from animals into humans has resulted from the advancing complexity of human culture.

The question of a unit of evolutionary selection A unit of universal selection is controversial

The designation of a "unit of evolutionary selection" is an attempt to pinpoint the primary substrate upon which the process of evolution directly acts; the unit of evolution defines the seminal entity whose evolution generates the life we see around us. Which entity is the prime mover of evolution and which entities are secondary consequences? In the beginning, Darwin proposed that individuals are the primary units driving evolution; according to Darwin the character of any species is a secondary result of the procreative success of the fittest individuals engaged in the struggle for survival¹. Dawkins, in contrast, argues that the gene is the primary replicator unit of evolution; individuals are merely transient phenotypic vehicles that bear the unit genes³⁹. Lewontin includes the whole spectrum of biological entities as units of evolution - genes, individuals, groups and species; according to this view, life at all its scales evolves together¹⁶¹. Zilber-Rosenberg and Rosenberg proposed the holobiont as a unit of selection⁶. Views and debates on the issue are reviewed by Lloyd in 162; the discussions nevertheless continue 11,163. These designations of a unit or units of evolution assume that survival-of-the-fittest is the driving mechanism of evolutionary change; how do we define the unit of evolution from the perspective of survival-of-the-fitted?

Above, we have argued that the mechanism of universal evolution involves the interplay of three universal properties of matter: energy, entropy and interaction. These three properties exert effects on the evolution of living entities across all scales - from molecules through ecosystems. From this perspective, no single entity is "the unit" of evolutionary selection. Survivalof-the-fitted is a mechanism that makes no distinctions between any of the entities susceptible to evolution; life is the expression of its energy-dependent interactions, and all interactions must pass the tests of entropy. We would agree with positions that argue that the whole spectrum of life's entities takes part in evolution. But we see particular entities-from cells to ecosystems-as secondary manifestations of the primary evolution of networks of interactions. If one wishes to maintain the concept of a unit of evolutionary selection, than the unit is not a particular entity like a gene or individual organism, but rather is the dynamic process of interaction itself. Indeed, the phenotype of a gene is formed by the interactions that generate gene expression, translation into proteins, post-translational modifications and all the rest. The organism, too, is a dynamic product of all its internal and external interactions. We may perceive entities, but all are snapshots of dynamic interactions. The progenitor of evolution is interaction.

Theories of evolution influence human behavior The concept of survival-of-the-fittest has been misused to justify cultural domination

Whether or not Natural Selection is a "law of nature" is controversial among biologists^{164,165}; nevertheless, extreme segments of Western Culture have usurped survival-of-the-fittest to justify an ideology of conquest and dominance. Lamentably, Natural Selection, at one time or another, has been invoked to justify colonialism, racial purity, eugenics, genetic cleansing, dictatorships, unfettered capitalism, unequal rights, and other harsh dissonances (see, for example, 166,167). Obviously, survival-of-the-fittest is a scientific theory and not a benchmark for human behavior; nevertheless, Natural Selection has been hijacked by agendas of exploitation.

Updating the mechanism of evolution, as we have done here, to highlight the roles of *cooperation, interaction* and *diversity* throughout biology might serve to encourage rational human behavior in the realms of governance, economics and social justice.

How can we reconcile the holobiont human with the individual human?

The discovery that all multi-cellular organisms are holobionts raises a paradox; at the biological level, there are no individuals; living systems, nevertheless, do manifest group individuality. Yet, the psychological core of each human is his or her feeling of an exclusive existence expressing private thoughts, feelings, and a specific history of being and relationship. Our internal perception of uniqueness is confirmed independently by our individual social and legal responsibilities. Can we resolve the seeming contradiction between our realities as both biological groups and functioning individuals?

Look at it this way: our psychological and institutional individuality, like our body, is created by group interactions and relationships, not merely with microbes, but with the people and the world in which we reside; our identity is the outcome of our history of bonding with others. Echoing Levins and Lewontin³⁶, we construct our niche by the nature of our interactions—we and our worlds co-evolve. We achieve a self by constructing a shared, interactive environment. Thy neighbor is part of thy evolving self.

Reconciling survival-of-the-fitted with Darwinian survival-of-(only)-the-fittest

Physicists point out that life, like all the rest of the material universe, must accommodate the Second Law of Thermodynamics and the phenomenon of $entropy^{27,29}$. Here, we have reasoned that the evolution of life, a dynamic expression of life itself, must also accommodate entropy. Consequently, stable outcomes of the evolutionary process must provide some solution to the spontaneous dissolution of order; entropy, in one way or another, must act as a selective factor in evolution – it would not be reasonable to conclude otherwise.

What about the concept of Natural Selection, a cornerstone of modern biology? One may ask, why not conclude from the above discussion that entropic selection also serves survivalof-the-fittest, without invoking survival-of-the-fitted. In other words, perhaps Natural Selection would select, among others, those individuals who counteract entropy most efficiently and, as a result, leave more progeny than others. This, however, does not work for at least four reasons: as we have discussed above, (i) there is no competition in dealing with entropy; a successful individual or species need not retard entropy most efficiently or resist entropy better than others, but only fit in sufficiently with a supportive network of interactions; (ii) not only the one most reproductively successful variant survives, but many others readily find their niches; (iii) many obviously advantageous features are not rewarded with reproductive advantage; and (iv) increased reproduction is at times more of a problem than a desired outcome.

It is conceivable that different manifestations of evolution might be assigned to a combination of fitness (Natural Selection) and fittedness (Entropic Selection). Entropy would seem to account for the more universal adaptations of evolution we have discussed here. In contrast, Natural Selection might drive more local, species-restricted adaptations. Certainly, the quantitative assessment of each mechanism would be different: the advance of evolution by individual fitness would require measurement of the birth rates of generations of descendants of particular test individuals - a difficult, if not impossible task; fittedness based on the strength of ongoing cooperative interactions could be measured, at least at the molecular level, by the amount of energy needed to disrupt the interaction: For example, the strength of the interactions that hold together dsDNA strands is reflected in the amount of heat that has to be applied to disrupt the interaction, as in the PCR reaction¹⁶⁸. In any case, a fresh look at the mechanisms of evolution is in order.

Analysis and modeling

A theory can be invoked to explain any phenomenon; a theory is scientific when it proposes a platform for continued rational analysis and when it can support testable predictions. How may we proceed to analyze survival-of-the-fitted?

Fittedness can be analyzed using tools of systems biology

Survival-of-the-fitted fits generally into the realm of systems biology, including dynamic complex systems and their attractor states. The perspective and the methods of systems biology go beyond a linear chain of discrete causes and effects; advancing information technologies have extended the scope of biology research, enabling analyses of complex interactions that generate dynamic self-organizing systems. "Big data" are collected, defined and analyzed for associations involving networks, modules, emergent properties and other organizational structures and principles. Systems' understanding emerges from simulations and mathematical models^{169,170}.

Such modeling could build upon the known generation of orderly structures from chaotic thermodynamic behavior at non-equilibrium states, at multiple scales (see, for example Turing¹⁷¹, Prigogine¹⁷², Kauffman³³, Jimenez¹⁷³, and England¹⁷⁴). We plan to model how such erratic structures will survive longer if they chance to fit into supporting interactions with existing structures; this mechanism could well provide novel arrangements for advancing evolution.

Stuart Kauffman has modeled the emergence of life and the biosphere using the concepts of living organisms as *autonomous agents* who navigate *fitness landscapes* to *make their living* with the aid of genetic *code scripts*^{33,88}. These concepts well suit a survival-of-the-fittest mechanism. However, our proposed evolutionary mechanism of entropic selection of holobionts and cooperative interactions for fittedness does not require the concepts of autonomous agents, fitness landscapes and gene-centered competitive success. Our nascent models are likely to differ from those considered in the past. In any case, the experiments of evolution have been and are being done by nature, analyses of the results need human computation. A system's biology of evolution is in the offing.

Predictions

Here, we propose some experiments and predictions, not all of which are currently feasible, to test survival-of-the-fitted:

1. Germ-free mice or rats are raised behind barriers and deprived of gut microbiomes, but they can be reconstituted by feeding them with defined clones of bacteria; one group could be fed with a single monoclonal gut bacterium and other groups could be fed with combinations of diverse bacterial clones. The survival-of-the fittest paradigm would predict that the microbiomes developing in the groups of mice fed with diverse clones of bacteria would evolve over time into microbiomes populated with only a single clone of the fittest bacterium. The microbiomes in the animals fed with a single clone, in the absence of competition, would not evolve.

In contrast, the survival-of-the-fitted paradigm would predict that the microbiomes of the animals fed only with a single monoclonal species of bacteria would in time respond to entropy to evolve into a diversity of bacteria expressing different genes. The animals fed a diversity of bacteria would continue to express a diverse microbiome. Thus, the contending *fitness and fittedness* theories of evolution would each lead to contradicting predictions.

- 2. Evolution will take place in the absence of competition and in environments with unlimited resources; contrary to Darwinian teachings, a struggle for fitness is not required because entropy prevails in the best of worlds. Diversification can be tested under controlled experiments *in vitro*^{35,175}. Monoclonal, initially invariant yeast, bacteria or *C. elegans* could be grown under conditions of "optimal" nutrition and density in monitored cultures and observed for the evolution of genetic and phenotypic diversification in the absence of struggle.
- 3. Systems that foster diversity and individuality will thrive better than systems limited to an "optimal" archetype; cooperation and symbiosis will emerge in diversified collectives. This prediction can be tested today in a variety of systems at various scales, from single cells to societies and cultures.
- 4. Sexual reproduction (or an equivalent system of reproducing, programmed diversification) will evolve in multi-cellular living systems. Testing this might require the development of novel *in vitro* culture systems.

- 5. Complexity will increase automatically, even in an *in vitro* culture of initially uniform populations. Observations of new interactions could be made using long term culture systems.
- 6. Life on other planets will be seen to have evolved, provided that entropy holds sway there.
- 7. All living systems anywhere will evolve programmed renewal.

Predictions # 6 and # 7 will require the development of suitable probes of habitable planets distant from earth. Perhaps extraterrestrial life may one day be studied at a distance by telemetric analysis of metabolism or other manifestations of life.

Finally, we would predict that the well-being of the biosphere will be aided by realizing intelligent interactive cooperation and shunning selfish struggle. This prediction is now being tested by observing our changing world, obviously without a control group. Let us each work to fulfill this prediction.

Data availability

No data is associated with this work.

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References

- Darwin C: On the Origin of Species by Means of Natural Selection, or the Preservation of Favoured Races in the Struggle for Life. Annals and Magazine of Natural History. John Murray, 1860; 5(26): 132–143. Publisher Full Text
- Stearns SC: Principles of Evolution, Ecology and Behavior. Open Yale Course EEB 122, 2009. Reference Source
- Mayr E: Cause and effect in biology. Science. 1961; 134(3489): 1501–1506.
 PubMed Abstract | Publisher Full Text
- Koonin EV: The Logic of Chance: the Nature and Origin of Biological Evolution. FT press, 2011.
 Reference Source
- Methé BA, Nelson KE, Pop M, et al.: A framework for human microbiome research. Nature. 2012; 486(7402): 215-21.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Zilber-Rosenberg I, Rosenberg E: Role of microorganisms in the evolution of animals and plants: the hologenome theory of evolution. *FEMS Microbiol Rev.* 2008; 32(5): 723-735.
 PubMed Abstract | Publisher Full Text
- Rosenberg E, Zilber-Rosenberg I: The hologenome concept of evolution after 10 years. Microbiome. 2018; 6(1): 78.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Rosenberg E, Zilber-Rosenberg I: The Hologenome Concept: Human, Animal and Plant Microbiota. Springer, 2014. Reference Source
- Bhattarai Y, Kashyap PC: Germ-free mice model for studying host-microbial interactions. In Mouse Models for Drug Discovery. Springer, 2016; 123–135. Publisher Full Text
- Moeller AH, Suzuki TA, Phifer-Rixey M, et al.: Transmission modes of the mammalian gut microbiota. Science. 2018; 362(6413): 453–457. PubMed Abstract | Publisher Full Text
- Stencel A, Wloch-Salamon DM: Some Theoretical Insights Into the Hologenome Theory of Evolution and the Role of Microbes in Speciation. Theory Biosci. 2018; 137(2): 197–206.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Scott-Phillips TC, Laland KN, Shuker DM, et al.: The Niche Construction Perspective: A Critical Appraisal. Evolution. 2014; 68(5): 1231–1243. PubMed Abstract | Publisher Full Text | Free Full Text
- Waddington CH: Paradigm for an evolutionary process. Towards a theoretical biology. 1969; 2: 106–128.
 Publisher Full Text
- Gould SJ: Darwinism and the Expansion of Evolutionary Theory. Science. 1982; 216(4544): 380–387.
 PubMed Abstract | Publisher Full Text
- Laland KN, Uller T, Feldman MW, et al.: The Extended Evolutionary Synthesis: Its Structure, Assumptions and Predictions. Proc Biol Sci. 2015; 282(1813): 20151019.
 PubMed Abstract | Publisher Full Text | Free Full Text

- Laland K, Uller T, Feldman M, et al.: Does Evolutionary Theory Need a Rethink? Nature. 2014; 514(7521): 161–4.
 PubMed Abstract L Publisher Full Text
- Cohen IR: Tending Adam's Garden: evolving the cognitive immune self. Elsevier, 2000.
 Publisher Full Text
- Cohen IR: Informational Landscapes in Art, Science, and Evolution. Bull Math Biol. 2006; 68(5): 1213–1229.
 PubMed Abstract | Publisher Full Text | Free Full Text
- 19. Cohen IR: Updating Darwin: Information and Entropy Drive the Evolution of Life. *F1000Res.* 2016; **5**: 2808.
- PubMed Abstract | Publisher Full Text | Free Full Text 20. Doige CA, Day T: A typology of undergraduate textbook definitions of heat provide a dissipling of the California California (California) and California California (California) and California California (California) and California California (California) and California (Californi
- across science disciplines. Int J Sci Educ. 2012; **34**(5): 677–700. Publisher Full Text
- 21. Merriam-Webster Dictionary: **On-line dictionary**. 2019; (accessed 6/2019). Entry for "Energy".
- 22. Wehrl A: General properties of entropy. *Rev Mod Phys.* 1978; **50**(2): 221. Publisher Full Text
- Oxford English Dictionary (OED): Online Dictionary. 2019; (accessed 6/2019). Entry for "Entropy".
- Martin JS, Smith NA, Francis CD: Removing the entropy from the definition of entropy: clarifying the relationship between evolution, entropy, and the second law of thermodynamics. Evol Educ Outreach. 2013; 6(1): 30. Publisher Full Text
- Feynman R: The Feynman Lectures. Accessed May 2020. Lecture 1: Atoms in Motion. 1963.
- Cohen IR, Harel D: Explaining a complex living system: dynamics, multiscaling and emergence. J R Soc Interface. 2007; 4(13): 175–182.
 PubMed Abstract | Publisher Full Text | Free Full Text
- 27. Atkins PW: The second law. Scientific American Library, 1984.
- Boltzmann L: The second law of thermodynamics. In: Theoretical Physics and Philosophical Problems: Selected writings. Springer Science & Business Media, 2012.
- Schrödinger E: What is life? The physical aspect of the living cell and mind. Cambridge University Press Cambridge, 1944. Reference Source
- Berry S: Entropy, irreversibility and evolution. J Theor Biol. 1995; 175(2): 197–202.
 PubMed Abstract | Publisher Full Text
- Styer DF: Entropy and evolution. Am J Phys. 2008; 76(11): 1031–1033. Publisher Full Text
- Demetrius LA: Boltzmann, Darwin and directionality theory. Phys Rep. 2013;
 530(1): 1–85.
 Publisher Full Text
- Kauffman S: Investigations. Oxford University Press, 2000. Reference Source

- Schneider ED, Sagan D: Into the cool: Energy flow, thermodynamics, and life. University of Chicago Press, 2005. Reference Source
- 35. Elena SF, Lenski RE: **Evolution experiments with microorganisms: the** dynamics and genetic bases of adaptation. *Nat Rev Genet.* 2003; 4(6): 457–69. PubMed Abstract | Publisher Full Text
- Levins R, Lewontin RC: The Dialectical Biologist. Harvard University Press, 1985.
 Reference Source
- Gleick J: Genius: The life and science of Richard Feynman. Vintage, 1993. In Prologue.
 Reference Source
- Rovelli C: Carlo rovelli all reality is interaction. 2017; Accessed May 2020. WNYC Radio Program. Reference Source
- 39. Dawkins R: **The Selfish Gene**. Oxford university press, 1976.
- 40. Dawkins R: The Extended Phenotype: The Long Reach of the Gene. Oxford University Press, 1982.
- Reference Source
 41. Hilbe C, Wu B, Traulsen A, *et al.*: Evolutionary performance of zerodeterminant strategies in multiplayer games. *J Theor Biol.* 2015; 374: 115– 124.
 PubMed Abstract | Publisher Full Text | Free Full Text
- 42. Margulis L: Symbiosis in Cell Evolution: Life and its Environment on the Early Earth. W. H. Freeman, 1981.
- Reference Source
 Embley TM, Martin W: Eukaryotic evolution, changes and challenges. Nature. 2006; 440(7084): 623–30.
- PubMed Abstract | Publisher Full Text

 44.
 Mbeunkui F, Johann DJ Jr: Cancer and the tumor microenvironment: a review
- of an essential relationship. Cancer Chemother Pharmacol. 2009; 63(4): 571– 582. PubMed Abstract | Publisher Full Text | Free Full Text
- Blaser MJ: The microbiome revolution. J Clin Invest. 2014; 124(10): 4162–4165. PubMed Abstract | Publisher Full Text | Free Full Text
- Blaser MJ, Webb GF: Host demise as a beneficial function of indigenous microbiota in human hosts. mBio. 2014; 5(6): e02262–14.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Kinross JM, Darzi AW, Nicholson JK: Gut microbiome-host interactions in health and disease. Genome Med. 2011; 3(3): 14.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Leigh EG Jr: The group selection controversy. J Evol Biol. 2010; 23(1): 6–19. PubMed Abstract | Publisher Full Text
- Onaka T, Takayanagi Y, Yoshida M: Roles of oxytocin neurones in the control of stress, energy metabolism, and social behaviour. J Neuroendocrinol. 2012; 24(4): 587–598.
 PubMed Abstract | Publisher Full Text
- Gruber CW: Physiology of invertebrate oxytocin and vasopressin neuropeptides. Exp Physiol. 2014; 99(1): 55–61.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Varian BJ, Poutahidis T, DiBenedictis BT, et al.: Microbial lysate upregulates host oxytocin. Brain Behav Immun. 2017; 61: 36–49.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Rizzolatti G, Craighero L: The mirror-neuron system. Annu Rev Neurosci. 2004; 27: 169–192.
 - PubMed Abstract | Publisher Full Text
- Baird AD, Scheffer IE, Wilson SJ: Mirror neuron system involvement in empathy: a critical look at the evidence. Soc Neurosci. 2011; 6(4): 327–335.
 PubMed Abstract | Publisher Full Text
- Zentall TR: Imitation by animals: How do they do it? Curr Dir Psychol Sci. 2003; 12(3): 91–95.
 Publisher Full Text
- MacDonald KB: Separation and its discontents: Toward an evolutionary theory of anti-Semitism. Praeger Publishers/Greenwood Publishing Group, 1998.
 Reference Source
- Nagasawa M, Mitsui S, En S, et al.: Oxytocin-gaze positive loop and the coevolution of human-dog bonds. Science. 2015; 348(6232): 333–336.
 Publisher Full Text
- Mi S, Lee X, Li X, et al.: Syncytin is a captive retroviral envelope protein involved in human placental morphogenesis. Nature. 2000; 403(6771): 785–9. PubMed Abstract | Publisher Full Text
- Raymond J, Blankenship RE: Horizontal gene transfer in eukaryotic algal evolution. Proc Natl Acad Sci U S A. 2003; 100(13): 7419–7420.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Soucy SM, Huang J, Gogarten JP: Horizontal gene transfer: building the web of life. Nat Rev Genet. 2015; 16(8): 472–82.
 PubMed Abstract | Publisher Full Text
- Nowak MA: Evolving cooperation. J Theor Biol. 2012; 299: 1–8. PubMed Abstract | Publisher Full Text

- Maynard Smith J: Evolution and the theory of games: In situations characterized by conflict of interest, the best strategy to adopt depends on what others are doing. *Am Sci.* 1976; 64(1): 41–45. Reference Source
- Maynard Smith J, Price GR: The logic of animal conflict. Nature. 1973; 246(5427): 15–18.
 Publisher Full Text
- Axelrod R, Hamilton WD: The evolution of cooperation. Science. 1981; 211(4489): 1390–1396.
 PubMed Abstract | Publisher Full Text
- Maynard Smith J, Szathmary E: The major transitions in evolution. Oxford University Press. 1997.
 Reference Source
- Szathmáry E: Toward major evolutionary transitions theory 2.0. Proc Natl Acad Sci U S A. 2015; 112(33): 10104–10111.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Wolf YI, Katsnelson MI, Koonin EV: Physical foundations of biological complexity. Proc Natl Acad Sci U S A. 2018; 115(37): E8678–E8687. PubMed Abstract | Publisher Full Text | Free Full Text
- 67. Pal A, Levy Y: **Structure, stability and specificity of the binding of ssDNA and ssRNA with proteins.** *PLoS Comput Biol.* 2019; **15**(4): e1006768. **PubMed Abstract | Publisher Full Text | Free Full Text**
- Clausen-Schaumann H, Rief M, Tolks-dorf C, et al.: Mechanical stability of single DNA Molecules. Biophys J. 2000; 78(4): 1997–2007.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Hofreiter M, Serre D, Poinar HN, et al.: Ancient DNA. Nat Rev Genet. 2001; 2(5): 353–9.
- PubMed Abstract | Publisher Full Text
- Myers AH, Robinson EG, Van Natta ML, et al.: Hip fractures among the elderly: factors associated with in-hospital mortality. Am J Epidemiol. 1991; 134(10): 1128–1137.
 PubMed Abstract | Publisher Full Text
- Zisberg A, Shadmi E, Sinoff G, et al.: Low mobility during hospitalization and functional decline in older adults. J Am Geriat Soc. 2011; 59(2): 266–273. PubMed Abstract | Publisher Full Text
- Dave D, Rashad I, Spasojevic J: The effects of retirement on physical and mental health outcomes. Technical report, National Bureau of Economic Research, 2006.
 Publisher Full Text
- Smith JM: Ireland's Magdalen Laundries and the Nation's Architecture of Containment. Manchester University Press, 2008. Publisher Full Text
- Fernhall Bo: Long-term aerobic exercise maintains peak VO², improves quality of life, and reduces hospitalisations and mortality in patients with heart failure. J Physiother. 2013; 59(1): 56–56.
 PubMed Abstract | Publisher Full Text
- O'brochta D: Easi-CRISPR: Long ssDNA donors greatly increase efficiency of HR. 2017. Reference Source
- Homan A: Rhythms of nature in the Barycz Valley. 2010; accessed 12/2019. Reference Source
- Milo DS: Good Enough: The Tolerance for Mediocrity in Nature and Society. Harvard University Press, 2019. Reference Source
- Fierer N, Jackson RB: The diversity and biogeography of soil bacterial communities. Proc Natl Acad Sci U S A. 2006; 103(3): 626–631.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Swiatczak B, Cohen IR: Gut feelings of safety: tolerance to the microbiota mediated by innate immune receptors. *Microbiol Immunol.* 2015; 59(10): 573–585.
 - PubMed Abstract | Publisher Full Text
- Lewis NS, Nocera DG: Powering the planet: Chemical challenges in solar energy utilization. Proc Natl Acad Sci U S A. 2006; 103(43): 15729–15735. PubMed Abstract | Publisher Full Text | Free Full Text
- Edwards KJ, Becker K, Colwell F: The deep, dark energy biosphere: intraterrestrial life on earth. Ann Rev Earth Plan sci. 2012; 40(1): 551–568. Publisher Full Text
- Brown JH, Gillooly JF, Allen AP, et al.: Toward a metabolic theory of ecology. Ecology. 2004; 85(7): 1771–1789.
 Publisher Full Text
- Shiraki N, Shiraki Y, Tsuyama T, et al.: Methionine metabolism regulates maintenance and differentiation of human pluripotent stem cells. *Cell* Metab. 2014; 19(5): 780–794.
 PubMed Abstract | Publisher Full Text
- Palmer CS, Ostrowski M, Balderson B, et al.: Glucose metabolism regulates T cell activation, differentiation, and functions. Front Immunol. 2015; 6: 1. PubMed Abstract | Publisher Full Text | Free Full Text
- Ma C, Han M, Heinrich B, et al.: Gut microbiome-mediated bile acid metabolism regulates liver cancer via NKT cells. Science. 2018; 360(6391): eaan5931.
 PubMed Abstract | Publisher Full Text | Free Full Text

- Moore S, Evans LDB, Andersson T, et al.: App metabolism regulates tau proteostasis in human cerebral cortex neurons. Cell Rep. 2015; 11(5): 689–696.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Ginestier C, Monville F, Wicinski J, et al.: Mevalonate metabolism regulates basal breast cancer stem cells and is a potential therapeutic target. Stem Cells. 2012; 30(7): 1327–1337.
 PubMed Abstract | Publisher Full Text
- Kauffman S: Answering Schrödinger's "What Is Life?" Entropy. 2020; 22(8): 815.
 Publisher Full Text
- Berg JM, Tymoczko JL, Stryer L: Biochemistry. W H Freeman, Section 1.3, Chemical Bonds in Biochemistry. 2002. Reference Source
- 90. Zahavi A: Mate selection—a selection for a handicap. J Theor Biol. 1975; 53(1): 205-214.
 - PubMed Abstract | Publisher Full Text
- Penn DJ, Számadó S: The handicap principle: how an erroneous hypothesis became a scientific principle. *Biological Reviews*. 2019; 95(1): 267–290.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Maturana HR, Varela FJ: Autopoiesis and cognition: The realization of the living. Springer Science & Business Media. 1991; 42.
 Publisher Full Text
- Metcalfe NB, Monaghan P: Growth versus lifes-pan: perspectives from evolutionary ecology. Exp Gerontol. 2003; 38(9): 935–940.
 PubMed Abstract | Publisher Full Text
- Mangel M, Kindsvater HK, Bon-sall MB: Evolutionary analysis of life span, competition, and adaptive radiation, motivated by the pacific rockfishes (sebastes). Evolution. 2007; 61(5): 1208–1224.
 PubMed Abstract | Publisher Full Text
- Diamantis A, Magiorkinis E, Sakorafas GH, et al.: A brief history of apoptosis: from ancient to modern times. Onkologie. 2008; 31(12): 702–706. PubMed Abstract | Publisher Full Text
- Kerr JFR: A histochemical study of hypertrophy and ischaemic injury of rat liver with special reference to changes in lysosomes. *J Pathol Bacteriol*. 1965; 90(2): 419–435.
 - PubMed Abstract | Publisher Full Text
- Sulston JE, Horvitz HR: Post-embryonic cell lineages of the nematode, Caenorhabditis elegans. Dev Biol. 1977; 56(1): 110–156.
 PubMed Abstract | Publisher Full Text
- Sandow BA, West NB, Norman RL, et al.: Hormonal control of apoptosis in hamster uterine luminal epithelium. Am J Anat. 1979; 156(1): 15–35.
 PubMed Abstract | Publisher Full Text
- Koonin EV, Aravind L: Origin and evolution of eukaryotic apoptosis: the bacterial connection. *Cell Death Differ.* 2002; 9(4): 394–404.
 PubMed Abstract | Publisher Full Text
- Hughes T, Rusten TE: Origin and evolution of self-consumption: autophagy. Adv Exp Med Biol. In: Eukaryotic Membranes and Cytoskeleton. Springer, 2007; 111–118.
 - PubMed Abstract | Publisher Full Text
- 101. Ahmed MS, Ikram S, Bibi N, et al.: Hutchinson-gilford progeria syndrome: a premature aging disease. Mol Neurobiol. 2018; 55(5): 4417–4427. PubMed Abstract | Publisher Full Text
- 102. Spalding KL, Arner E, Westermark PO, et al.: Dynamics of fat cell turnover in humans. Nature. 2008; 453(7196): 783–7. PubMed Abstract | Publisher Full Text
- Nguyen LV, Vanner R, Dirks P, et al.: Cancer stem cells: an evolving concept. Nat Rev Cancer. 2012; 12(2): 133–43.
 PubMed Abstract | Publisher Full Text
- 104. 1000 Genomes Project ConsortiumAuton A, Brooks LD, et al.: A global reference for human genetic variation. Nature. 2015; 526(7571): 68–74. PubMed Abstract | Publisher Full Text | Free Full Text
- 105. Swanton C: Intratumor heterogeneity: evolution through space and time. Cancer Res. 2012; 72(19): 4875–4882. PubMed Abstract | Publisher Full Text | Free Full Text
- 106. Bunzl M: Causal overdetermination. J Philos. 1979; 76(3): 134–150. Publisher Full Text
- 107. Doolittle WF, Booth A: It's the song, not the singer: an exploration of holobiosis and evolutionary theory. *Biol Philos.* 2017; 32(1): 5–24. Publisher Full Text
- Freund J, Brandmaier AM, Lewejohann L, et al.: Emergence of individuality in genetically identical mice. Science. 2013; 340(6133): 756–759.
 PubMed Abstract | Publisher Full Text
- Stern S, Kirst C, Bargmann CI: Neuromodulatory control of long-term behavioral patterns and individuality across development. *Cell*. 2017; 171(7): 1649–1662.e10.
 PubMed Abstract | Publisher Full Text
- Milnor J: On the concept of attractor. In: The theory of chaotic attractors. Springer, 1985; 243–264.
 Publisher Full Text
- Pao W, Girard N: New driver mutations in non-small-cell lung cancer. Lancet Oncol. 2011; 12(2): 175–180.
 PubMed Abstract | Publisher Full Text

- 112. Goldschmidt R: The material basis of evolution. Yale University Press, 1940. Reference Source
- 113. Bell G: The masterpiece of nature: the evolution and genetics of sexuality. Routledge, 2019. Reference Source
- 114. Burke NW, Bonduriansky R: Sexual conflict, facultative asexuality, and the true paradox of sex. Trends Ecol Evol. 2017; 32(9): 646–652. PubMed Abstract | Publisher Full Text
- 115. Monk JD, Giglio E, Kamath A, et al.: An alternative hypothesis for the evolution of same-sex sexual behaviour in animals. Nat Ecol Evol. 2019; 3(12): 1622–1631. PubMed Abstract | Publisher Full Text
- Livnat A, Papadimitriou CH: Sex as an algorithm: the theory of evolution under the lens of computation. *Commun ACM*. 2016; 59(11): 84–93.
 Publisher Full Text
- 117. Prum RO: The Evolution of Beauty: How Darwin's Forgotten Theory of Mate Choice Shapes the Animal World and Us. Doubleday, 2017. Reference Source
- 118. Santtila P, Sandnabba NK, Harlaar N, et al.: Potential for homosexual response is prevalent and genetic. Biol Psychol. 2008; 77(1): 102–105. PubMed Abstract | Publisher Full Text
- 119. Kimura M: Evolutionary rate at the molecular level. *Nature*. 1968; **217**(5129): 624–626.
 - PubMed Abstract | Publisher Full Text
- Kimura M: The neutral theory of molecular evolution. Cambridge University Press, 1983.
- 121. Jensen JD, Payseur BA, Stephan W, et al.: The importance of the neutral theory in 1968 and 50 years on: A response to Kern and Hahn 2018. Evolution. 2019; 73(1): 111–114. PubMed Abstract | Publisher Full Text | Free Full Text
- Kern AD, Hahn MW: The neutral theory in light of natural selection. Mol Biol Evol. 2018; 35(6): 1366–1371.
- PubMed Abstract | Publisher Full Text | Free Full Text 123. Wikipedia: Entry for "Corporation". 2019. (accessed 6/2019).
- 124. Mayr E: What is a Species, and What is Not? Philos Sci. 1996; 63(2): 262–277. Reference Source
- Gevers D, Cohan FM, Lawrence JG, et al.: Re-evaluating prokaryotic species. Nat Rev Microbiol. 2005; 3(9): 733–9.
 PubMed Abstract | Publisher Full Text
- Barnes AD, Jochum M, Lefcheck JS, et al.: Energy flux: The link between multitrophic biodiversity and ecosystem functioning. *Trends Ecol Evol.* 2018; 33(3): 186–197.
 PubMed Abstract | Publisher Full Text | Free Full Text
- 127. Smith CR, Glover AG, Treude T, et al.: Whale-fall ecosystems: recent insights into ecology, paleoecology, and evolution. Ann Rev Mar Sci. 2015; 7: 571–596. PubMed Abstract | Publisher Full Text
- Lyons TW, Reinhard CT, Planavsky NJ: The rise of oxygen in earth's early ocean and atmosphere. Nature. 2014; 506(7488): 307–15.
 PubMed Abstract | Publisher Full Text
- 129. Jäättelä M: Heat shock proteins as cellular lifeguards. Ann Med. 1999; 31(4): 261–271.
 PubMed Abstract | Publisher Full Text
- Wood RD, Mitchell M, Sgouros J, et al.: Human DNA repair genes. Science. 2001; 291(5507): 1284–1289.
 PubMed Abstract | Publisher Full Text
- 131. Rimer J, Cohen IR, Friedman N: Do all creatures possess an acquired immune system of some sort? *Bioessays*. 2014; 36(3): 273–281. PubMed Abstract | Publisher Full Text
- 132. Vandermeer JH: Niche theory. Annu Rev Ecol Syst. 1972; 3(1): 107–132. Publisher Full Text
- Estrela S, Libby E, Van Cleve J, et al.: Environmentally mediated social dilemmas. Trends Ecol Evol. 2019; 34(1): 6–18.
 PubMed Abstract | Publisher Full Text
- 134. Prüfer K, Munch K, Hellmann I, et al.: The bonobo genome compared with the chimpanzee and human genomes. Nature. 2012; 486(7404): 527–531. PubMed Abstract | Publisher Full Text | Free Full Text
- Odling-Smee FJ, Laland KN, Feldman MW: Niche construction: the neglected process in evolution (MPB-37). 2013; 61: Princeton University Press, Reference Source
- Day RL, Laland KN, Odling-Smee FJ: Rethinking adaptation: the nicheconstruction perspective. Perspect Biol Med. 2003; 46(1): 80–95.
 PubMed Abstract | Publisher Full Text
- 137. Hasson O: Pursuit-deterrent signals: communication between prey and predator. Trends Ecol Evol. 1991; 6(10): 325–329. PubMed Abstract | Publisher Full Text
- Ritchie EG, Johnson CN: Predator interactions, mesopredator release and biodiversity conservation. *Ecol lett.* 2009; 12(9): 982–998.
 PubMed Abstract | Publisher Full Text
- McNally L, Brown SP: Building the microbiome in health and disease: niche construction and social conflict in bacteria. *Philos Trans R Soc Lond B Biol Sci.* 2015; 370(1675): 20140298.
 PubMed Abstract | Publisher Full Text | Free Full Text

- 140. Blaser MJ, Kirschner D: The equilibria that allow bacterial persistence in human hosts. Nature. 2007; 449(7164): 843–849. PubMed Abstract | Publisher Full Text
- 141. Cohen IR: Rain and Resurrection How the Talmud and Science Read the World. CRC Press, 2010. Publisher Full Text
- 142. Tung J, Barreiro LB, Burns MB, et al.: Social networks predict gut microbiome composition in wild baboons. eLife. 2015; 4: e05224. PubMed Abstract | Publisher Full Text | Free Full Text
- Blaser MJ: Missing Microbes: How the Overuse of Antibiotics Is Fueling Our Modern Plagues. Henry Holt and Company, New York, 2014. Reference Source
- De Laubenfels MW: Dinosaur extinction: one more hypothesis. J Paleontol. 1956; 30(1): 207–212. Reference Source
- 145. Gokhman D, Lavi E, Prüfe K, et al.: Reconstructing the DNA methylation maps of the Neandertal and the Denisovan. Science. 2014; 344(6183): 523-527. PubMed Abstract | Publisher Full Text
- Thompson D: What in the world is causing the retail meltdown of 2017? The Atlantic. 2017; 10(04).
 Reference Source
- Lyons SK, Smith FA, Brown JH: Of mice, mastodons and men: humanmediated extinctions on four continents. Evol Ecol Res. 2004; 6(3): 339–358. Reference Source
- Bar-On YM, Phillips R, Milo R: The biomass distribution on earth. Proceedings of the National Academy of Sciences. 2018; 115(25): 6506–6511.
 Publisher Full Text
- McShea DW: Complexity and evolution: what everybody knows. Biol Philos. 1991; 6(3): 303–324.
 Publisher Full Text
- Wolf YI, Koonin EV: Genome reduction as the dominant mode of evolution. Bioessays. 2013; 35(9): 829–837.
 PubMed Abstract | Publisher Full Text | Free Full Text
- 151. Yahalomi D, Atkinson SD, Neuhof M, *et al.*: A cnidarian parasite of salmon (myxozoa: *Henneguya*) lacks a mitochondrial genome. *Proc Natl Acad Sci.* 2020; 117(10): 5358–5363. PubMed Abstract | Publisher Full Text | Free Full Text
- Barnosky AD, Matzke N, Tomiya S, *et al.*: Has the earth's sixth mass extinction already arrived? *Nature*. 2011; 471(7336): 51–7.
 PubMed Abstract | Publisher Full Text
- DeLong EF, Pace NR: Environmental diversity of bacteria and archaea. Syst Biol. 2011; 50(4): 470–478.
 PubMed Abstract | Publisher Full Text
- Bronstein JL, Armbruster WS, Thompson JN: Understanding evolution and the complexity of species interactions using orchids as a model system. New Phytol. 2014; 202(2): 373–375.
 PubMed Abstract | Publisher Full Text
- Quintana FJ, Cohen IR: The HSP60 immune system network. Trends Immunol. 2011; 32(2): 89–95.
 PubMed Abstract | Publisher Full Text
- 156. Gould SJ, Vrba ES: Exaptation—a missing term in the science of form. Paleobiology. 1982; 8(1): 4–15. Publisher Full Text
- Hodgkin J: What does a worm want with 20,000 genes? Genome Biol. 2001; 2(11): comment2008.1-comment2008.4.
 PubMed Abstract | Publisher Full Text | Free Full Text

- Ezkurdia I, Juan D, Rodriguez JM: Multiple evidence strands suggest that there may be as few as 19 000 human protein-coding genes. *Hum Mol Genet.* 2014; 23(22): 5866–5878.
 PubMed Abstract | Publisher Full Text | Free Full Text
- Davidson EH, Erwin DH: Gene regulatory networks and the evolution of animal body plans. Science. 2006; 311(5762): 796–800.
 PubMed Abstract | Publisher Full Text
- Olson MV, Varki A: Sequencing the chimpanzee genome: insights into human evolution and disease. Nat Rev Genet. 2003; 4(1): 20–28. PubMed Abstract | Publisher Full Text
- 161. Lewontin RC: The units of selection. Annu Rev Ecol Syst. 1970; 1(1): 1-18.
- 162. Elisabeth Lloyd: Units and levels of selection. 2005. Reference Source
- 163. Roughgarden J, Gilbert SF, Rosenberg E, et al.: Holobionts as units of selection and a model of their population dynamics and evolution. *Biological Theory*. 2018; **13**(1): 44–65. Publisher Full Text
- 164. Reed ES: The lawfulness of natural selection. Am Nat. 1981; 118(1): 61–71. Publisher Full Text
- Byerly HC: Natural selection as a law: Principles and processes. Am Nat. 1983; 121(5): 739–745.
 Publisher Full Text
- 166. Paul DB: Darwin, social darwinism and eugenics. The Cambridge Companion to Darwin. 2003; 214. Publisher Full Text
- 167. Bergman J: The Darwin Effect: It's influence on Nazism, Eugenics, Racism, Communism, Capitalism & Sexism. New Leaf Publishing Group. 2014. Reference Source
- Mullis KB: The unusual origin of the polymerase chain reaction. Sci Am. 1990; 262(4): 56–65.
 PubMed Abstract | Publisher Full Text
- 169. Efroni S, Harel D, Cohen IR: Toward rigorous comprehension of biological complexity: modeling, execution, and visualization of thymic T-cell maturation. Genome Res. 2003; 13(11): 2485–2497. PubMed Abstract | Publisher Full Text | Free Full Text
- 170. Al-Shahrour F, Minguez P, Tárraga J, et al.: Babelomics: a systems biology perspective in the functional annotation of genome-scale experiments. Nucleic Acids Res. 2006; 34(Web Server issue): W472-W476. PubMed Abstract | Publisher Full Text | Free Full Text
- 171. Turing A: The chemical basis of morphogenesis. Philosophical transactions of the Royal Society of London B. 1952; 237(641): 37–72. Publisher Full Text
- Prigogine I: Time, structure, and fluctuations. Science. 1978; 201(4358): 777–785.
 Publisher Full Text
- Jiménez J: Coherent structures in wall-bounded turbulence. J Fluid Mech. 2018; 842.
 Publisher Full Text
- Horowitz JM, Jeremy LE: Spontaneous fine-tuning to environment in manyspecies chemical reaction networks. *Proc Natl Acad Sci U S A*. 2017; 114(29): 7565–7570.
 PubMed Abstract | Publisher Full Text | Free Full Text
- 175. Pesce D, Lehman N, de Visser JAGM: Sex in a test tube: testing the benefits of in vitro recombination. *Philos Trans R Soc B Biol Sci.* 2016; 371(1706): 20150529. PubMed Abstract | Publisher Full Text | Free Full Text

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Version 2

Reviewer Report 10 August 2020

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Susan Stepney 🔟

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Cohen and Marron discuss a view of evolution in the context of entropy: that the main driver underpinning change and evolution at all scales in living organisms is the need to resist the inevitable decay of that organisation, the need to resist the increase in entropy. They provide extensive examples of the benefits of organisation, how there are no isolated organisms, and how viewing biological systems, from cells to ecosystems, through the lens of interactions allows understanding of the existence of various universals in biology. They conclude by suggesting investigations to test this view.

I find the argument compelling and thought-provoking as it stands. However, there are a few places where I believe links to other work in literature could help further demonstrate both the novelty of the approach, with its focus on evolution specifically, and its connection to other thermodynamic takes on biology.

In particular I think it would be useful to include a little on the work of Kaufmann (constrained release of energy; non-equilibrium flow into the adjacent possible; fourth law of thermodynamics; etc), and of Schneider & Sagan (open systems and non-equilibrium thermodynamics; life reducing energy gradients and thereby *accelerating* the increase in entropy; etc).

- Stuart A. Kauffman. Investigations. Oxford University Press. 2000
- Eric D. Schneider, Dorion Sagan. *Into the Cool: energy flow, thermodynamics, and life*. University of Chicago Press. 2005

Also, prediction #2 would benefit from a comparison with Lenski's long term evolution experiment.

There are also places where the terminology and argument could be tightened up somewhat:

1. p3: although there are clues in the subsequent text, it might be useful to add a couple of sentences saying explicitly why the evolution of unicellular organisms is not dealt with.

2. p4: the statement of the 2nd law would better read: *disorder in an isolated system does not*

decrease. (In thermodynamics, a *closed* system can still exchange energy with its surroundings, eg a constant temperature heat bath, whereas an *isolated* system cannot. Also, entropy does not continue to increase past thermodynamic equilibrium.)

- 3. p5, col2: "everything made of matter is made of ... atoms"; this is true only under certain physical conditions: the sun, for example, is a plasma; a neutron star is made of degenerate matter.
- 4. p8, col2: "Bonding between reacting entities reduces ... the total amount of entropy." It reduces the entropy of the atoms, but does not reduce in total: the rest is carried away by energy released by the exothermic reaction and absorbed by the constant temperature heat bath.
- 5. p9, col1: "define a living entity as one that is able to exploit energy, by way of metabolism, to renew itself." The autopoiesis work of Maturana and Varela could be referenced here.
- 6. p13, col1: deer, barn owls, etc, co-existing with human environments, are called "feral" creatures. Feral rather means "living wild, having escaped from captivity or domestication" (as with the Mynah birds on p14); here a better term would be "synanthropic".
- 7. p13, co1: all the examples of animals constructing niches: bird nests fit here too.
- 8. p13, col1: "each species interacts in its own unique way with that space": this is reminiscent of the William James quotation: "Other sculptors, other statues from the same stone! Other minds, other worlds from the same monotonous and inexpressive chaos!
- 9. p14, col2: on the biomass of domesticated species, the point could be made even more dramatically by saying: "the biomass of livestock outweighs by many fold the biomass of all wild mammals combined, and even outweighs the biomass of humans". If "survival of the fittest" holds, cattle are fitter than humans!
- 10. p17, prediction 6: "provided that entropy holds sway there"; why would entropy *not* hold sway there? All the arguments in this paper assume 2nd Law inescapability.

There are also a couple of typos.

1. p3, revision box, point 4: Szathmary's name is misspelled.

2. p15, line 1: repeated "by".

Is the topic of the opinion article discussed accurately in the context of the current literature?

Yes

Are all factual statements correct and adequately supported by citations?

Yes

Are arguments sufficiently supported by evidence from the published literature?

Yes

Are the conclusions drawn balanced and justified on the basis of the presented arguments? $\ensuremath{\mathsf{Yes}}$

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: Complex systems, artificial life

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard.

Author Response 25 Aug 2020

Assaf Marron, Weizmann Institute of Science, Rehovot, Israel

We thank Susan Stepney for her expert review, positive comments, and valuable suggestions, and for drawing our attention to additional relevant publications.

Version 3 is now amended to address many of the comments and suggestions. Specifically:1. In the introduction, we stated that we plan to address the mechanism of unicellular evolution in a future paper.

2. In the section "Entropy is life's nemesis and evolution's facilitator", we edited our statement of the second law of thermodynamics, and added published recitations of two other brief statements of the law, all of which address the comment that entropy does not always increase.

3. In the above section, we added to the list of papers exploring the connections between evolution and entropy citations of "Investigations" by Kauffman and "Into the cool: Energy flow, thermodynamics, and life" by Schneider and Sagan. We have elaborated on the comparison with Kauffman: In the section "Analysis and modeling," A key point in this comparison is the fact that our proposed evolutionary mechanism of entropic selection of holobionts and cooperative interactions for fittedness does not require the concepts of autonomous agents, fitness landscapes, and gene-centered competitive success. Our nascent models are likely to differ from those considered in the past.

4. In the section "Interactions", we reordered and rephrased the sentences to clarify and add context to Feynman's quote that "everything is made of atoms".

5. In the section "programmed renewal," we cited the work of Maturana and Varela on autopoiesis and the definition of living systems.

6. In the section "Metabolism protects against entropic dissolution," we added citations to Kauffman's discussion of the constrained release of energy.

7. In the above section, we clarified that bonding releases entropy into the environment as heat.

8. In the section "Humans are responsible for widespread niche deconstruction", we explicitly included humans in the discussion of animal biomass.

9. We kept our use of the word "feral" in the sense of "not domesticated", in line with the main definition of the OED and meaning 2a in Merriam-Webster (though both also include the narrower meaning offered in the review).

10. We agree with Susan Stepney's association to William James inspiring quotation, but decided not to add it to the paper.

11. We kept the cautious conditional regarding planets in other galaxies "provided that

entropy holds sway there"12. We corrected several typos.

We thank Susan Stepney again for the detailed and valuable review. Irun R. Cohen and Assaf Marron

Competing Interests: No competing interests were disclosed.

Reviewer Report 07 August 2020

https://doi.org/10.5256/f1000research.28164.r68397

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Eugene V. Koonin 匝

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I appreciate Cohen's and Marron's response to my review of this paper. The major issues that I raised have been addressed, in the least, by siting the appropriate literature. We still do not see eye to eye with regard to 'survival of the fitted' vs 'survival of the fittest', where I fail to recognize a substantial distinction. However, the authors are entitled to their position.

Is the topic of the opinion article discussed accurately in the context of the current literature?

Yes

Are all factual statements correct and adequately supported by citations?

Yes

Are arguments sufficiently supported by evidence from the published literature? $\ensuremath{\mathsf{Yes}}$

Are the conclusions drawn balanced and justified on the basis of the presented arguments? $\ensuremath{\mathsf{Yes}}$

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: Evolutionary biology

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard.

Version 1

Reviewer Report 13 July 2020

https://doi.org/10.5256/f1000research.26968.r65123

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了 🛛 Eugene V. Koonin 匝

National Center for Biotechnology Information (NCBI), National Library of Medicine (NLM), National Institutes of Health (NIH), Bethesda, MD, USA

Cohen and Marron present a concept of evolution that centers on interactions on different level molecular, cellular, organismal etc. - which are perceived as a universal anti-entropic force. Entropy is understood as a tendency for increasing disorder and is viewed as the principal selective factor in evolution. The antientropic evolution is postulated to be 'universal', hence no neutral evolutionary processes. The authors, then, forward the principle of 'survival of the fitted' whereby all biological entities that counter entropy with sufficient success to remain in an ordered state survive.

Although not exactly new (indeed, it goes back to Boltzmann who the authors duly cite!), the notion of the fundamental nature of the antientropic tendencies in evolution is worth putting forward. More specifically, the emphasis on interactions permeating the entire biological organization as a universal antientropic factor is worthy and might appeal to many biologists as a fresh perspective eon the general character of evolution.

However, I find several areas within and around these where concepts where I feel compelled to mount criticisms, and not altogether minor ones.

First, to me, the 'survival of the fitted' appears to be a non sequitur. Let us say, everything is fine with entropy as a selective factor and interactions, but why not survival of the fittest, that is, those that counteract entropy most efficiently and, as a result, leave more progeny than others?

Second, in all their discussion of entropy, the authors refer only to classical,equilibirum thermodynamics. As far as I can see, this discussion cannot be complete without taking into account the concepts of non-equilibrium thermodynamics, in particular, maximization of entropy production and dissipative structures. The literature on these subjects is extensive, starting with Prigogine.

Third, to me, it is inexplicable how and why, when discussing the pivotal roles of interactions at different levels in evolution, the authors fail to include Maynard Smith-Szathmary's concept of major transitions in evolution (Ref. 1¹ and references therein) and, especially, the generalized form of this concept that comes uncannily close to the concept of interactions-driven evolution discussed here².

Fourth, I believe that, along with the key role of interaction, it is essential for developing a general concept of evolution, to consider competition between interactions on different scales and the resulting frustrated states².

Finally, more with respect to the presentation of the material. The figures in the current manuscript are over-simplified and, overall, underwhelming. Figure 1 is a trivial illustration of the concept of entropy and is superfluous (readers unfamiliar with the second law will not benefit from reading this article) whereas for Figures 2 and 3, more elaborate schemes are advisable.

References

1. Szathmáry E: Toward major evolutionary transitions theory 2.0.*Proc Natl Acad Sci U S A*. 2015; **112** (33): 10104-11 PubMed Abstract | Publisher Full Text

2. Wolf Y, Katsnelson M, Koonin E: Physical foundations of biological complexity. *Proceedings of the National Academy of Sciences*. 2018; **115** (37): E8678-E8687 Publisher Full Text

Is the topic of the opinion article discussed accurately in the context of the current literature?

No

Are all factual statements correct and adequately supported by citations?

No

Are arguments sufficiently supported by evidence from the published literature? No

Are the conclusions drawn balanced and justified on the basis of the presented arguments? Partly

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: Evolutionary biology

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard, however I have significant reservations, as outlined above.

Author Response 21 Jul 2020

Assaf Marron, Weizmann Institute of Science, Rehovot, Israel

We appreciate Eugene Koonin's positive opening remarks and his expert review and suggestions for improving the paper.

The specific recommendations for improvement are five:

First, "everything is fine with entropy as a selective factor...but why not survival of the fittest, that is, those that counteract entropy most efficiently and, as a result, leave more progeny than others?"

This is an important question and we have addressed it directly in the revision. Briefly, we reason that dealing with entropy does not directly require competition and struggle; one need only to fit in "sufficiently"; moreover, variants that are not "the fittest" readily find other niches, and many advantageous phenotypes are not necessarily associated with reproductive advantage; indeed, increased reproduction is at times more of a problem than a desired outcome. This is, of course, in addition to the distinction the paper makes between universal adaptations driven by entropic selection and local adaptations, which may well be based on reproductive advantage and Darwinian natural selection. See the section titled "Reconciling survival-of-the-fitted with Darwinian survival-of-(only)-the-fittest".

Second, "the authors refer only to classical, equilibrium thermodynamics." We appreciate the reference to the work of Prigogine and others on the emergence of order in multiple scales from chaotic thermodynamic behavior both in equilibrium and non-equilibrium states. In our revision, we have added the suggested references and others as important elements in quantitative analysis, simulation, and prediction. See the section titled "Analysis and modeling".

We thank Eugene Koonin for calling our attention to these ideas.

Third, the first draft of the paper neglected to cite the concepts of Maynard Smith and Szathmary on Major Transitions in Evolution. Our revision now refers to this work which, among others, highlights the pivotal roles of interaction and the relentless increase of complexity. See subsection "Cooperation retards entropy" in section "Interactions".

Fourth, the Reviewer suggested that the paper should "consider competition between interactions on different scales and the resulting frustrated states" as a formative element in cooperative interaction. In the revised paper, we now incorporate this idea and cite the relevant paper by Wolf, Katznelson and Koonin. See subsection "Cooperation retards entropy" in section "Interactions".

We also added references to Maynard Smith & Szathmary and to Wolf, Katznelson and Koonin in the section titled "the emergence of complexity".

Fifth and finally, the Reviewer suggested that the Figures are over-simplified and superfluous (Figure 1) and "for Figures 2 and 3, more elaborate schemes are advisable." In the revised paper, the captions clarify that the Figures are meant to serve only as simplified conceptual anchors or mnemonics for the principles of the theory.

We thank Eugene Koonin for raising these points which we feel have improved the paper

considerably. Irun R. Cohen and Assaf Marron

Competing Interests: No competing interests were disclosed.

Reviewer Report 23 June 2020

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Eugene Rosenberg

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Cohen and Marron offer a highly original concept of evolution, a subject fundamental to biology and philosophy. They begin by providing a clear and well-written description of neo-Darwinism, recent extended theories of evolution, which take into account epigenetics and microbiomes, and entropy. They then posit their main point- that the evolution of complexity is driven by cooperation, which is a mechanism to avoid entropic destruction. They argue that this cooperation is required at all biological levels, from molecules to ecosystems. For example, they point out the cooperation of individual strands of DNA to form double-stranded DNA helps avoid entropic destruction.

Like most revolutionary ideas in science, the evolution of universal adaptations of life driven by the properties of matter: energy, entropy, and interactions, will be difficult for evolutionary biologists to accept. However, this article should be read and seriously considered by biologists. The authors end the article by suggesting some interesting experiments to test their concept of evolution.

Is the topic of the opinion article discussed accurately in the context of the current literature?

Yes

Are all factual statements correct and adequately supported by citations?

Yes

Are arguments sufficiently supported by evidence from the published literature? Yes

Are the conclusions drawn balanced and justified on the basis of the presented arguments? Yes

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Competing Interests: No competing interests were disclosed.

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard.

Author Response 21 Jul 2020

Assaf Marron, Weizmann Institute of Science, Rehovot, Israel

We are gratified to see that Eugene Rosenberg judges the ideas presented in the paper to be original and fundamental, and that the proposed experiments are interesting and relevant.

We thank Eugene Rosenberg for this expert review.

Irun R. Cohen and Assaf Marron

Competing Interests: No competing interests were disclosed.

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