



First principles in the life sciences: the free-energy principle, organicism, and mechanism

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Received: 20 November 2017 / Accepted: 3 September 2018 / Published online: 10 September 2018
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Abstract

The free-energy principle states that all systems that minimize their free energy resist a tendency to physical disintegration. Originally proposed to account for perception, learning, and action, the free-energy principle has been applied to the evolution, development, morphology, anatomy and function of the brain, and has been called a *postulate*, an *unfalsifiable principle*, a *natural law*, and an *imperative*. While it might afford a theoretical foundation for understanding the relationship between environment, life, and mind, its epistemic status is unclear. Also unclear is how the free-energy principle relates to prominent theoretical approaches to life science phenomena, such as organicism and mechanism. This paper clarifies both issues, and identifies limits and prospects for the free-energy principle as a first principle in the life sciences.

Keywords Adaptation · Free energy · Life · Mechanism · Organicism

1 Introduction

According to the free-energy principle (FEP), all systems that minimize their free energy resist a tendency to physical disintegration. Originally proposed to explain how sensory cortex infers the causes of its inputs and learns causal regularities, FEP has been used to elucidate the function of action, perception, and attention, and to account for organisms' evolution and development (Friston 2003, 2009, 2010a, 2013; Friston et al. 2006; Friston and Stephan 2007).

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Advocates have claimed FEP offers a “framework within which to explain the constitutive coupling of the brain to the body and the environment,” which provides “a normative, teleological essence to the synthesis of biology and information,” and which may illuminate the continuity between life and mind (Allen and Friston 2018, p. 2476).

Advocates also present FEP as a “mandatory principle” or “imperative” for biological systems, and as a principle enjoying a “fundamental status” in neuroscience (Friston et al. 2006, p. 71; Friston and Stephan 2007). The principle purportedly “applies to any biological system [...] from single-cell organisms to social networks” (Friston 2009, p. 293).

Although these dramatic ascriptions have attracted attention in philosophy and the life sciences, FEP’s epistemic status remains opaque, along with its exact role in biological and neuroscientific theorizing. Conspiring against its accessibility are the varying formalisms and formulations of FEP, the changing scope of application, reliance on undefined terms and stipulative definitions, and the lack of clarity in the logical structure of the reasoning leading to FEP.

Regarding its role in theorizing, FEP conflicts with tenets of two of the most prominent contemporary theoretical approaches in the life sciences: organicism (Gilbert and Sarkar 2000; Soto et al. 2016) and mechanism (Brandon 1984; Bechtel and Richardson 1993/2010). Regarding its epistemic status, one worry is that FEP lacks explanatory power because it “is divorced from the biophysical reality of the nervous system” (Fiorillo 2010, p. 605). The principle’s unifying power has been called into question, too. Some have suggested that FEP provides an implausible model of the functional roles of perception and action (Gershman and Daw 2012; Colombo 2017; Klein 2018); others have argued that FEP doesn’t deliver a grand unifying theory, and a plurality of modeling approaches is preferable for explaining complex neurophysiological and cognitive phenomena (Marblestone et al. 2016; Colombo and Wright 2017). Also controversial is how FEP illuminates the continuity between life and mind, synthesizing biology and information—especially if FEP is committed to some form of cognitivism (Hohwy 2016; Kirchhoff and Froese 2017).

According to organicists (aka ‘holists’), phenomena studied in the life sciences should be explained by appealing to whole organisms. Organicists appeal to the principles of biological autonomy and adaptivity (Varela et al. 1974; Di Paolo and Thompson 2014; Moreno and Mossio 2015), organization as closure of constraints (Maturana 1975; Mossio et al. 2016), and variation as extended criticality (Montévil et al. 2016). While FEP borrows the formalism of random dynamical systems to explain the behavior and nature of organisms, organicists maintain that the formalisms borrowed from current theories in physics are not apt for the representation of life science phenomena (Longo et al. 2012).

According to mechanists, phenomena studied in the life sciences should be explained by appealing to the component parts and operations of mechanisms, where a mechanism is a spatiotemporally-organized composite system producing a phenomenon (Bechtel and Richardson 1993/2010; Darden 2006). While FEP is used to logically derive claims about structural and functional properties of the brain “a priori, on the basis of purely theoretical considerations” (Friston 2003, p. 1325), mechanistic

philosophers argue that laws and logical deduction do not adequately capture explanation in the life sciences (Glennan 2017).

In this paper, we clarify the epistemic status of FEP by answering two sets of questions. First: what are the inferential steps leading to FEP? And how do they relate to one another? Second: what's the relationship between FEP and central tenets of the organicist and mechanistic philosophies when it comes to adequate scientific representation of the phenomena of life?

To answer these questions, we carefully reconstruct the reasoning leading to FEP (Sect. 2) and then deploy a contrastive strategy to bring out salient assumptions of FEP. We argue that FEP is inconsistent with a tenet of organicism, specifically that the formalism and concepts of current physical theories are not apt for the scientific representation of organisms and their behaviors (Sect. 3). We also show that FEP is inconsistent with mechanistic approaches in the life sciences, which eschew laws and theories and take the explanatory power of scientific representations to be dependent on the degree of relevant biophysical detail they include. After we clarify the status of FEP as a first principle (Sect. 4), we conclude by suggesting that understanding phenomena in the life sciences should allow for incompatible approaches with varying degrees of idealization (Sect. 5). In particular, the axiomatic, idealizing, first-principles approach favored by free-energy theorists should be pursued in the life sciences alongside the synthetic approaches favored by organicists, and the analytic approaches favored by mechanists.

2 The transcendental argument

Free-energy theorists formulate FEP in various ways. Depending on the exact formulation, the scope of application of FEP varies. One formulation has FEP focused on brains, aiming to clarify the functional significance of their activity and structural connectivity (Friston 2003, 2009; Friston et al. 2006). Another has FEP ranging over any biological entity, process, or complex system—including brainless organisms like single cells and plants, evolutionary processes by natural selection, and ecosystems—and aiming to explain how biological systems can maintain their physical integrity in a changing environment (Friston 2013; Hobson and Friston 2016). In yet another formulation, FEP applies to any complex adaptive system, including physical information systems and non-biological systems like social networks and artefacts (Friston 2009, p. 293).

Under any formulation, the reasoning leading to FEP has the form of a transcendental argument for the conclusion that FEP is a condition on the very possibility of existence of adaptive systems. At its most general, the argument proposes that adaptive systems would be impossible unless FEP were true. In Friston's words: "systems that do not minimize free energy cannot exist" (Friston 2013, p. 2). While asserted to be a condition of possibility on systems' existence, FEP is offered to help leverage 'that-actually' explanations of biological systems' capacity for maintaining their physical integrity by behaving adaptively.

The transcendental deduction of FEP involves six main steps organized in two batches:

1. If a system Σ acts selectively on the environment to avoid phase transitions and is in a non-equilibrium steady-state, then Σ behaves adaptively.¹
 2. Σ behaves adaptively only if Σ preserves its physical integrity by maintaining its “characteristic” variables within homeostatic bounds despite environmental fluctuations.²
 3. Σ acts selectively on the environment to avoid phase transitions and is in a non-equilibrium steady-state just in case Σ preserves its physical integrity by maintaining its “characteristic” variables within homeostatic bounds despite environmental fluctuations.
 4. Σ preserves its physical integrity by maintaining its “characteristic” variables within homeostatic bounds despite environmental fluctuations just in case Σ places an upper bound on the informational entropy (*average surprise*) of its possible sensory states.
 5. If Σ minimizes the free energy of its possible sensory states, then Σ places an upper bound on the informational entropy of its possible sensory states.³
 6. Any system Σ that places an upper bound on the informational entropy of its possible sensory states will preserve its physical integrity by maintaining its “characteristic” variables within homeostatic bounds despite environmental fluctuations.
- ∴ 7. Any system Σ that minimizes the free energy of its possible sensory states will preserve its physical integrity by maintaining its “characteristic” variables within homeostatic bounds despite environmental fluctuations.

The first three claims comprise the first batch, where (1) states a thermodynamically sufficient condition on adaptive behavior, (2) states a homeostatically necessary condition, and (3) functions as a bridge principle that connects physical and biological predicates expressing properties of a system’s behaving *adaptively*. The bridge is derivable from the conjunction of its left-to-right conditional, which follows directly from (1) to (2), and the stipulation of its suppressed right-to-left converse. The next three claims comprise the second batch, where (4) stipulatively connects homeostasis to information-theoretic quantities, and (5) states that free-energy minimization is a sufficient condition for placing an upper bound on the informational entropy of the

¹ As explained in Sect. 2.1, *equilibrium* refers to thermodynamic equilibrium. A system is in a steady state if the variables defining the behavior of the system are unchanging with time. An equilibrium state is a special case of a steady state. (A system in equilibrium is one in a steady state; the converse isn’t necessarily true.)

² Values of these characteristic variables would be observable properties of “the extended phenotype of the organism—its morphology, physiology, behavioural patterns, cultural patterns, and designer environments,” and would be phase functions of an attracting set of the organism’s states (Ramstead et al. 2018, p. 3). The idea is that any living system possesses an ‘attracting set’, i.e., a set of states towards which it will tend to evolve, for a wide variety of values of its initial states, and “this set of attracting states can be interpreted as the extended phenotype of the organism” (ibid). Free-energy theorists’ transcendental deduction assumes that an attracting set in a system open to its external milieu is formally equivalent to a steady-state solution that is far from equilibrium.

³ According to some formulations, any system that conserves its boundaries “can be described as” modeling its external milieu or “can be cast as” minimizing free energy (Hobson and Friston 2016, p. 246; Ramstead et al. 2018, p. 2). In other formulations, the system itself is said to be the modeler (Friston 2011). We address this confusion in Sect. 3.

sensory states of a system.⁴ Claim (6) expresses a sufficient condition for a system's preserving its physical integrity. Claim (7) expresses FEP, and follows from the two batches of claims.

The modal force of FEP, as well as its claim to being an a priori first principle, depend on free-energy theorists' stipulations about the semantic equivalence between concepts from different disciplines, like the equivalences between the concepts of *adaptive value*, *surprisal*, and *model evidence*, and on their interpretation of probability. We'll consider their understanding of adaptive living systems in Sect. 2.2, and put into focus the other assumptions underlying their reasoning in Sect. 3.

2.1 Thermodynamics

The argument for FEP begins with the observation that some systems maintain their physical integrity, displaying adaptive behavior amid a changing environment. Such systems are thermodynamically open, persisting far from their thermodynamic equilibrium state (Friston et al. 2006, pp. 71–72; Friston and Stephan 2007, pp. 421–422). Systems that are thermodynamically open exchange matter and energy with their surroundings. For example, both snowflakes and bacteria exchange chemicals and energy with their surroundings. Snowflakes acquire and lose matter and heat under the causal pressures of their environment, and bacteria allow matter and energy to cross their cytoplasmic membranes. Closed systems have boundaries that matter from their surroundings cannot cross. Earth, for example, exchanges energy with its surroundings, but hardly any matter. Isolated systems exchange neither energy nor matter with their surroundings. The universe and a closed thermos bottle are examples of isolated systems.

⁴ Premise 5 should be treated with care. First, in machine learning, free energy is a variational bound on informational entropy that often plays a central role in variational Bayesian inferences. Where X is an observed variable (e.g., a sensory sample) and Z a hidden random variable of interest (e.g., an environmental cause of a sensory sample), the idea behind variational Bayesian inference is to find some approximation distribution density $q(Z)$ that is tractable and that is as close as possible to the true posterior distribution $p(Z|X)$. As the approximation distribution can have its own parameters $q(Z|\theta)$, the problem of inferring the value of Z given the value of X can be understood as an optimization problem, in which one aims to find the parameter values that minimize some objective function and make q as close as possible to the posterior of interest. To measure the closeness of the two distributions $q(Z)$ and $p(Z|X)$, a common metric is the Kullback–Leibler divergence. Second, free-energy theorists argue that organisms cannot minimize the informational entropy of their sensory states (i.e., 'surprisal') directly, because this involves the intractable problem of computing marginal probabilities. Instead, they argue that organisms can tractably minimize the informational entropy of their sensory states indirectly, by minimizing a bound called 'variational free energy'. As Buckley et al.'s (2017) helpful mathematical review makes clear, this claim requires qualification, since variational free energy is a tight bound on surprisal only when the organism's current 'best guess' of the causes of its sensory input (represented by a 'recognition density function' $q(Z)$) is identical with the posterior density of environmental states given the organism's sensory input $p(Z|X)$. Finding an optimal recognition density that is identical with the posterior is non-trivial, and requires further assumptions about its form, and about the form of the dynamics in the environment. "Furthermore, while this process furnishes the organism with an approximation of surprisal it does not minimise it. Instead the organism can minimise variational free energy (VFE) further by minimising surprisal indirectly by acting on the environment and changing sensory input" (Buckley et al. 2017, p. 59). We'll return on both these points in Sect. 2.4. We are grateful to anonymous reviewers, including a key architect of FEP, for pressing us to clarify this reconstruction.

Rather than properties of Σ —like being thermodynamically open, or being in equilibrium—it may be useful to speak of (sets of) states. The term *state* here refers to a property instance at a time. Velocities, for example, are states: different objects can have different velocities over time but only a single velocity at any given time. A system's states are specified by the values of sets of measurable macroscopic variables, including geometric, thermal, mechanical, and chemical properties at a given time scale, and can typically be associated with mathematical representations, e.g., vectors in a vector- or state-space. Such variables are related to one another in law-like ways, and so provide a way to deduce changes in the system's behavior over time.

When systems are left to themselves and external conditions are unchanging, their states change until there are no net flows of matter or energy either within the system or between it and its surroundings. This state is thermodynamic equilibrium. Although most systems found in nature aren't in equilibrium, there are important differences across systems. In comparison to snowflakes left to themselves, bacteria are far from their equilibrium state: while their chemical and metabolic properties change, the bacteria can acquire energy and nutrients, and maintain a non-equilibrium steady state to avoid thermodynamic equilibrium.

When systems change states, they undergo a process. The succession of a system's states defines the process's path or trajectory. Some processes involve abrupt discontinuous transitions between solid, liquid, or gaseous states of matter. These transitions are called *phase transitions*. When systems undergo phase transitions, some of their physical properties change—often resulting from changes in temperature, pressure, or other surrounding conditions. For example, snowflakes will phase transition into water droplets when external temperature reaches its melting point. Many biological systems also undergo phase transitions, where their morphological and metabolic profiles change dramatically. These transitions depend on phenotypic plasticity, which is the ability of biological systems with one genotype to change their phenotype in response to changing environmental conditions. Examples include temperature-dependent sex determination in some fish and reptiles, eye-spot formations in some butterflies, and insect metamorphosis.

Because of the second law of thermodynamics, the path of the process undergone by a system isolated from all external influences eventually goes into a state of thermodynamic equilibrium. There are several formulations of the law (Uffink 2001). One formulation refers to thermodynamic entropy, which is a state variable measuring the amount of “disorder” (or randomness) in a system, and implies that entropies of isolated systems not in equilibrium typically increase over time, approaching maximum value at equilibrium. It is a function of a system's state, and captured by the formula: $S = k \log(W)$. This formula describes the entropy S of a system in terms of the logarithm of the number of possible microstates $W = \{w_1, w_2, \dots, w_n\}$ that are consistent with the macroscopic states of the system, where k is the Boltzmann constant.

Comparing snowflakes with water vapor offers one intuitive way of visualizing the notion of “disorder” associated with thermodynamic entropy. Water vapor in a container can have many possible arrangements of individual molecules consistent with the macroscopic properties of the gas like its volume and pressure. Because snowflakes' molecules are constrained by crystalline bonds, the number of possible configurations of individual molecules consistent with the macroscopic properties of

snowflakes is smaller. Snowflakes have less entropy than water vapor: they are less “disordered” (random).

2.2 From thermodynamics to homeostasis

Free-energy theorists claim that biological systems apparently “resist” or “violate” the second law of thermodynamics (or more accurately, the fluctuation theorem) because they maintain their physical integrity in the face of random fluctuations in the environment (see e.g., Friston and Stephan 2007, pp. 421–422; Ramstead et al. 2018, 2 ff.).

Biological systems are open systems that maintain order (thermodynamic entropy) by exchanging energy and matter with their surroundings. Upon considering all such exchanges, the total entropy of the system and its environment increases over time in ways that can be described by thermodynamical laws. It’s this capacity for “negative entropy”—acting selectively upon their environments and metabolizing food, which distinguishes living from non-living systems (Collier 1986; Schrödinger 1992; Collier and Hooker 1999; Morowitz and Smith 2007; Bailly and Longo 2009).

A system’s capacity to act selectively upon its environment enables the system to maintain its physical integrity amid changing environmental conditions. Friston and Stephan (2007) illustrate this point with a fictitious example. They compare regular snowflakes with winged snowflakes that can act on their environment. Regular snowflakes are passively pushed around by environmental forces until their temperatures reach a certain threshold, where they undergo phase-transitions, losing their integrity and turning into water droplets. Because winged snowflakes can fly and keep certain altitudes, they can maintain their temperature within bounds and away from their melting point. Maintaining their temperature within certain bounds is a necessary condition for the winged snowflakes to keep a non-equilibrium steady state so that they may avoid a phase transition and disintegration. So, winged snowflakes can maintain a “relatively constant milieu” despite environmental changes—that is, maintain homeostasis—via behaviors that adjust to new or changing conditions while maintaining their macroscopic properties within bounds. Because their behaviors enable winged snowflakes to maintain homeostasis, those behaviors, unlike regular snowflakes, are said to be adaptive.

We have two conditions on a system’s adaptive behavior. One is in terms of thermodynamics: if systems act selectively upon their environments to preclude phase transitions and stay away from thermodynamic equilibrium, then they behave adaptively. Another is in terms of homeostasis: if systems behave adaptively, then they change their relationships with their environments to maintain vital physiological variables within certain bounds.

Free-energy theorists link adaptivity directly to viability, understood in terms of homeostasis, similarly to Ashby’s (1960, p. 58) account of adaptation as ultrastability did. For free-energy theorists, characteristics for phase-dependent measurement function—or as Ashby called them, “essential variables”—must be kept within viable limits to prevent the system from dying rapidly or disintegrating from phase transi-

tions. Like Ashby (1960), free-energy theorists relate thermodynamic and biological formulations of adaptive processes.

Specifically, free-energy theorists assume that any living system possesses a random dynamical attractor—a set of states towards which a dynamical system tends to evolve for a wide variety of initial conditions of the system’s state. This attractive set is interpreted as the system’s extended phenotype, which includes characteristics defining a kind of biological system. Under appropriate conditions, any system possessing a random dynamical attractor can be shown to be formally equivalent to any system at a steady state far from equilibrium, where the system’s “characteristic” variables are within homeostatic bounds (Friston 2012; Ramstead et al. 2018, p. 2).⁵ In other words, the paths of the processes of adaptive (living) systems fall within a specific, relatively narrow region of all possible states in their phase space. For Friston, no less than Ashby, survival is equivalent to the system’s being in that narrow region.

With these physical and biological formulations articulated, the argument’s next step is to deploy the mathematics of random dynamical systems theory and information theory to answer the following question. What characteristics must biological systems possess to maintain their path within a specific (homeostatic) region that precludes phase transitions?

2.3 From homeostasis to surprise

Physical systems can be represented as sets of variables. Different values of these variables pick out different states of the systems. The set of all possible states of a system can be represented as a state space, which allows one to describe the system and its changes in time. A phase space is a continuous state space described with a smooth manifold. A space’s dimensionality depends on how many variables are needed to describe the target system and its dynamics. Each state of the system is represented with a point in the state space. Given the state of the system at any moment, one can use an evolution rule, which can be either deterministic or stochastic, to describe the next states of the system in state space.

Friston (2012, 2013) represents biological systems as random dynamical systems, with state spaces partitioned into external and internal states. External states correspond to environmental causes that generate sensory samples (also known as *sensory input*, *sensory outcomes*, *sensory data*, or *evidence*), which affect the system’s internal state. A subset of states (known as *Markov blanket*—more on this momentarily) can ground a separation between the internal states of the system and the external states of the environment. A further subset of these separating (blanket) states is distinguished as its so-called *active states*, where different values of an active state determine different positions of the system in the environment.

The justification for partitioning these states appeals to the construct MARKOV BLANKET (Friston 2013). Pearl (1988) introduced this concept in relation to Markov

⁵ Equating adaptation and viability renders Ashby’s concept of essential variables unclear. If essential, in what sense can its values be transgressed without causing death? Or, if such transgression is possible, then in what sense are they essential? The construct ALLOSTASIS may offer, here, a way to understand how organisms can operate outside of normal set-points and approximate equilibria values, without leading to the total cessation of all physiological function.

networks for representing probabilistic knowledge. Roughly, given a set of random variables N , the Markov blanket for a variable $x \in N$ is the subset M containing all random variables that “shield” x from all the other variables in N . Fixing the values of the variables in M leaves x conditionally independent of all other random variables; hence, the Markov blanket of a random variable is the only knowledge one may need to predict the behavior of that variable (Pearl 1988, 97 ff.).

In machine learning, Markov blankets help address problems with constructing sets of causal models from sample data that are as small as possible, and help search these sets to find a true causal model (Spirtes et al. 2000). Friston (2013), however, supposes further that Markov blankets are objective features of the real world separating the states internal to biological systems from those external to them (more on this reification in Sect. 3). His mathematical representation of system’s exchanges with their environments involves four basic types of quantities (for a helpful review of the mathematical details, see Buckley et al. 2017):

1. external states $\Psi = \{\psi_1, \dots, \psi_n\}$ standing in for the environmental causes of sensory states;
2. active states $A = \{a_1, \dots, a_n\}$ that change what external states the system samples;
3. sensory states (or samples) $D = \{d_1, \dots, d_n\}$ that depend upon active and internal states of the system;
4. a generative model M defined in terms of its parameters or sufficient statistics, and would be “encoded” by the system’s internal states.

For biological systems, free-energy theorists assume that an organism’s phenotype, defined as “the repertoire of physiological and sensory states in which an organism can be” (Friston 2010a, p. 127), is constituted by the organism’s internal states and its Markov blanket, which comprises active and sensory states.

Free-energy theorists represent the organism’s internal states with the parameters (or sufficient statistics) of a posterior probability distribution over external states generating sensory samples (Friston and Stephan 2007, p. 424). The parameters would be physically realized as weights and biases of networks in the organism’s brain, and are optimized—where optimization consists in finding values for these parameters that minimize or maximize an objective function—in relation to a generative model.

Biological systems’ phenotypes are said to “entail” the generative model (Friston 2012), where *entailment* here means that external states can be (inductively) inferred on the basis of sensory states, and that this inference implicitly assumes a generative model (i.e., a probabilistic specification of how sensory states are generated by external states). Given how free-energy theorists define *phenotype*, the generative model M provides the probability that a certain kind of system obtains any possible state in state space. In short, according to free-energy theorists, for any phenotype, there is a generative model M that renders the internal states of the phenotype as the sufficient statistics of posterior densities of external states under M .

In the winged snowflake example, external states include ambient temperature, wind direction, and other environmental factors that generate sensory samples (or inputs) influencing the snowflake’s internal state. Internal states include its temperature and the strength of its microcrystals’ electrostatic bonds, but also active states like its local position in the environment at a time. The snowflake’s active state includes its position

in the environment, so that changing this active state will lead to a change in position; and, by changing position in the environment, the snowflake receives different sensory samples (or experiences different sensory states). Given the kind of system a winged snowflake happens to be, it's improbable that states where its temperature is higher than $0\text{ }^{\circ}\text{C}$ will obtain.

With respect to cells, external states include ambient temperature and pH. These states generate sensory samples (i.e., energy arrays impinging on organisms' sensory surfaces) that influence the state of transmembrane receptors. The cell's internal states include the concentration of intracellular metabolites, but also active states, like the motion of flagella at times, that change how the cell's environment influences the cell's receptors. Given the kind of system a cell happens to be, it's improbable that states where its temperature is higher than $50\text{ }^{\circ}\text{C}$ will obtain.

Given the four basic quantities (for external, sensory, active, and internal states of a biological system) and their relationships of probabilistic conditional (in)dependence, one can examine what characteristics a random dynamical system must possess so that its physical states “are confined to a bounded subset of states and remain there indefinitely” (Friston 2012, p. 2106). This subset of states in state space is a random dynamical attractor corresponding to a set of (non-equilibrium) steady states, where levels of variables representing the sensory states of the system—like temperature, pH, glucose, blood oxygenation, etc.—are within homeostatic bounds. To learn what characteristics biological systems must possess if they are to maintain their path within a specific (homeostatic) region that precludes phase transitions, free-energy theorists ask what dynamics such systems must exhibit for a random dynamical attractor to obtain.

If one then stipulates that “biological systems move around in their state space, but revisit a limited number of states” that correspond to homeostatic steady states far from equilibrium (Friston 2013, p. 11), then, of all possible obtaining states, there's a small number that they will achieve in their lifetime with a high probability. All other states will be obtained with an exceedingly low probability.

To capture this idea formally, free-energy theorists use the information-theoretic concept SURPRISE. The average surprise of sampling some outcome corresponds to Shannon's entropy, which is formally similar to the thermodynamic concept ENTROPY. Specifically, the surprise of sampling some sensory outcome (or experiencing some sensory state) can be represented with the negative log probability: $-\log p(D=d_{t+1}|a_t, M)$. This measure quantifies the improbability that a sensory outcome d_{t+1} is sampled, given action a_t and the generative model M . If the sensory outcome is “incompatible” with M and a_t , then the sensory sample d_{t+1} is surprising. If there's a high probability that biological systems are found at any point in their lifetime in homeostatic states, then environmentally-generated sensory samples will be unsurprising. Sensory samples generated by all other external states in the environment will be highly surprising.

2.4 From surprise to free energy

If one stipulatively defines *adaptively-behaving system* as any system whose behavior minimizes the average surprise of its sensory states, then actual systems behaving adap-

tively must sample unsurprising sensory outcomes. That is, the system's objective is to maximize model evidence $p(D=d_{t+1}|M)$, or minimize surprise $-\log p(D=d_{t+1}|a_t, M)$. This means, formally, that the system must select actions that optimize the function:

$$G(a_t) = \log \int p(D = d_{t+1}, \Psi = \psi_{t+1} | A = a_t, M) d\Psi$$

where $p(D=d_{t+1}, \Psi = \psi_{t+1} | A = a_t, M)$ is the joint density of sensory samples (i.e., sensory states) D and their generating causes Ψ in the environment, conditioned on action and on a generative model. This density factors into a likelihood $p(D=d_{t+1} | \Psi = \psi_{t+1}, A = a_t, M)$ and prior density $p(\Psi = \psi_{t+1} | M)$, which jointly specify the generative model “entailed by” the system's phenotype. Optimizing $G(a_t)$ requires changes in actions of systems with phenotype corresponding to the generative model M .

For most problems, optimizing $G(a_t)$ involves an intractable marginalization over (hidden) environmental states Ψ . To overcome computational intractability, a variational (or recognition) density $q(\Psi, \mu)$ can be introduced to define another quantity that is greater than surprise. This quantity—called *free energy*—provides a bound on the integral mentioned above, and is a function of sensory samples and internal states of the system. It is defined thus:

$$F(d_{t+1}, \mu_{t+1} | a_t) = -\langle \log p(D = d_{t+1}, \Psi = \psi_{t+1} | A = a_t) \rangle_q + \langle \log q(\Psi = \psi_{t+1}, \mu_{t+1}) \rangle_q$$

Under certain assumptions (cf., Dayan et al. 1995; MacKay 1995), optimizing $F(d_{t+1}, \mu_{t+1} | a_t)$ is computationally tractable. In the context of FEP, this optimization involves changes only to internal parameters μ or to the action parameter a , which can be controlled by the system.

Because the free-energy function F is greater than G , by acting on the environment to minimize the free energy of their sensory samples, biological systems would avoid surprising sensory states. If they avoid surprising sensory states, biological systems may attain a homeostatic state; and by selecting actions that attain homeostatic states, biological systems will thereby behave adaptively and preserve their physical integrity. And by FEP, any system that minimizes the free energy of its sensory states with respect to action, or its internal parameters, will avoid phase transitions, which would make its physical disintegration unlikely.

3 FEP and organicism

Having articulated the steps involved in the reasoning to FEP, let's examine two dimensions along which FEP and organicism appear inconsistent: their understanding of how, adaptive organisms should be represented, and their interpretation of probability.

3.1 How to represent organisms?

Free-energy theorists formulate FEP using the traditional modeling tools of random dynamical systems in thermodynamics, and represent organisms' adaptive dynamics

as trajectories through attractive non-equilibrium states in phase space. For organicists, however, organisms' adaptive dynamics cannot be adequately represented with this tool predefined over the "characteristic" variables individuating kinds of biological systems (Longo et al. 2012).

Organicists have different options for justifying this claim. They may deny the existence of an organism's characteristics, or deny that characteristic variables (if they exist) can be reliably identified for any biological system. Another option is to emphasize that the mathematical tools used to represent and explain biological phenomena are merely that—abstract mathematical tools that can be used to describe how one state evolves into another state over time. For instance, Chater and Oaksford forewarn about the risk of conflating scientific models and their targets of the models with this example:

the theory of aerodynamics is a crucial component of explaining why birds can fly. But clearly birds know nothing about aerodynamics, and the computational intractability of aerodynamic calculations does not in any way prevent birds from flying. Similarly, [systems] do not need to calculate their optimal behavior functions to behave adaptively. They simply have to use successful algorithms; they do not have to be able to make the calculations that would show that these algorithms are successful. (2000, p. 110)

The suggestion is that the properties of the models used to represent organisms should not be directly ascribed to the organisms' biological or cognitive equipment. If so, this would bear directly on the posit of *free energy*, which free-energy theorists introduce because the optimization problem described in Sect. 2.4 is intractable. Yet, if there's no imperative for organisms themselves to compute solutions to the optimization problem, and the optimization problem is a problem life scientists need to solve to understand organism's behavior, then the motivation to advance FEP dissipates.

Free-energy theorists assume that organisms are ergodic: their phase averages (i.e., average values of specified functions of their microscopic states) are identical to averages over time of quantities measurable from microscopic states. For ergodic systems, the dynamics of a system's microstates—e.g., its biomolecular kinematics—are sufficiently random, and the coupling between the system and the external states in its environment is sufficiently slow, such that the microscopic dynamics of the system's states can be replaced by a random sample from the ensemble density of the microscopic states. This ensemble density assigns a probability to each possible microscopic configuration, and can be used to derive predictions about macroscopic properties of the system as those expected in the ensemble. Given ergodicity, for any region of phase space, the average time the system spends in that region is proportional to the region's "attractiveness." Ergodic systems will repeatedly revisit the neighborhoods of attracting states. So, if organisms are ergodic, they can be represented as random dynamical attractors in phase space. If such ergodic dynamical systems also possess a Markov blanket, then "[they'll] appear to actively maintain their structural and dynamical integrity" (Friston 2013, p. 10). Under certain conditions, such systems will appear to minimize their free energy.

Organicists will be quick to note that we cannot assign meaningful probabilities to the microscopic states of organisms because organisms, at all levels of organization,

cannot explore all possible paths in phase space. “Not only will we not make all possible proteins of length 200 or 2000, we will not make all possible organs, organisms, social systems, [...] here is an indefinite hierarchy of non-ergodicity as the complexity of the objects we consider increases” (Kauffman 2013, p. 167). For organicists, ergodicity is biologically irrelevant simply because organisms are non-ergodic (Longo et al. 2012).

If ergodicity were biologically irrelevant, and the average of any measure of the state of an organism doesn’t converge over a sufficient period of time, then it is a misleading idealization to represent an organism’s phenotype with an invariant ergodic ensemble density (assuming it can be meaningfully defined) that specifies the probability, for any possible microstate, that the organism is in a certain macrostate. And it would also be biologically irrelevant to note that, under ergodic assumptions, the long-term average of surprise is entropy.

While organicists may think of FEP as misleading and biologically irrelevant, free-energy theorists would counter by asserting that the notion of an attracting set itself implies ergodicity. So, if all living systems possess an attracting set, then they must be ergodic too; and if those systems are ergodic, then they will possess characteristic, relatively stable, measurable properties. For free-energy theorists, the dynamics of such systems will appear to place an upper bound on their informational entropy, and to maximize the evidence for a model M of external states “entailed” by their characteristic properties. This behavior—they would conclude—can be expressed as approximate Bayesian (active) inference about the causes of sensory states in terms of minimizing variational free energy.⁶

On the other hand, organicists have argued that organisms constitute historically grounded constraints on energy flows, where their phase space continually changes, and so organicists deny that living systems are aptly represented with a classical phase space. Since attracting sets are subsets of classically predefined phase spaces, organicists deny the assumption that all living systems’ characteristic behavior is aptly represented with an attracting set. Hence, for organicists, the implication from attracting sets to ergodicity is a red herring.

Representing organisms and their dynamics while idealizing their non-ergodic status would prevent taking seriously the historical considerations of lineage that are essential to understanding what organisms are, and how they change over time. If historical considerations and lineage matter to understanding organisms and their dynamics, then biological systems should be represented as “specific” and their trajectories as “generic.” Instead, for organicists, free-energy theorists get it backward: physical systems are “generic,” while their trajectories “specific” (Longo and Montévil 2013: ch. 7).

If physical systems were “generic,” then different types of systems could be individuated by properties such as mass, charge, temperature, or momentum. Position and momentum, for example, characterize mechanical systems. The mathematical representation of physical systems of the same type will preserve formal symmetries

⁶ This inference-optimization bridge, which we also mentioned in fn. 4, is a powerful feature of variational methods that treat statistical inference problems as optimization problems. Minimizing variational free energy can thus be understood as approximate Bayesian inference, in the sense that minimizing free energy is formally equivalent to optimizing a variational bound on Bayesian model evidence.

between their defining characteristics.⁷ Each symmetry in mathematical representation implies that certain physical properties of the system, such as total kinetic energy, are conserved and remain unchanged as the system evolves over time (Gross 1996). However, for tokens of the same type of physical system, trajectories (or paths) would be “specific” since they are uniquely determined by the rules of the system’s evolution given its initial state.

In free-energy theorists’ representations, organisms are “generic.” Their representation in terms of actions, internal states, and generative models preserves symmetries in formulations of FEP. Trajectories in the phase space of organisms would be “specific” according to FEP, because solutions of FEP for a given kind of organism yield a unique trajectory in phase space given the initial state of the organism.⁸ Properties of organisms like lineage and heritability would just be an expression of a specific trajectory on a generic manifold, namely the attracting set.

For Longo et al. (2012), organisms should instead be represented as “specific,” while their trajectories “generic.” Because of their historicity and materiality, organisms wouldn’t possess general characteristics that allow for mathematically invariant representations. They write,

In biology, symmetries at the phenotypic level, are continually changed, beginning with the least mitosis, up to the “structural bifurcations” which yield speciations in evolution. Thus, there are no biological symmetries that are a priori preserved. [...] There are no sufficiently stable mathematical regularities and transformations to allow an equational and lawlike description entailing the phylogenetic and ontogenetic trajectories. (2012, p. 1390)

So, from organicists’ perspective, organisms cannot—unlike non-living physical systems—be represented with a fixed phase space and rules of evolution predefined over some set of “characteristic,” mathematically invariant variables.

The disagreement between free-energy theorists and organicists cuts deeper still. Free-energy theorists take an adaptationist and selectionist standpoints in biology, whereby “selection explains how biological systems arise and the only outstanding issue is what characteristics they must possess” (Friston and Stephan 2007, p. 423). Organicists reject the adaptationist and selectionist perspectives, which—they argue—are insufficient for explaining the autonomy of living beings, and their capacities to regulate their processes in relation to environmental conditions registered as viable or unviable, improving or deteriorating (Di Paolo and Thompson 2014; Moreno and Mossio 2015).

⁷ For example, in Newtonian mechanics, given two bodies with the same mass starting from rest and moving in opposite directions with different velocities along the same axis, the total kinetic energy of the systems comprised of the two bodies remains the same if the velocities are interchanged. That is, the solution of the equation yielding the total kinetic energy will be the same.

⁸ Friston (2012) relates this result to the principle of least action (PLA). In Hamilton’s formulation, *action* is defined as the integral along possible paths of a system’s process connecting two specified states. According to PLA, the actual path of the process between initial and final states in a specified time is a dynamical system’s trajectory in phase space, which is found “by imagining all possible trajectories that the system could conceivably take, computing the action for each of these trajectories, and selecting one that makes the action locally stationary” (Gray 2009).

Furthermore, following Ashby (1940), FEP emphasizes homeostatic stability as the core feature of organisms. Accordingly, organisms should be represented as random dynamical attractors. Contemporary organicists like Longo and colleagues emphasize that organisms are fundamentally ever-changing processes, maintained in relatively stable conditions by further processes. Longo and Montévil (2013), for instance, advance the idea that organisms should be represented as attaining extended critical phase transitions. Free-energy theorists generally construe undergoing a phase transition as equivalent to biological disintegration and death—although there are several examples of phase transitions in biology, such as metamorphosis, that are consistent with an attracting set, and, arguably, with FEP too (Clark 2017).

Even so, Longo and Montévil (2013) argue that organisms' adaptive behavior should be represented as a continuous critical transition from one phase to another, whereby organisms are continuously reconstructed with variations in their dynamic couplings with ecosystems. The basic idea is that ancestry continuously co-constitutes and reshapes organisms as well as by their interactions with ecosystems. This idea fits a growing wealth of evidence concerning phenotypic plasticity, whereby organisms with the same genotype can generate differing phenotypes through their interactions with the environment, which recreate novel conditions of existence passed on to their descendants (Montévil et al. 2016).

Free-energy theorists would again note that evolution, and natural selection in particular, is also a free-energy minimizing process (Hobson and Friston 2016; Ramstead et al. 2018). But, for organicists, selectionism, and the fixation on surprise-minimizing processes obscure the historically-grounded, environmentally co-constituted nature of biological adaptivity.

One last dimension of disagreement concerns the interpretation of the probabilities involved in scientific representations of changing organisms. Probability plays a central role in free-energy theorists' accounts. Adaptive organisms are said to be embodied generative models of their environments, where a generative model M specifies the probability that a certain external state, sensory input, and internal state occur together (Friston 2013). Organisms are said to have Markov blankets separating their internal states from the external environment, defined over a set of random variables and a probability measure (Kirchhoff et al. 2018). Furthermore, Friston (2009, 2010b) claims FEP entails the Bayesian brain hypothesis, which implies that nervous systems represent probability distributions and store generative models and prior probabilistic knowledge about the world, and that neural networks can perform statistical inferences based on these probabilities (Knill and Pouget 2004; Colombo and Seriès 2012). Finally, free energy itself is a bound on surprise (or information entropy), and surprise is a probabilistic measure of the uncertainty of sampling some sensory data, given a generative model.

Neglect of both the differing scope of distinct formulations of FEP (as applied to non-biological adaptive systems, organisms, or brains at a certain time scale) and the interpretations of the probabilities in different formulations of FEP has generated disagreement—particularly about whether FEP is committed to positing mental representations and to an essentially inferential picture of cognition. To move the debate forward, at least two questions should be distinguished. First, should we understand FEP as a modeler's tool to characterize and predict adaptive behavior, or should it

be understood as an objective feature of target systems? Second, in determining the scope of FEP, how should the probabilities it posits be interpreted? Should they be understood epistemically or physically?

Some researchers in the organicist tradition seem to interpret the probabilities involved in FEP physically, as the frequency or propensity of the occurrence of some event (e.g., Bruineberg et al. 2018; Kirchhoff and Froese 2017). When FEP targets brains, these researchers reject the idea that brains literally represent probabilities and draw inferences. Other researchers have interpreted the probabilities involved in FEP epistemically, as rational degrees of belief in the occurrence of some event and the willingness to act on this belief. When FEP targets brains, these researchers tend to suggest that brains literally represent probabilities, possess Markov blankets, and draw inferences (e.g., Hohwy 2016). For his part, Friston (2013) seems to interpret the probabilities involved in FEP as objective features of real-world systems; they aren't just modeling tools useful to capture the fluctuating dynamics of biological systems. Risking conflation of scientific models and their targets, Friston (2011, 2013) claims that brains themselves represent probabilities about the occurrence of events, and that brains make inferences about the causes of their sensory inputs, based on neurally-encoded statistical models.

Let's consider the first idea. In formulating FEP, one need initially specify a system's phase space (or sample space) and a desired non-equilibrium steady state distribution in it. One can then optimize the non-equilibrium steady-state distribution by minimizing the free energy of samples generated by the environment with respect to actions and a generative model. The system—claims Friston (2011, 2013)—will then appear to sample its environment as if it were aiming at maximizing the evidence for its own existence. However, the construction of 'as if' explanations falls short when the goal is construed as specifying the real nature of biological explananda. Where the functional capacities attributed to those systems prove intractable, resorting to such explanations won't circumvent the problem, even when the computations involved are construed as subsymbolic, offline, heuristic, or approximate (van Rooij et al. 2018).

The desired steady-state distribution of an organism is, by definition, just the organism's evolved extended phenotype. This means that minimizing free energy would be equivalent to maximizing expected adaptive value: surprising states would just be, by definition, maladaptive. One way to support this definition is to note that organisms, and brains, are immersed in a "statistical bath" of energy arrays from the environment impinging on their sensory surfaces. These energy arrays would sculpt phenotypes, associated with a certain non-equilibrium steady-state distribution, which the organism would update as a function of its sensory samples.

However, organicists will argue that defining *adaptivity* as unsurprisingness is misguided and relies on implausible assumptions. With Longo et al. (2012), one may note that a desired non-equilibrium steady state distribution is only definable on a predefined phase space. If it's not possible to predefine a phase space for biological systems, the definition will be undercut—and with it, the justification for believing that the probabilities involved in FEP are not mere modeling tools but are objective features of living systems.

Furthermore, leaving on the side free-energy theorists' selectionist perspective, one may find implausible that surprising events should always be maladaptive. Using

Gershman and Daw’s examples, “[s]hould the first amphibian out of water dive back in? If a wolf eats deer not because he is hungry, but because he is attracted to the equilibrium state of his ancestors, would a sudden bonanza of deer inspire him to eat only the amount to which he is accustomed?” (2012, p. 306). If *adaptivity* is stipulatively defined in terms of minimization of surprise, then some instances of adaptive behavior will consist in seeking out novel situations that may provide systems the opportunity to resolve expected surprise.

Let’s assume that the probabilities involved in FEP aren’t simply modelers’ tools, and turn to the second question about how to interpret these probabilities. In its maximal scope formulation, the probabilities involved in FEP should be interpreted physically, as objective propensities or frequencies. (After all, it makes little sense to say that an amoeba has a certain degree of belief that a bacterium is in the premises.) Physical interpretations of probabilities cohere both with how probabilities are generally understood in statistical mechanics, as well as with the suggestion that inference talk is inapposite to the behavior of organisms like an amoeba or bacterium, understood as dynamical systems coupled with their environment (Bruineberg et al. 2018).⁹ Finally, a propensity-based interpretation clarifies the sense in which FEP isn’t a tautology, just like it isn’t a tautology to say that dice produce odd numbers more often than threes—more on tautology and FEP in Sect. 4.1. Propensities of free-energy minimizing organisms to survive, or of dice to fall equally often on each side, permit fallible predictions about their behaviors.

Yet, a propensity interpretation of the probabilities involved in FEP is questionable. For starters, it’s opaque what sort of property a propensity is. If propensities are causal tendencies, then they should be asymmetric relationships just like causal relationships; but then propensities cannot be probabilities (Humphreys 1985). After all, probabilities can be reversed: given the probability of E given C , there is the probability of C given E , which is a theorem in the Kolmogorov’s axiomatization of probability. If propensities are probabilities that obey Kolmogorov’s axioms, they can also be reversed: e.g., a given Marans chicken has the propensity to lay brown eggs, just as a given brown egg would have a propensity to have been laid by a Marans chicken (which makes no causal sense).

⁹ The notion of inference in FEP also requires clarification. Bruineberg et al. claim that, “[w]ithin the Free Energy framework, the notion of ‘inference’ is much more minimal and does not involve any propositions: any dynamical system A coupled with another B can be said to “infer” the “hidden cause” of its “input” (the dynamics of B) when it reliably co-varies with the dynamics of B and it is robust to the noise inherent in the coupling” (2018, p. 2436). They invoke Huygens’s case of the synchronization of two pendulum clocks to illustrate this “minimal” sense of inference. Three points in response. First, whether or not the concept INFERENCE in FEP involves propositions depends how to understand the probabilities featuring in the free energy framework. If these probabilities are modelers’ tools without actual counterparts in target systems, then INFERENCE may involve propositions, viz., those propositions entertained by scientists when they make inferences about target systems on the basis of the mathematics of random dynamic systems theory. Second, reliable covariance robust to noise falls within the extension, not of the concept INFERENCE, but of some successor notion; so Bruineberg et al. have not illustrated a “minimal sense” so much as they’ve just changed the subject. Third, Huygens observed phase/opposition coupling between two pendulum clocks hanging from a beam or from a board sitting on two chairs. But FEP is formulated within the mathematics of random dynamical systems, and Huygens’s case has not been aptly represented as a problem in parameter estimation in dynamical systems—a problem that FEP is supposed to help solve (see Oliveira and Melo 2015). Ultimately, Bruineberg et al.’s claim does not settle the question of how INFERENCE should be understood within the free-energy framework.

If propensities are instead long-run relative frequencies, then a reference class should be defined. For biological organisms, such a reference class would correspond to a phase space; but if a phase space cannot be predefined for an organism, the probabilities involved in FEP cannot be defined.

Suppose instead that the probabilities in FEP are understood epistemically as rational degrees of belief. This interpretation is most plausible when FEP is restricted to whole, cognitively-sophisticated animals, i.e., to creatures possessing mental representations and capacities for rational inference. Organisms would thus act adaptively by minimizing a free-energy bound defined over internal, graded, epistemic states representing events in the environment.

While this interpretation wouldn't mitigate the problem of defining a phase space (or sample space) for organisms, Friston and colleagues surmise that "sustained exposure to environmental inputs causes the internal structure of the brain to recapitulate the causal structure of those inputs. In turn, this enables efficient perceptual inference" (Friston et al. 2006, p. 77). This suggestion risks obfuscating that adaptive pressures on biological observers are generally unrelated to the accuracy of their epistemic states. From an evolutionary perspective, good prior beliefs should promote adaptive fitness, not simply match relative frequencies in the environment. From a statistical perspective, prior beliefs that merely "recapitulate" observed environmental statistics do not obviously "enable efficient inference." They instead lead to overfitting, which in turn "leads to a fragility of performance in future encounters with the same class of environments, which is maladaptive" (Feldman 2017, p. 31). More entropic priors that do *not* recapitulate the statistics of the environment will avoid overfitting, and may generally be more apt to "optimize the fit between mind and world" (ibid.).

4 FEP and mechanism

We now turn to mechanistic philosophy, and argue that FEP is inconsistent with mechanism along two dimensions of representation: the dependence of explanatory force on describing mechanisms and the rejection of the idea that life science phenomena can be adequately explained through an axiomatic, physics-first approach.

4.1 How to explain life phenomena?

Free-energy theorists appeal to FEP in explanations of various phenomena, including Hebb's rule and spike-timing dependent plasticity, the multiplicity and hierarchical organization of cortical layers, their reciprocal connection with distinct feedforward and feedback properties, and the existence of adaptation and repetition suppression (Friston 2010a). Such explanations are thought of axiomatically, as logical deductions from sets of axioms and formulae (Friston 2012). Stipulative definitions, like *living system* as an attracting set in a phase space or *adaptive* behavior as behavior that reduces average surprise, provide the bridge principles that connect theoretical predicates from different disciplines, and that allow free-energy theorists to attempt the deductions needed to claim reductions of other principles to FEP. In their sweeping attempt to

explain these and other phenomena, and to reduce their theories and principles to FEP, free-energy theorists have also claimed for their theory another virtue: a grand theoretical unification (Friston 2010a; Hohwy 2014).

In presenting these features, FEP is apparently at odds with mechanists' emphasis that life science phenomena should be explained by appeal to mechanisms, and that adequate strategies for explanation in the life sciences should involve decomposing these mechanisms into component parts and operations and providing an account of how these parts and operations work together to produce the phenomenon.

FEP and mechanism are related by their common emphasis on function. Both free-energy theorists and mechanists can agree that FEP provides an idealized functional principle. Where they diverge is over whether and when functional accounts of biological phenomena suffice for adequate explanation, or whether structural details of component parts and mechanistic organization are also necessary for such functional accounts to have explanatory power.

Progress in the life sciences—and especially the cognitive and behavioral sciences—often begins by empirically identifying and adequately describing a functional capacity. Descriptions of that capacity then allow for the method of functional analysis, wherein that capacity is functionally decomposed in terms of its constituent properties, processes, and components. Functional properties of a system have traditionally been individuated by their relations to inputs, outputs, and other internal properties of the system under investigation—that is, by their causal role. By redescribing systems' capacities in terms of their functional properties and dispositions, functional analysis offers scientists a way to tackle the target phenomenon. But it also offers the potential for prediction and explanation, as empirical applications of FEP demonstrate (e.g., Bastos et al. 2012).

FEP provides an analysis of adaptive behavior, in the sense that it assumes a thermodynamically sufficient condition on adaptive behavior, and a homeostatically necessary condition. In the abstract, FEP requires an interplay between four sets variables: sensory samples D that influence a system Σ 's internal states, and active states A that influence the external states of the local environment Ψ . The mathematical dependencies between Σ (D , A), and Ψ define free energy over Σ as a function of D and a probabilistic representation of its causes—the thought being that a system's capacity to adapt to its environment can be functionally analyzed in terms of suppression of free energy, via internal representations that readjust state changes in its transducers (to maintain or improve perceptual fidelity) or effectors (to maintain or promote successful control).

For their part, mechanists have argued that functional analyses lack explanatory power, as they are mechanism sketches (Piccinini and Craver 2011). Mechanism sketches are conceived of as incomplete representations of (the function of) a mechanism, in which some relevant aspects—either component parts or their operations, or their organization—are omitted from the explanation. Biological or cognitive functions may be decomposed into their constituent properties, processes, or subroutines; and decomposition may detail a capacity as a nexus of functional relationships between variables standing for component operations, without thereby specifying how the capacity is actually realized. Mechanists acknowledge that functional decompositions of capacities into modeled causal and non-causal operations within a mechanism

constrain the possible structures and configurations that might perform those operations; but they are equally keen to emphasize that structural decompositions into modeled components within a mechanism can also constrain the possible functions and configurations performed. Details about the relevant physiological and anatomical components of mechanisms are necessary to filling in mechanism sketches by localizing each operation to its respective component part (Bechtel and Richardson 1993/2010).

Free-energy theorists' functional analyses appeal to states and dynamics that idealize away from the biophysical details of the structural complexity of actual systems. While functional analyses may be construed as mechanism sketches, FEP itself doesn't provide in any obvious sense a sketch of a mechanism, since it swings free of the need to supply any microstructural, biophysical, or anatomical detail. Mechanists would therefore conclude that FEP lacks explanatory power (Kaplan and Craver 2011).

Free-energy theorists may reply that FEP defines a class of process models that provide hypotheses about how spatiotemporally organized components and operations in biological systems might carry out free-energy minimization. For example, some have suggested that FEP is algorithmically carried out by predictive coding, and that predictive coding is physically realized by specific hierarchically-structured neuronal assemblies (Friston 2009; Bastos et al. 2012). Accordingly, certain higher-order neuronal assemblies would output predictions of the states of lower-order assemblies, which are then compared with the actual states of the lower-order assemblies to form prediction errors that are passed back up the hierarchy to update the predictions from higher-order neuronal assemblies. The recurrent exchange of signals between adjacent neural assemblies resolves prediction error at each level, resulting in hierarchically deep, neurally-encoded accounts of sensory inputs.

While it may go some way toward meeting the demands of filling out a mechanism sketch, this suggestion is inadequate for mechanists. First, predictive coding is only one of several possible algorithms that may be used to optimize energy functions. Since many of them might be empirically adequate but difficult to disentangle, concerns of underdetermination may emerge. Second, while predictive coding has been used to model some aspects of visual perception (Rao and Ballard 1999), "the experimental evidence for it seems currently inconclusive in the sense that it does not rule out Bayesian inference with a direct variable code, potentially in combination with a variety of non-probabilistic processes" (Aitchison and Lengyel 2017, p. 224). Third, because FEP is intended to generalize beyond organisms with a nervous system, predictive coding would need to be a mechanism by which all adaptive systems work—from bacteria, to winged snowflakes and bladder cells, to plants and social networks. Obviously, for such a wide array of systems, appealing to message-passing in neuronal hierarchies as a relevant mechanism by which they operate is insufficiently general. Fourth and finally, even if issues of underdetermination were put aside and the experimental evidence for predictive coding were overwhelming, and even if it were shown to be the generalized mechanism by which all adaptive systems work, mechanists would be positioned to claim that filling out the mechanism sketch is what matters: appeals to the mechanism of predictive coding—not FEP—are what provides explanatory depth. For mechanists, functional descriptions of capacities that require

mechanistic analysis to achieve this depth can be important principles, but not foundational ones that do the heavy lifting in ultimately explaining biological phenomena.

Free-energy theorists may simply reject the idea that adequate scientific representation of life science phenomena must target the component parts and operations and internal organization of mechanisms. Exemplifying this rejection is Chirimuuta's (2018) contention that several explanatorily adequate models in computational neuroscience are non-mechanistic. The models she considers would be instances of "efficient coding explanation," which, abstracting away from biophysical specifics, answer why certain neuronal systems should behave in the ways described by the models. Based on design principles informing the model, such explanations would thus identify the functional utility of general patterns of behaviors instantiated by neural systems.

Similarly, free-energy theorists may argue that FEP is an optimality principle in the life sciences (cf., Rice 2015). FEP, along with the class of process models it defines, provides us with idealized, coarse-grained descriptions of certain functional variables, which leverage varieties of realization types against the drive to detail lower-level biomechanical structures essentially involved in producing target explananda-phenomena. FEP, and the class of process models it defines, would omit causal-mechanical detail and make biologically unrealistic assumption to focus attention on very general observable patterns displayed by biological systems, but also by any other system behaving adaptively. Given that aim—they might claim—it's misguided to charge FEP for not providing us with mechanistic information.

4.2 FEP as a first principle?

As we have seen, explanations derived from FEP abstract away and distort most of the mechanistic features of their target phenomena. But if FEP doesn't aim at uncovering mechanisms or difference-makers, then what's its epistemic status vis-à-vis mechanistic explanation, exactly?

Unfortunately, FEP's epistemic status is muddled. It has been called an *unfalsifiable platitude*, an *imperative*, a *tautology*, a *stipulative definition*, *paradigm*, *law of the life sciences*, *law of nature*, an a priori *first principle*, a *unifying explanation*, and a *simple postulate or axiom*. Wiese and Metzinger assert that "FEP can be regarded as the fundamental theory, which can combine the different features of predictive processing described above within a single, formally rigorous framework" (2017, p. 12). Friston and collaborators contend that "free energy minimization may be an imperative for all self-organizing biological systems" (2012, p. 2117), and that "the whole point of [FEP] is to unify all adaptive autopoietic and self-organizing behavior under one simple imperative; avoid surprises and you will last longer [...]", which is a principle so basic that "there is no need to recourse to any other principles" (Friston et al. 2012), and "The tautology here is deliberate, it appeals to exactly the same tautology in natural selection (Why am I here?—because I have adaptive fitness: Why do I have adaptive fitness?—because I am here). Like adaptive fitness, the free-energy formulation is not a mechanism or magic recipe for life; it is just a characterization of biological systems that exist" (ibid.). For his part, Allen (2018, p. 19) characterizes FEP as a *normative*

theory, an axiomatic, self-evidently true natural law, and a tautologically true axiom, likening it to a *paradigm, framework, and a research programme* as well.

This rhetorical jumble makes it harder to understand the status of FEP as a first principle, and thus the contrast between FEP and mechanism. So, some clarification is called for. To begin, one should not follow Allen or Wiese and Metzinger in confusing *principles* with *theories, paradigms, and research programmes*, since these technical terms refer to different species of scientific representations, with different properties and scientific and philosophical purposes.

Likewise, that any self-conserving system, via environmental exchanges, must, as a condition of possibility on its existence, minimize an information-theoretic bound on a negative log probability is an intriguing thought—but not one that qualifies as a platitude or a truism, under any normal understanding of those concepts.

One plausible thought is that FEP is a first principle because it's an axiom or postulate: “[t]he free-energy principle is a simple postulate that has complicated ramifications” (Friston 2011, p. 91), and again, “FEP derives [is?] a normative, a priori first principle from a provable definition of living systems” (Allen and Friston 2018, p. 2473).¹⁰ And in fact, as mentioned, free-energy theorists attempt an enormous variety of derivations from FEP. In that sense, FEP may be said to play the role of a first principle. But while they utilize an axiomatic approach grounded in the mathematics of theoretical physics (Friston 2012), FEP is a principle that is itself derived from other statements and definitions. So, it is not an underived axiom or postulate. By implication, principles like FEP need not be underived axioms to play the role of first principle in the life sciences.

Free-energy theorists often characterize FEP as a principle in the life sciences because it's “tautological,” even “unfalsifiable.” Such claims are not easily interpreted. If FEP were a tautology, then free energy theorists may have a triviality problem. As Klein remarks, “[a]ppeal to apparent tautologies should trouble you. For whatever tautologies do, they don't explain why things happen” (2018, p. 2552). Tautologies may offer a starting point for explaining the end goal states of optimal systems, but the empirical adequacy of models with added biophysical causal detail is far removed from triviality. And the attempted explanations relying on FEP do not involve anything like the decomposition and localization of biophysical mechanisms underlying adaptive behavior.

Similarly, falsifiability is normally treated as a hallmark of any scientific claim; so if principles must be falsifiable to be scientific and FEP is unfalsifiable, then FEP is not a scientific principle.¹¹ What's intended cannot be that FEP is unfalsifiable because it's merely stipulated; for while there are stipulative definitions involved in the transcendental argument, FEP is not one of them. What's intended cannot be that FEP is unfalsifiable because it fails to be truth-apt either, since it would then not be

¹⁰ It's unclear in what sense the definition of *living system* as an attracting set is “provable.” Definitions are commonly restatements that do not stand in need of proof, or are even capable of being proved. More plausibly, Friston and colleagues want to claim that the definition they propose is legitimate—though it's not obvious what criteria of legitimacy should be in place to evaluate this claim.

¹¹ One reviewer encourages free-energy theorists to bite the bullet: “if a principle has to be falsifiable, then FEP has no epistemic status.”

a law-like generalization, and could not serve as the conclusion of a transcendental argument.

Presumably, then, FEP is like all other scientific principles in being truth-apt. But unlike other principles such as Galileo's principle describing the periodicity of pendula, which subsequently enjoyed more accurate and precise formulations, it seems that what's intended is that principles like FEP or Hamilton's PLA cannot but help survive all scrutiny of their pedigree and have truth-values that cannot be improved upon.

In this sense, FEP might be a constraint that mechanist explanations in the life sciences must honor, in so far as life scientists aim to determine which values of some energy functional constitute the best available solution—given certain design constraints—to the problem of maintaining the path of a target biological system in state space within a specific (homeostatic) region that precludes phase transitions.

Unlike what advocates have claimed (e.g., Hohwy 2014), FEP-based theorizing is not consistent with mechanists' idea that the power of an explanation depends on its capacity to uncover the mechanism of a target phenomenon. But unlike what mechanists will reply, the response is not obviously just to claim, "so much the worse for FEP". This apparent inconsistency is methodologically good: the boom of research relying on FEP just highlights there is room for deductive systematization and physics-first approaches in life science theorizing (see also Gurova 2011). Rather than aiming to represent difference-makers of life science phenomena, FEP instead aims to represent, in the language of information theory and random dynamical systems theory, what characteristics complex systems must possess for self-maintenance and self-regulation.

5 Conclusion: a plurality of principles

Living organisms are complex, adaptive systems that present both robust regularities but also constant variation. Understanding their regularities and variations requires conceptual frameworks in which knowledge from physics, chemistry, biology, ecology, and ethology can be synthesized, using tools from mathematics and computational theory. The diversity of expertise involved in understanding brains and organisms, and the fragmentation in present-day neuroscience and biology, highlights the need for principles that could afford a common intellectual framework for researchers from different communities to work together to answers questions of common concern.

FEP is an impressive candidate for one such first principle aiming to ground general, idealized models for tracking one fundamental pattern underlying the robust regularities and constant variation displayed by complex and diverse phenomena in the life sciences. FEP symbolizes what a first-principle, physics-first, axiomatic approach to the life sciences looks like, while it has crystallized the notion of prediction-error minimization in philosophical and scientific debates as a central theoretical posit to understanding life and mind.

In this paper, we have examined numerous disagreements between FEP and basic tenets of organicism and mechanism concerning the scientific representation of life phenomena. The incompatibility between these different approaches should not suggest that only one of these approaches can aptly represent the phenomena of life. These

different approaches are meant to fulfill different epistemic aims of different scientific communities; and these aims may be best pursued separately, with a plurality of modeling tools (Mitchell 2002).

Life scientists and philosophers can, and often do, productively argue about the aptness of modeling tools for their intended purposes. But they sometimes engage in speculative metaphysics, by directly reading off metaphysical conclusions about the nature of life and mind from usage of models they implicitly assume to be uniquely apt. Some of the confusions in recent debates surrounding FEP, organicism, and mechanism depend on indulging this sort of metaphysics without carefully considering the epistemic and pragmatic roles that “rampant and unchecked” idealizations, like those involved in FEP, play in science (see Potochnik 2017 on the role of “rampant and unchecked” idealizations in science).

Acknowledgements We would like to thank Marshall Abrams, Joe Dewhurst, Giuseppe Longo, Matteo Mossio, Galen Pickett, the participants of the 2018 workshop “The Promise of Predictive Processing” at Tufts University, and two anonymous referees for helpful conversations and comments on previous versions of this paper. In addition, MC gratefully acknowledges financial support from the Alexander von Humboldt Foundation.

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