

Sloppy model analysis identifies bifurcation parameters without normal form analysis

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Bifurcation phenomena are common in multidimensional multiparameter dynamical systems. Normal form theory suggests that bifurcations are driven by relatively few combinations of parameters. Models of complex systems, however, rarely appear in normal form, and bifurcations are controlled by nonlinear combinations of the bare parameters of differential equations. Discovering reparameterizations to transform complex equations into a normal form is often very difficult, and the reparameterization may not even exist in a closed form. Here we show that information geometry and sloppy model analysis using the Fisher information matrix can be used to identify the combination of parameters that control bifurcations. By considering observations on increasingly long timescales, we find those parameters that rapidly characterize the system's topological inhomogeneities, whether the system is in normal form or not. We anticipate that this novel analytical method, which we call time-widening information geometry (TWIG), will be useful in applied network analysis.

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I. INTRODUCTION

This paper provides a method for extracting bifurcation parameters from a set of dynamic equations by combining information geometry and bifurcation theory. Both are useful for modeling multiparameter systems and systems with multiple regimes of behavior respectively, but together they provide methods for data-driven analysis of a wide array of natural phenomena. By creating an explicit connection between the information in the signal (model output) and the model parameters, we identify the combinations of parameters responsible for topological change in the dynamics, the codimension of the bifurcation, and the timescale necessary to resolve this information. The information further provides the directions normal to the separatrix, which divides behavioral regimes of the system.

Traditionally, when confronted with a high-dimensional, multiparameter system of dynamic equations, bifurcation analysis proceeds by attempting to simplify the system to a manageable size. Center manifold reduction exploits the Hartman-Grobman theorem [1] to create a lower-dimensional linear map in the region of a critical point that is locally accurate and is a rapid way to determine the system stability. Shoshitaivishili extended this method to nonhyperbolic equilibria, creating a container for critical modes to straighten out nonlinear terms and, ideally, drop some of them [2]. Such methods have been used to describe phenomena as diverse as neural network optimization and foraging decisions in monkeys [3,4].

A related approach is the method of Poincaré-Birkhoff normal forms. It uses appropriately centered manifolds to

analyze which nonlinear terms are essential and must remain even under optimal coordinate transformations. Such transformations are useful, because the reduced normal-form equations typically have greater symmetry than the initial problem, a property that can be exploited by many analytical tools. Though powerful, “in practice lengthy calculations may be necessary to extract the relevant normal-form coefficients from the initial equations” [2]. Even if such coefficients can be found, neither their interrelationship nor their relative sensitivities are always apparent. It is often the case that some parameters differ by many orders of magnitude in their effect on long-term dynamics, and a method that does not distinguish among them is suboptimal for most applications.

The method of Lyupanov exponents is an admirably general tool for analyzing the global stability of a system. Unfortunately, it provides little information about which specific parameter combinations lead to system (in)stability. For the purposes of bifurcation analysis, it is therefore sometimes paired with sensitivity analyses based on the global sensitivity metrics of Sobol' [5]. These measures, along with useful extensions such as the Fourier amplitude sensitivity test and importance measures [6–8], are able to determine exactly how much of a model's variability is due to each of its parameters. While this often works in practice, there are two potential pitfalls in this approach. First, it assumes that the parameters responsible for variability are also responsible for instability, which is not always the case. Second, if the bifurcation is caused by combinations of many parameters (as frequently happens), then variability will often be high across all these parameters even though the bifurcation itself has a low codimension. In other words, a low-dimensional bifurcation surface generally cuts diagonally across parameter space unless appropriately reparameterized. Once such a transformation is applied and the system is reduced to a normal form (see Sec. III), then the codimension should be apparent, but finding that reparameterization is still likely

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to be cumbersome, if not impossible, in closed form. Just one such transformation can require several papers, as in the case of high-dimensional diffusion-activated processes from Kramers, through Langer, and finally to one dimension, derived using iterations of singular value decomposition by Berezhkovskii [9].

A third, independent line of analysis comes from renormalization group (RG) methods, which are usually applied to study universal power laws near critical points. Feigenbaum [10] was the first to note such universalities in bifurcations of the discrete period-doubling type, a result that he and others extended until it included all major bifurcation types [11–15]. Working from the other direction, scientists investigating critical phenomena with RG (e.g., many behaviors of quantum chromodynamics) have discovered bifurcations and used the tools of one to analyze the other [16,17]. A remarkable study found deep equivalence between RG transformations and normal form theory, showing that the difficult transformation of an ODE system into a normal form could often be accomplished to at least second order by applying three RG transforms [18].

More broadly, universal scaling laws and RG analysis of critical points is often associated with emergence and the systematic irrelevance of many degrees of freedom. Recent work has extended these ideas to a broader class of systems known as “sloppy models” [19–23]. The moniker “sloppy” is meant to convey that these systems have a few combinations of parameters that are many orders of magnitude more influential than other parameter combinations. More precisely, one unit change in a “stiff” parameter direction has as much influence as a million or more unit change in a different “sloppy” direction. Sloppy model analysis relies heavily on the techniques of information geometry [20,24,25] and in this paper we use the terms interchangeably. These techniques have motivated novel reduction algorithms by removing unimportant, sloppy parameters [25–27]. Recent work [28] demonstrates that as coarse-graining of RG models proceeds, the flow causes information of “relevant” parameter combinations to be maintained while “irrelevant” parameters are compressed and become sloppy. These ideas share a common goal with bifurcations analysis in which many diverse systems are collected into a few universal, normal forms. This paper closes the loop, showing how information geometry applies directly to bifurcation analysis without passing through the “middleman” of renormalization group theory. The usefulness of such an analysis, which we call time widening information geometry (TWIG), also circumvents the need for the other types of analyses described above.

In this work, we demonstrate similar notions of “relevant” and “irrelevant” parameters near a bifurcation using the formalism of information geometry and sloppy models. The intuition behind this approach is as follows. Topological inhomogeneities in the flow field produce trajectories containing different information on either side of a bifurcation. For example, on one side of a Hopf bifurcation, all trajectories collect into a central fixed point, while they flow into an orbit (limit cycle) on the other side. TWIG works by measuring the information content in these trajectories at increasingly long timescales and identifying those combinations of parameters to which the trajectory is most sensitive. At long timescales,

these are the parameters responsible for the bifurcation, while parameters that cause only local variability have less impact.

Information geometry can be applied to complex systems from many disciplines—but especially systems biology—to iteratively “reverse engineer” optimal statistical models by removing parameters whose value has little influence on the macroscopic behavior of the system [19,21,26,29]. However, it was recently appreciated that such reverse engineering can be done even if the underlying system bifurcates into qualitatively different behaviors, because the information geometry of parameters participating in the bifurcation show a characteristic “sand dune” shape when crossing from one behavioral state to another [30]. These results imply that if the functional form of the system is known, then it should be even easier to determine bifurcation parameters than if the system’s equations need to be inferred.

This paper is organized as follows: In Sec. II, we provide background information on bifurcations and information geometry generally and, specifically, how we conceptualize them for the purposes of applying the latter to the analysis of the former. In Sec. III, we show how an IG analysis of the normal form bifurcations rapidly provides insight into the structure of bifurcations simple enough to be understood by other methods. Section IV shows how this analysis extends to more difficult bifurcations, the implications of which are summarized in Sec. VI.

II. BACKGROUND AND PROBLEM FORMULATION

A. Bifurcations

Bifurcations frequently arise in the analysis of dynamical systems, where one typically characterizes the flow field with special attention to any fixed points or stable oscillations [31]. Consider a generalized system of n -coupled dynamic equations, where each equation is of the form $\dot{\mathbf{y}} = f(\mathbf{y}; \theta)$, where θ is a vector of m parameters. Small changes to any of the θ_i values typically result in correspondingly small changes to the n -dimensional vector field, such as small changes to the position of a fixed point or radius of a limit cycle. Such deformations are topologically equivalent (meaning the number and properties of the attractors and repellers in the field do not change) and homeomorphic (continuous with a continuous inverse). However, there may be critical parameter values where a small change causes new fixed points to emerge from old ones, two fixed points to approach and be mutually annihilated, or limit cycles to be broken. Such nonhomeomorphic transformations are generically called bifurcation no matter their exact form.

Several types of simple bifurcations have been identified and reduced to their simplest possible mathematical expression. These are the “normal forms” and are enumerated in the section below. These forms are convenient starting points for analysis, since they have clearly defined rate parameters that are unambiguously responsible for causing topological inhomogeneities. However, even elegant mathematical descriptions of real-world dynamical systems rarely conform exactly to one of the normal forms.

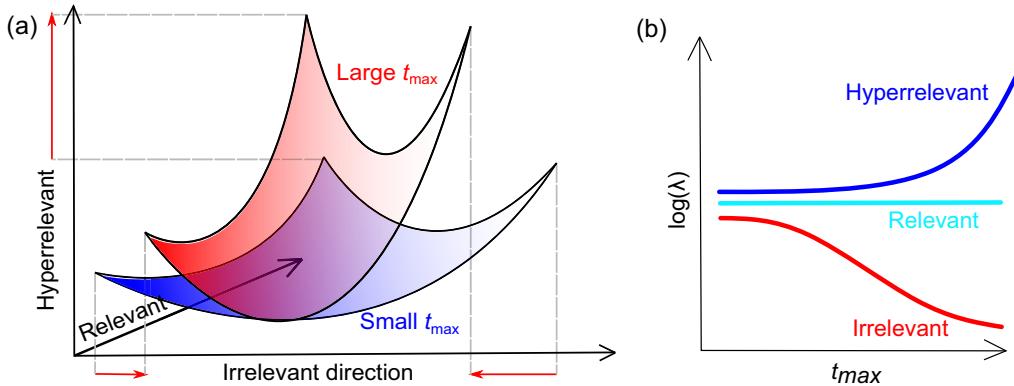


FIG. 1. (a) The model manifold in data space represents all values that can be reached. The axes represent directions that are distorted in characteristic ways as t_{\max} increases. They can be contracted (irrelevant), expanded (hyperrelevant), or unchanged (relevant). (b) Relevance can be quantified by observing the eigenvalues of the Fisher information matrix as t_{\max} is increased. Eigenvalues that do not change at longer timescales retain their relevance, while those that increase or decrease become either more or less relevant.

Bifurcation parameters for physical models often do not align with the bare parameters. In the classic example of boiling liquid, the bifurcation parameter is some combination of temperature, pressure, salinity, and others. In general, a reparameterization to a single, unambiguous bifurcation parameter may be possible in principle but often requires either substantial additional physical insight, mathematical sophistication, or both. Some researchers have even recommended building an analogous physical circuit as the fastest method to detect the bifurcation [32]. Complex models can have hundreds of coupled dynamic equations with thousands of parameters (e.g., models of sophisticated mobile phone circuit boards [33] or complex protein networks [34]). How can we determine which parameter (or, more likely, combination of parameters) is responsible for the bifurcation in such cases?

B. Information geometry

The fundamental object of information geometry is the Fisher information matrix (FIM or \mathcal{I}), which quantifies the information that the observations \mathbf{y} contain about the parameters θ of a dynamical system. Here we introduce the FIM for dynamical systems.

Consider a system of ordinary differential equations where the parameters are tuned to be exactly at their critical values, i.e., the system is at (one of) its bifurcation point(s). The system is allowed to evolve, and the trajectory of one of its equations y_j is sampled at several time points $y_j(t_i)$, where $t_i = t_0 + \frac{i}{n}t_{\max}$. To help visualize this process, let us imagine a one-dimensional system,

$$y(t) = \theta_1 + e^{-\theta_2 t} + e^{\theta_3 t}, \quad (1)$$

sampled at $t = \{1, 2, 3\}$ to create a vector of three observations $\mathbf{y} = \{y(t_1), y(t_2), y(t_3)\}$ which we plot in \mathbb{R}^3 , i.e., data space. If $\theta_3 > 0$, then there is no equilibrium; if $\theta_3 = 0$ and $\theta_2 > 0$, then the equilibrium is at $\theta_1 + 1$ or $\theta_1 + 2$ if $\theta_2 = 0$. As the parameters of θ change, the position of \mathbf{y} will also change, but except for extreme values of θ_i , it cannot reach all possible values in \mathbb{R}^3 . The space filled by values of \mathbf{y} that can be reached for a given range of parameter values defines

the model manifold. A schematic of such a manifold is drawn in Fig. 1(a).

The Fisher information is most commonly defined in probabilistic terms as the expected Hessian matrix of the log-likelihood:

$$\mathcal{I} = -E \left[\frac{\partial^2}{\partial \theta^2} \log \mathcal{L}(\theta | d) \right], \quad (2)$$

where θ is a vector of parameters and d is the data. For deterministic systems such as we consider here, it is standard practice to assume that measurements are obscured by additive Gaussian noise,

$$d_i = y(t_i) + \zeta, \quad (3)$$

where $y(t_i)$ is the (deterministic) output of the model at time t_i and ζ is standard normal random variable $\zeta \sim \mathcal{N}(0, 1)$. This assumption defines a probability distribution to which Eq. (2) can be applied [24]. Because this construction is so common in information theory, it is often referred to as the sensitivity Fisher information matrix or sFIM [35] for reasons that will soon be apparent. In general \mathcal{I} can be expressed in terms of the first derivatives only

$$\mathcal{I} = -E \left[\frac{\partial^2}{\partial \theta^2} \log \mathcal{L}(\theta | d) \right], \quad (4)$$

$$= E \left[\frac{\partial}{\partial \theta} \log \mathcal{L}(\theta | d) \frac{\partial}{\partial \theta} \log \mathcal{L}(\theta | d) \right]. \quad (5)$$

Using the second form, one can show that sFIM becomes

$$\mathcal{I}_{i,j} = \sum_{k=1}^M J_{k,i} J_{k,j} = (J^T J)_{i,j}, \quad (6)$$

where we have introduced the Jacobian or sensitivity matrix $J_{k,j} = \frac{\partial y_k}{\partial \theta_j}$ whose entries denote the sensitivity of prediction k to changes in parameter j . In Eq. (6), M denotes the number of observations.

The entries of the FIM indicate the sensitivity of the model's trajectory to changes in each pair of parameters. A high score indicates that a parameter pair has a strong influence on model dynamics, while a small score indicates a “sloppy” direction (parameter values can change a great deal

without much changing \mathbf{y}). The curvature of the likelihood function converts distances in parameter space to distances on the manifold in data space, making the FIM a Riemannian metric tensor on the model manifold in data space. It is important to note that the physical units of parameters can strongly affect the values of the FIM. For this reason, it is common to perform dimensional analysis before sloppy model analysis as we do throughout this study.

In general the curvature of the likelihood surface does not align with the bare parameters. Rather, the characterization of the model's sloppiness aligns with the eigenvectors of \mathcal{I} . Eigenvalues of the FIM are related to the singular value decomposition of $J = U \Sigma V^T$:

$$\mathcal{I} = V \Sigma^2 V^T, \quad (7)$$

where U and V are matrices of the left and right singular vectors of J and Σ is the diagonal matrix of its singular values. This implies that the right singular vectors of the Jacobian V are also the eigenvectors of the FIM. The eigenvectors of \mathcal{I} "orient" the parameter space into the parameter combinations most relevant for changing the model's behavior.

Imagine now that we coarsen the sampling rate by changing t_{\max} . In our simple example, increasing t_{\max} from 3 to 6 means the model \mathbf{y} is sampled at $t = \{2, 4, 6\}$. This procedure stretches the manifold in some directions and compresses it in others. This distortion is measured by an increase or decrease in the eigenvalues of \mathcal{I} , respectively. Compression of the manifold (i.e., decreasing eigenvalue) with increasing t_{\max} indicates that the combination of parameters is less important to the long-term dynamics. We call the corresponding eigendirection "irrelevant." Similarly, if the manifold stretches (i.e., increasing eigenvalue), then we call the corresponding direction "hyperrelevant." Directions that are neither compressed nor stretched are called "relevant" directions [Fig. 1(b)]. Returning to the example in Eq. (1), θ_1 is relevant since its effect on the model's output is unchanged with observation time. In contrast, θ_2 is irrelevant since the exact rate of the decay matters less as timescales become very large, and θ_3 is hyperrelevant since small changes have large effects at large t . Note that θ_2 and θ_3 are functionally interchangeable if either is negative.

This procedure is similar to coarse-graining under RG flow described in Ref. [28] and is used to generate their Fig. 1. In our case, however, because we are coarsening the sampling rate, the total observation time increases and includes new information, i.e., observations at later times. As such, it is not a true coarse-graining and introduces the possibility of hyperrelevant directions, i.e., directions that become increasingly important such as θ_3 . We will see that hyperrelevant directions are associated with the stability or instability of the equilibrium.

This method is also somewhat analogous to studies that use Sobol' sensitivity analysis to track importance at different timescales, either bare parameters or eigenvalue combinations. Such methods are excellent at providing estimates of model variability at a given point in parameter space and have noted both increasing and decreasing importance for model parameters of biophysical systems [36,37]. Critics note that these methods are computationally expensive, even when

implementing Morris acceleration [38], and the implications for bifurcation analysis are not immediately obvious.

In addition to characterizing bifurcations, TWIG analysis reveals two other features of bifurcating systems. First, there can be parameters (or combination of parameters) that move the location of a fixed point without causing a bifurcation. Such parameter combinations appear as "relevant" eigendirections, as the new equilibrium appears in long-time observations. These parameters need to be removed in order to correctly identify the codimension of the bifurcation. We do this by solving for the location of the fixed point with a numerical root-finding algorithm and subtracting it from the trajectory at every point. This effectively translates the fixed point to the origin and is analogous to the recentering step of center manifold analysis. For limit-cycle trajectories, we recenter by subtracting off the (unstable) fixed point that must exist within the cycle (according to the Poincaré-Bendixson theorem [39]).

The second feature arises in such oscillating systems. Parameters that change the phase or frequency of oscillation without destroying the equilibrium itself appear as hyperrelevant as the accumulating phase difference becomes increasingly important at late times. Previous research has shown that such systems frequently cause problems in an information geometry framework by introducing "ripples" into the likelihood surface of Eq. (2). The solution is to perform a coordinate transformation so the period itself becomes a parameter. In one formulation of the FIM, this causes the manifold to "unwind," creating a smooth likelihood surface [40] and thereby eliminating a misleading eigendirection.

Four important pieces of information come from this TWIG analysis. First, the number of hyperrelevant and relevant directions corresponds to the codimension of the bifurcation system. Second, the square of each element of the eigenvector matrix V_{ij} indicates the participation factor of each bare parameter θ_i in eigenvector j . This last fact follows because the participation factor $p_{ij} \equiv U_{ij} V_{ij} = V_{ij}^2$ as can be seen by combining the definition of a participation factor [41,42] with Eq. (7) above which shows that the left and right singular vectors of the FIM are identical. Third, the eigendirections themselves will change as t_{\max} increases and parameters that influence the short-term dynamics lose their salience at long timescales. If initial conditions are included as parameters, then their loss of relevance is a strong indicator that the system has been simulated "long enough" to capture equilibrium behavior. This is not a trivial concern in practice, where long numerical simulations are always fighting the accumulation of computer round-off error. Finally, at equilibrium the relevant eigendirections point along the (potentially) high-dimensional separatrix surface, and so the bifurcation can be mapped through all parameter space.

Note that this procedure works no matter the number of dynamical variables involved in the differential equation system. However, it presupposes that the model can be simulated on at least one side of the bifurcation to arbitrarily long times, i.e., it analyzes stable dynamics on the threshold of instability. A bifurcation that switches between two different forms of instability will not be easily detectable with this method, as trajectories will diverge on both sides of the bifurcation. In

the next section, we demonstrate how this procedure works for all common normal forms of bifurcations.

III. NORMAL-FORM BIFURCATIONS

Local bifurcations can be described mathematically in a potentially infinite number of ways, but nearly all of them can be reparameterized, at least locally, to one of five kinds of normal forms. These are as follows:

(1) *Saddle-node*: $\dot{x} = r + x^2$, where one stable and one unstable fixed point emerge from a previously uninterrupted flow at a critical value r_{crit}

(2) *Transcritical*: $\dot{x} = rx - x^2$, where a stable and an unstable fixed point exist everywhere but at the bifurcation, and swap stability at the critical value

(3) *Supercritical pitchfork*: $\dot{x} = rx - x^3$, where symmetric stable fixed points emerge from a single fixed point, which itself becomes unstable

(4) *Subcritical pitchfork*: $\dot{x} = rx + x^3$, symmetric unstable fixed points emerge from an unstable fixed point, which itself becomes stable

(5) *Hopf*: a stable limit cycle emerges from what had previously been a stable point attractor. Depending on the coordinate system, the normal form is $\dot{z} = z(a + b|z|^2)$ (complex), $\dot{x} = -y + x(\mu - r^2)$; $\dot{y} = x + y(\mu - r^2)$ (Cartesian), or $\dot{r} = r(\mu - r^2)$; $\dot{\theta} = -1$ (polar).

A method able to detect bifurcation parameters for these types of bifurcations will detect the overwhelming majority of bifurcations we are likely to encounter. The Fisher information as a function of t_{max} for each bifurcation type has a closed-form solution, which complements and validates the numerical results that we present here (see Appendix A). In each case, the sensitivity with respect to the bifurcation parameter, r , dominates the long-term dynamics of the system in the neighborhood of the bifurcation, no matter how many other higher-order parameters are added to the normal form.

For example, a supercritical pitchfork of the form $\dot{x} = rx - x^3 + \alpha_1 x^4 + \alpha_2 x^5 \dots$ experiences a bifurcation when $r = \alpha_i = 0$. At short timescales (e.g., where $t_{\text{max}} < 1$), the system's trajectory is strongly influenced by changes to its initial value x_0 and the higher-order α terms (for $x_0 > 1$). However, later dynamics show that changes to the α_i 's (and x_0) barely affect the trajectory of approach to equilibrium at 0, while small modifications to r move the equilibrium itself (Fig. 2). An eigenanalysis of the FIM (Fig. 3) quantifies these insights and clearly demonstrate the effect of coarse-graining on the system (i.e., increasing t_{max} while keeping the number of samples constant). At very short timescales ($t_{\text{max}} < .05$), x_0 and the highest-order α term are the main participants of the leading eigenvector, and x_0 soon falls off as t_{max} increases; recall from Fig. 2 that this high-order term was equivalently able to bend the trajectory significantly until $t \approx 1$. Around $t_{\text{max}} = 10$, r begins to have a noticeable influence on the observed trajectory, and correspondingly this is the point where r becomes the dominant participant in the leading eigenvector. For large t_{max} , the leading eigenvalue increases while all other eigenvalues decrease, indicating that the system's bifurcation is codimension one. Note that in this range, small changes to the initial value x_0 have fallen all the way to the last eigenvector, indicating that the system has been allowed to run long

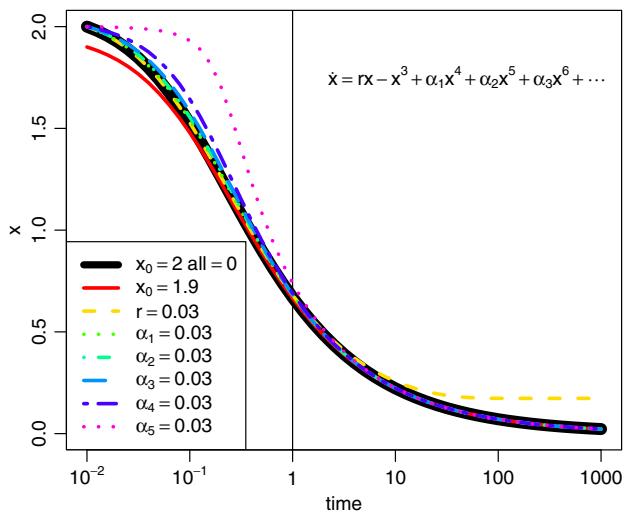


FIG. 2. The trajectory of a supercritical pitchfork at the bifurcation point (heavy black line), and slightly perturbed from it (thin colored lines). At short timescales, y_0 (thin red) and high-order parameters (long-dashed blue) appear relevant. But as the dynamics progress, r (short-dashed yellow) emerges as the only parameter that changes the long-term equilibrium point. This change from relevant to not (and vice versa) occurs at $t \approx 10$ and is reflected in the arch shape and changing colors of Fig. 3.

enough that transient dynamics are removed or at least have orders of magnitude less influence than any of the nuisance parameter α 's. There is no significance to the fact that in this and subsequent “rainbow diagrams,” the leading eigenvalue eventually begins to increase; this is simple case of an increasing line overtaking nonincreasing ones and nothing inherent about the highest eigenvalue at small timescales. This can be confirmed by the change in color, indicating that the parameter responsible for the leading eigenvector has changed.

Similar figures can be produced for the saddle-node, transcritical, and subcritical pitchfork bifurcation classes. In each case, the eigenanalysis of the FIM indicates

(i) how long the system should be simulated by the time it takes for the effect of the initial conditions to reach the least relevant eigenvector

(ii) the codimension of the bifurcation by the number of nondecreasing eigenvalues (= 1 for each normal form),

(iii) the participation factor of each parameter in the (hyper)relevant directions by the square of the corresponding eigenvectors (asymptotically approaching 100% r in each normal form)

(iv) the null space of the bifurcation surface, making it possible to track the bifurcation hypersurface through parameter space.

These are relatively simple bifurcations, where the separatrix is the hyperplane $r = 0$. In more complicated situations where the separatrix is a nonlinear combination of bare parameters, this analysis identifies the vector normal to the separatrix. In principle, this local characterization could be extended to map that separatrix [along the (hyper)relevant directions] through the high-dimensional parameter space.

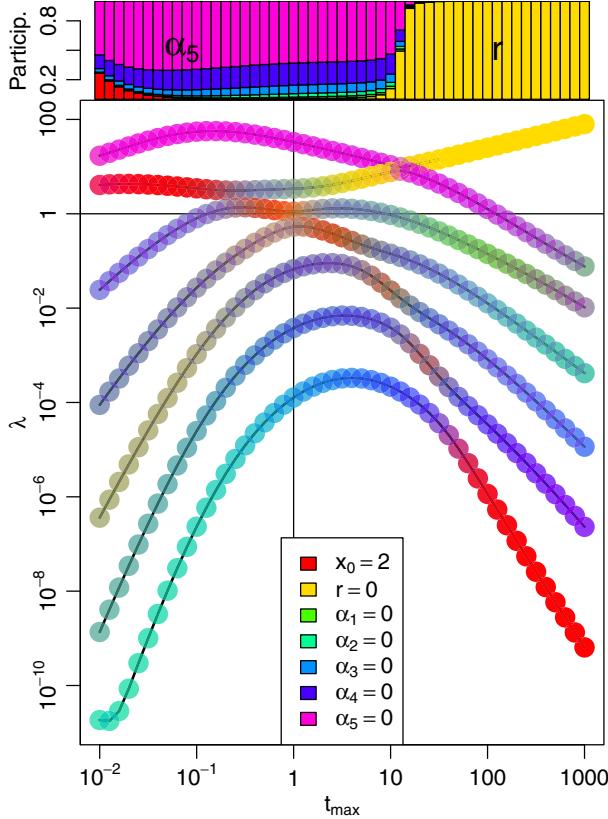


FIG. 3. The “rainbow diagram” of the system from Fig. 2, showing the FIM’s eigenanalysis at each t_{\max} . The top panel represents the participation of each parameter in the first eigenvector $[V_{i,1}^2]$ in Eq. (7)]. The leading eigenvector changes from α_5 dominant (purple) to r dominant (yellow) at $t_{\max} \approx 10$, i.e., just where the α_5 trajectory is replaced by the r trajectory as most divergent in Fig. 2. The large panel below shows all seven eigenvalues $[\lambda_i = \sqrt{\Sigma_{ii}^2}]$ in Eq. (7)] at each t_{\max} , colored as the weighted average RGB of each parameter’s participation factor. Thus, the top line, corresponding to the largest eigenvalue in the top panel, starts mostly purple (α_5) but turns yellow as r dominates the first eigenvalue at larger t_{\max} values. For all parameters, a small change to parameter values influences trajectory at short timescales (the rising limb) but, with the exception of r , not at long timescales (the descending limb). The red color in the bottom-right indicates that the initial value x_0 eventually becomes the least relevant parameter in the model. Pure colors indicate an eigenvector pointing along a parameter axis, while mixed colors like browns and grays indicate that many parameters participate in the eigenvector.

Hopf bifurcations present more of a challenge: The limit cycle that emerges from a fixed point can have both its radius *and* velocity manipulated by model parameters, which can potentially provide the false impression that the system has codimension 2 rather than the actual codimension of 1. Consider the following Hopf bifurcation in polar coordinates, where, as above, additional high-order terms have been added:

$$\begin{aligned} \dot{r} &= \mu r - r^3 + \alpha_1 r^4 + \alpha_2 r^5 \\ \dot{\theta} &= \omega + \beta r^2 + \alpha_3 r^3 + \alpha_4 r^4 + \alpha_5 r^5. \end{aligned} \quad (8)$$

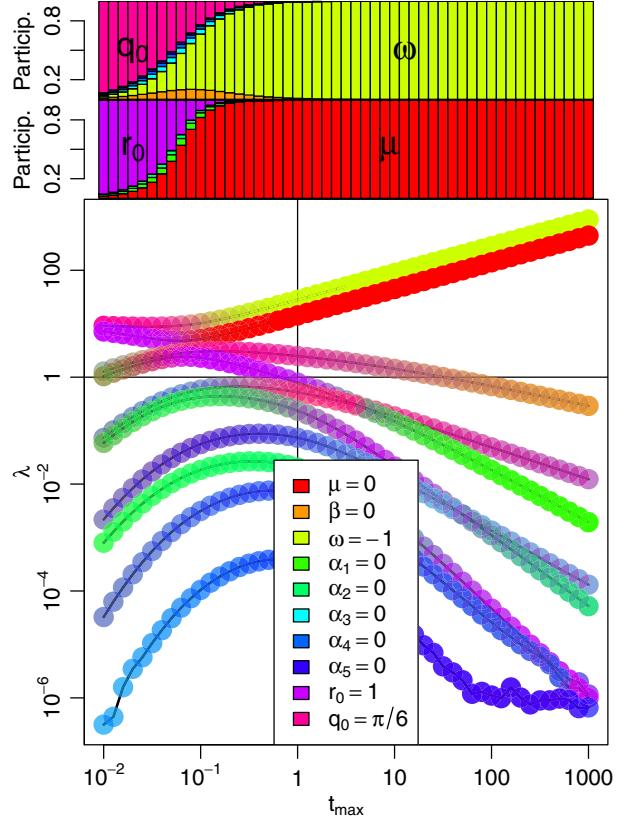


FIG. 4. TWIG analysis of the Hopf bifurcation. The first of the hyperrelevant (rising) eigenvalues comes from the periodicity of the trajectory, whose velocity is set by ω . The second hyperrelevant eigenvalue comes from the bifurcation itself, indicating that the Hopf bifurcation is codimension 1, and the bifurcation depends simply on μ , and not some complicated combination of parameters. Note that the Hopf bifurcation is far easier to simulate at long timescales in polar form than in Cartesian coordinates.

At the bifurcation point $\mu = 0$, a fixed point at the origin expands into a limit cycle. The velocity of trajectories around this cycle are primarily driven by ω , provided y values are small. Note that the periodicity of the Hopf bifurcation introduces a second hyperrelevance to long-term dynamics. Infinitesimal changes to velocity make little difference to the final position of the trajectory $F(t_{\max}; y, \theta)$ if t_{\max} is small but will have an increasing effect as t_{\max} grows. By contrast, μ is hyperrelevant because it is the bifurcation parameter. The increasing importance of these two parameters, in contrast to all others, is clearly illustrated in Fig. 4.

As noted above, this ability to characterize all normal-form bifurcations depends on the ability to isolate changes in information due to the bifurcation itself. This depends on the only source of variation in long-term behavior coming from the bifurcation, and so the preceding analyses were conducted for systems exactly at the bifurcation point. We now consider how the picture changes for dynamics near, but not exactly at, the bifurcation point. Applying TWIG just to the left and right of the bifurcation point of a pitchfork ($r = \pm .01$) shows characteristic patterns (Fig. 5). In these cases, we find

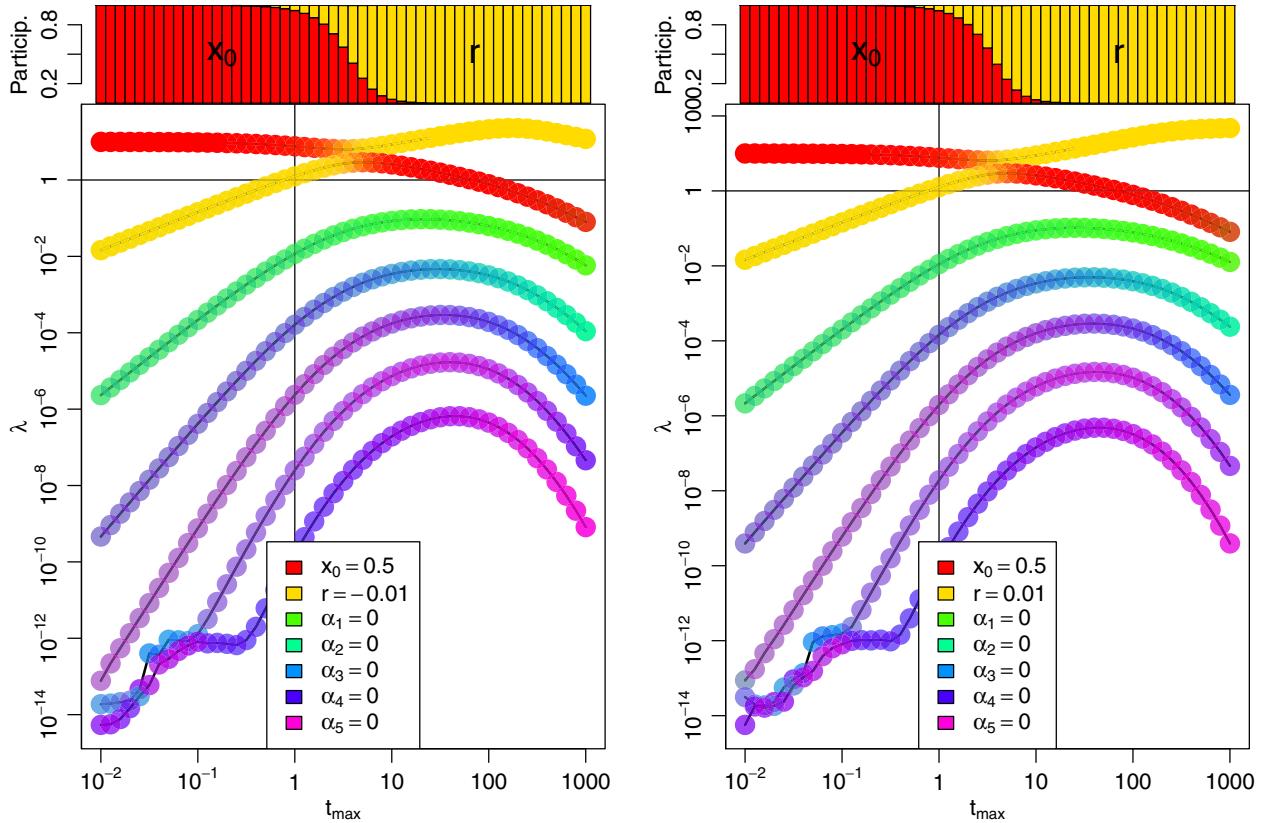


FIG. 5. TWIG analysis near-but-not-at the bifurcation values show the diagnostic pattern of an increasing eigenvalue at intermediate timescales, but then a decrease [in (a)] or an asymptote [in (b)]; top yellow curve in the $t_{\max} = 100$ to 1000 range in both]. It is still relatively easy to identify bifurcation parameters and the bifurcation's codimension, though the falloff becomes more pronounced, and thus the clarity of the analysis more obscured the further one moves away from the bifurcation in either direction.

that the bifurcation parameter is hyperrelevant on intermediate timescales ($10 < t_{\max} < 100$ in Fig. 5). However, on longer timescales ($t_{\max} > 100$), the leading eigenvalue either decreases [Fig. 5(a)] or asymptotes [Fig. 5(b)] once the trajectories have converged to the fixed point, depending on whether the location of the fixed point can or cannot be shifted by changing the parameter values, respectively. In other words, when approached from the $r < 0$ side, small changes to r do not move the equilibrium [$y(t) \rightarrow 0$ as $t \rightarrow \infty$], meaning the exact value of r is irrelevant. But approaching from the $r > 0$ side causes trajectories to run to $y(t) \rightarrow \pm\sqrt{r}$, meaning r is relevant. Moving the system closer to bifurcation, this intermediate regime extends further and further, until at $r = 0$ it occupies the entire trajectory and r is hyperrelevant at all times $t > 10$.

In general, being slightly off the bifurcation obscures the effect of the bifurcation parameter to an extent proportional to the distance from the bifurcation. This is particularly useful in the case of hemistable bifurcations, which need to be approached from the stable side or else test trajectories will diverge to infinity (and cause computer overflow). In the case of the subcritical pitchfork, at the bifurcation itself ($r = 0$) the system is unstable. However, at values of $r \rightarrow 0^-$, just

less than bifurcation value, TWIG can be performed and the bifurcation characterized as above (Fig. 6).

IV. BIFURCATIONS IN NON-NORMAL FORMS

Equations describing real systems are not typically written in one of these normal forms. So even when a researcher knows a system contains a bifurcation, it might not be apparent which one of these it is. For example, a model of a bead on a rotating hoop

$$mr \frac{\partial^2 \phi}{\partial t^2} = -b \frac{\partial \phi}{\partial t} - mg \sin \phi + mr\omega^2 \sin \phi \cos \phi$$

has a supercritical pitchfork bifurcation, though it might require simulating many values of r and ω to appreciate this [31]. Similarly, the equation

$$\dot{x} = r \ln x + x - 1 + \alpha_1(x - 1)^2 + \alpha_2(x - 1)^3 + \dots \quad (9)$$

contains a transcritical bifurcation at $x = 1$ when $r = -1$. However, this only becomes clear after reparameterizing the equation by $R = r + 1$, and $X = \frac{r}{2}(x - 1)$, when the equation assumes the normal form $\dot{X} = RX - X^2 + \mathcal{O}(X^3)$. Such a substitution might not be immediately apparent to

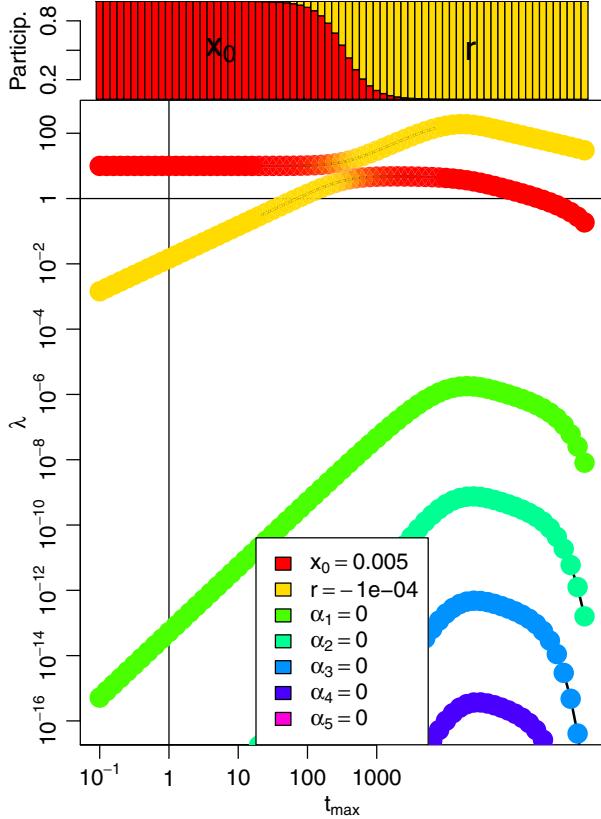


FIG. 6. The subcritical pitchfork cannot be analyzed using TWIG at the bifurcation point ($r = 0$) because the system is unstable. However, simulations slightly to the stable side of the bifurcation ($r \rightarrow 0^-$) reveal the bifurcation parameter, though because analysis happens off the bifurcation, the peak occurs at intermediate values instead of reaching an asymptote.

a researcher; however, time-widening information geometry clarifies the situation.

If the dynamics in Eq. (9) are run long enough, then we observe that one eigenvalue is relevant while all others are irrelevant. Furthermore, the corresponding participation factor becomes dominated exclusively by r (Fig. 7). This tells us that (1) the process has codimension 1 and (2) the reparameterization involves only r . We confirm that our analysis has converged since the initial condition y_0 is the dominant participation factor in the smallest eigenvalues. However, we note that convergence occurs at a somewhat larger value of t_{\max} than in the normal form examples above. Also note that transcritical bifurcations have a leading eigenvalue that is *relevant* rather than *hyperrelevant*, due to a quirk of the normal-form algebra. See Appendix B for a thorough explanation.

But what happens when the situation is not so straightforward? Modifying the above example to the equation

$$\dot{x} = r \ln(x) + a(x - \alpha) + b(x - \alpha)^2 + \dots \quad (10)$$

should still have a transcritical bifurcation for certain parameter values but no simple reparameterization to create a normal form exists. From above, we can recognize that when a transcritical bifurcation occurs at $x = 1$ for $r = -1$,

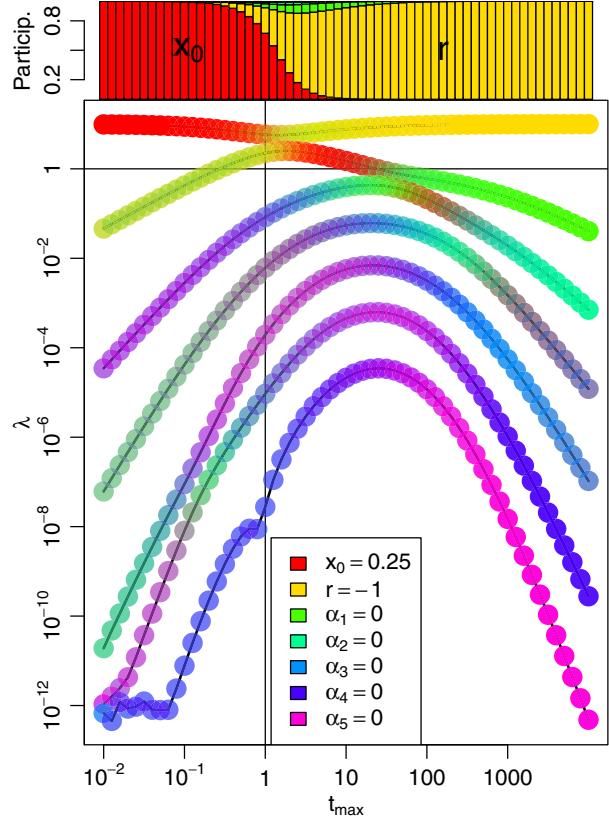


FIG. 7. Equations such as Eq. (9) that are not in normal form can be interpreted using the same procedure as for normal form bifurcations. As above, the presence of just one nondecreasing eigenvalue, whose corresponding eigenvector is dominated by the single parameter r , indicates that the system has codimension 1 and the bifurcation parameter involves only r . The relevant (not hyperrelevant) leading eigenvalue is characteristic of a transcritical bifurcation.

$\alpha = 1$. However, when $\alpha \neq 1$, in the neighborhood of $x = \alpha$ all the power terms are zero, but the term $r \ln(x) > 0$ if $\alpha < 1$, suggesting that no fixed point exists in that region. The appearance or disappearance of a fixed point is the hallmark of a saddle-node bifurcation and indicates that allowing a bit of variability in the fixed point's location has introduced a second codimension to the dynamical system. This is borne out by TWIG analysis (Fig. 8), which shows that the equation indeed produces a hyperrelevant eigenvector corresponding to the saddle-node parameter α , which controls the existence—not just the location—of an equilibrium. The transcritical bifurcation still exists and is controlled by r , as implied by the previous analysis. This example shows that even in situations with two different bifurcation classes, neither of which can be reparameterized into normal form, TWIG still allows us to efficiently and unambiguously identify codimension and bifurcation parameters.

A. A biophysical example

Glycolysis is a multistep process which uses the bond energy of glucose to catabolize energy-carrying biomolecules

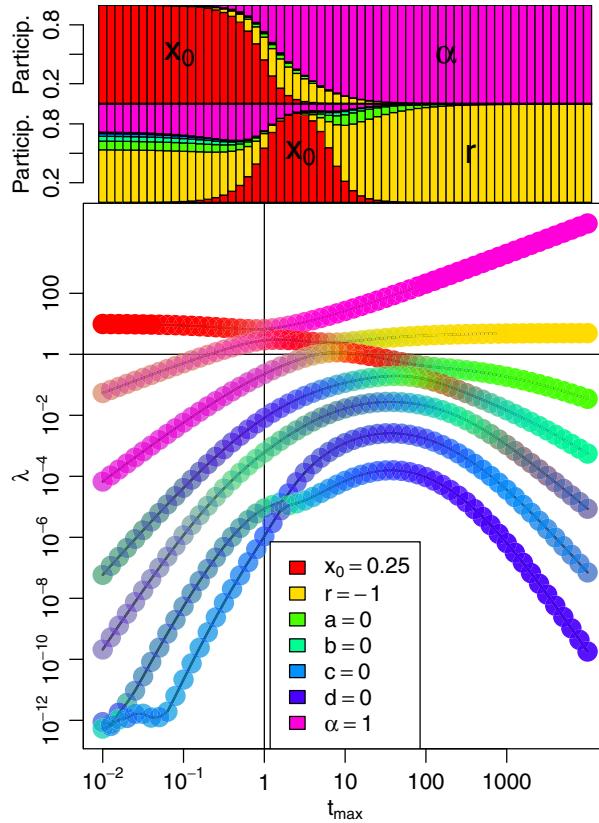


FIG. 8. A difficult non-normal-form transcritical bifurcation such as Eq. (10) can be extremely challenging to analyze analytically, but sloppy analysis indicates one hyperrelevant parameter (corresponding in this case to a saddle-node) and one relevant parameter (as usual, indicating transcritical bifurcation). This means that this system has a bifurcation of codimension two. Note that the participation factor of the two leading eigendirections runs to 1.0 in the direction of α and r respectively, indicating that the system can be placed into normal form without a complicated recombination of parameters.

easily usable by cells, which represents one of the dominant processes of all heterotrophic life on earth. A bottleneck in this crucial process is the phosphorylation of fructose-6-phosphate into fructose-1,6-bisphosphate catalyzed by the enzyme phosphofructokinase. The complicated five-species mass-action equation describing this reaction's kinetics can be simplified using Tikhonov's theorem and assuming low concentrations of ATP to the simple dimensionless system [43,44]:

$$\begin{aligned} \dot{x} &= -x + ay + c_1x^2y + c_2x^3 \\ \dot{y} &= b - ay + c_3x^2y + c_4y^2, \end{aligned} \quad (11)$$

where x and y are the concentrations of ADP and F6P respectively, and the four c_i constants are nuisance parameters added to mask the system dynamics. There is a curved bifurcation surface that separates the range of kinetic parameters a, b which lead to either a fixed point at $(b, \frac{b}{a+b^2})$ when $c_1 = 1, c_3 = -1$ as in the canonical model, or a stable limit cycle. The separation between the fixed point and limit-cycle regimes has the form $b^2 = \frac{1}{2}(1 - 2a \pm \sqrt{1 - 8a})$ [31]. The

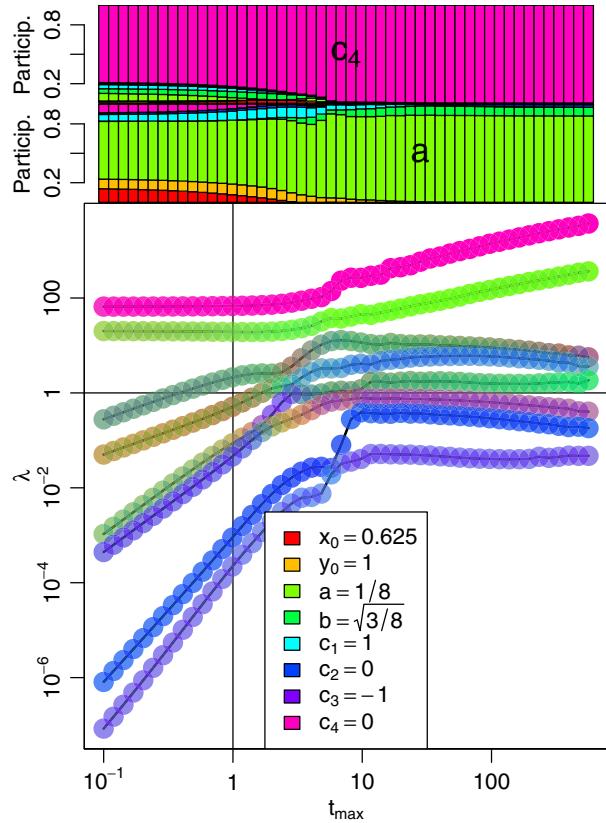


FIG. 9. Analysis of the “glycoscillator” bifurcation [Eq. (11)]. The frequency of the oscillations are driven by c_4 , while the radius of oscillations can be controlled with just one of the a, b parameters discovered by Sel'kov [43].

resulting oscillations in glycolytic activity predicted by this analysis have been observed *in vivo* since the early 1970s [45].

A TWIG analysis of this system provides several insights, summarized in Fig. 9. First, even though the separatrix between fixed point and limit cycle in a, b space is a nonlinear curve, because b can be reparameterized as a function of a , it is codimension one. Second, the “nuisance” parameter c_4 introduces a change in the period of the oscillations, which means that infinitesimal changes in its value cause larger deviations in final trajectory the longer the simulation runs. This shows up as a hyperrelevant direction in TWIG; however, as discussed above, it is *not* a second codimension.

V. CHAOTIC SYSTEMS

Systems showing chaotic behavior have long represented a challenge to traditional categories of thinking, and the difficulty of distinguishing deterministic chaos from randomness is practically its own subdiscipline [31,46–49]. In the context of TWIG analysis, there are two characteristics of the system that need to be considered carefully.

First, unlike other systems considered here, one hallmark of chaos is long-term sensitive dependence on initial conditions, or the “butterfly effect.” Because of this, a TWIG

analysis carried out in the chaotic regime, in contrast to Fig. 3 where the parameter x_0 becomes the least relevant, will classify initial condition parameters as relevant. Note that if the chaotic system produces a strange attractor, then the initial conditions will change the location of the system on the attractor at long timescales but not the shape of the attractor itself, which prevents these parameters from becoming hyperrelevant. That is, the maximum distance between two trajectories begun at slightly different initial conditions will eventually saturate on opposite sides of the attractor and not increase without bound.

Second, the four classic examples of chaotic systems approach chaos through a complicated series of bifurcations rather than a singular event as in the normal-form bifurcations above. The logistic map famously contains a period-doubling “bifurcation cascade,” with the distance between these bifurcation events decreasing geometrically by the Feigenbaum constant α universally [10,11,31]. There is thus a “fuzzy boundary” between the periodic behavior of, say, an 8-cycle and the chaotic region as we pass through the increasingly narrow 16-cycle region, 32-cycle region, and so forth. The Hénon map experiences a similar bifurcation cascade along the line $b = 0.3$ as a increases from 1 to 1.5 [50], while the Rössler attractor has a bifurcation cascade in the opposite direction on the plane $a = 0.2$, $c = 5.7$ as b decreases from 1.5 towards 0 [51]. As we show below for a Rössler system, these boundaries are not just fuzzy but also fractal. Most complex of all, the Lorenz system experiences a pitchfork bifurcation at $r = 1$, whose two stable points then experience Hopf bifurcations at $r \approx 24.74$, while the unstable point undergoes a “homoclinic explosion” at $r \approx 13.926$ that produces an “a thicket of infinitely many saddle-cycles and aperiodic orbits” [31]. If even these pedagogical “toy models” of chaos have such indeterminate boundaries, then it is likely that examples of chaotic systems encountered “in the wild” will as well.

While the FIM may be evaluated at any point in this fuzzy region between order and chaos, its interpretation is less clear. The eigenvectors, which indicate the direction normal to the separatrix in other systems, lose this meaning since there is no direction normal to a fractal surface. Note that this also holds true for the intermittency route to chaos as well. Abrupt changes to chaos, with or without smooth changes in fractal dimension, also exist and would be expected to give cleaner results in the TWIG analysis [52,53], but unfortunately are expected to be less common and less familiar to readers.

That being said, TWIG still can provide powerful qualitative insights into the nature of a period-doubling chaotic system. The Rössler attractor defined by

$$\begin{aligned}\dot{x} &= -y - z \\ \dot{y} &= x + ay \\ \dot{z} &= b + z(x - c)\end{aligned}\quad (12)$$

has a well-known period doubling map revealed by decreasing b along the parameter-space line $a = 0.2$, $c = 5.7$, with the fuzzy transition from an 8-cycle to chaos occurring in the region near $b \approx 0.70$. TWIG analysis carried out near this point reveals that while changes to b or c in this region can lead to long term divergent behavior, changes to a have a

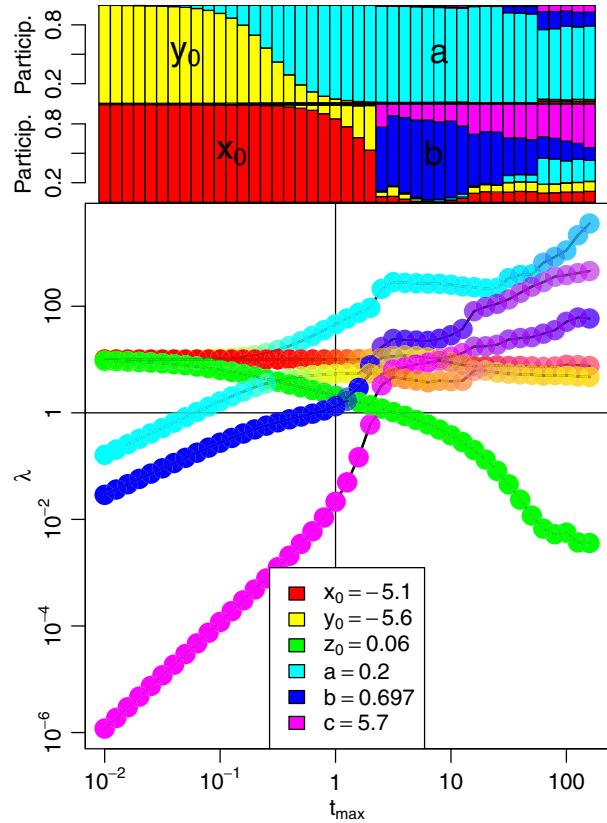


FIG. 10. TWIG analysis of the Rössler attractor, a chaotic system, evaluated in the region of rapid period doubling just before the onset of chaos. Due to the butterfly effect, the initial conditions remain relevant at long timescales and cannot be used to determine appropriate simulation length. However, excluding these from analysis, we are still able to qualitatively see that there is one hyperrelevant direction, dominated by a . This came as a surprise to the authors, because the bifurcation region was approached by changing values of b until a period-doubling cascade was observed, yet TWIG uncovered a greater sensitivity to a than b even in this region. This was confirmed by sampling the parameter space in Fig. 11.

much stronger effect (Fig. 10). In other words, even though we “walked up” to the bifurcation region in the b direction, TWIG was able to tell us that the fuzzy bifurcation boundary was strongly angled normal to the a direction. This insight is not found in the usual treatments of the Rössler attractor [31,51,54] but can be easily verified by simulating the system in Eq. (12) at many sample parameter values in the region around the bifurcation. This reveals flat “sheets” of periodic behavior sandwiched between strata of chaos in the a direction (Fig. 11); these sheets can eventually be encountered for a fixed value of a by moving far enough in b or c , which is essentially the process diagramed in the period-doubling map with which we started this exercise.

Above, we made the claim that TWIG analysis could be used to determine four characteristics of the system, the first being the length of time to run an analysis by the decay of sensitivity to initial conditions. For chaotic systems, this is no longer the case due to the butterfly effect. However, by

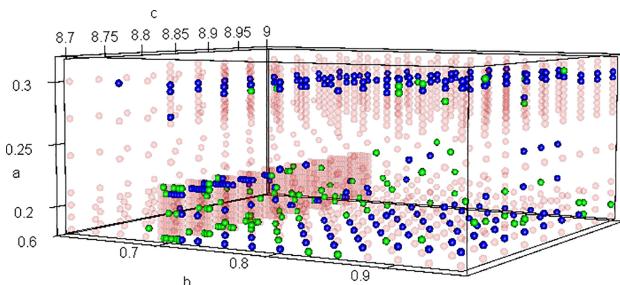


FIG. 11. The parameter space in the period-doubling region of the Rössler attractor shows flat sheets of 8-cycle behavior (solid blue spheres) sandwiched between chaos (transparent red) in the a direction. Green spheres are simulations difficult to classify as either 8-cycle or chaotic.

removing the initial values as parameters, we see that TWIG can still be used *qualitatively* to determine the other three characteristics: bifurcation co-dimension, the null space, and the (fuzzy) normal to the bifurcation region.

VI. CONCLUSION

Gradual time-dilation of the Fisher information matrix as realized by our TWIG analysis is an efficient way of characterizing bifurcations in a dynamical system. Researchers have long used eigenanalysis of \mathcal{I} to characterize the “sloppiness” of a system, i.e., its exponential range of sensitivities to parameter changes, and recently leveraged this accumulated expertise with coarse-graining to understand phenomena occurring at distinct timescales [28,55]. Building on these insights, we here demonstrate that as t_{\max} increases, the changing eigenvalues of \mathcal{I} (and the composition of the corresponding eigenvectors) allow us to (1) characterize the codimension of the bifurcation, (2) quantify the participation of each bare parameter in the bifurcation, (3) characterize the bifurcation’s hypersurface, and (4) have an internal check on the length of time necessary to simulate the system to reach equilibrium. These are substantial insights to be gained relatively cheaply. Sloppy bifurcation analysis constitutes a powerful tool to supplement traditional analytical analysis [31,56] and other specialized analytical tools for high-dimensional problems [6,8,10,34,54,57,58].

Insights derived from TWIG are useful not just for theoreticians interested in characterizing a bifurcation or reparameterizing a system to emphasize the bifurcation; it is also critical for the process of fitting parameter values. The rainbow plots in this paper demonstrate that at the bifurcation point, simulations frequently show a separation of over 10 orders of magnitude in their parameter sensitivity, a gap that gets larger the longer the simulations run (or the more data are collected, in an experimental context). If researchers care about fitting all parameters, then it is crucial to recognize that the effect of hyperrelevant parameters will overwhelm the others, so only if these parameters are fixed in the experiment can the less relevant ones be inferred [30,58]. Future work

may naturally extend the method to large systems including those derived from partial differential equations.

Our TWIG analysis has some inherent limitations. It presupposes that the model can be simulated on at least one side of the bifurcation to arbitrarily long times, i.e., it analyzes stable dynamics on the threshold of instability. A bifurcation that switches between two different forms of instability will not be easily detectable with this method, as trajectories will diverge on both sides of the bifurcation. However, such doubly unstable bifurcations may be of limited practical interest anyway, as loss of stability is generally a far more common real-world problem than a change in the type of instability of a system that never was stable to begin with. Hemistable points (as in saddle-node or subcritical pitchfork bifurcations) are easily analyzed when approached from the stable side (see Fig. 6); otherwise, test trajectories can diverge beyond computer tolerance at moderate timescales. A notable limitation of the method as presented here is the inability to analyze hyperbolic fixed points. Future work may additionally leverage center manifold techniques to investigate bifurcations in such systems. We note here that absolutely unstable fixed points (i.e., where every eigenvalue is positive) can be conveniently analyzed in TWIG simply by running time backwards, and analyzing trajectories at ever-closer instants to the initial divergence from the instability.

Because it is a particularly efficient method of determining important information about high-dimensional bifurcations, we anticipate that TWIG will be useful in situations with many components where one or a few bifurcations are expected in each component. These include power grids, circuit boards, interatomic models, complex protein regulatory networks, and ecosystem-based management systems of multiple interacting populations. Such complexity presents substantial difficulties for closed-form analysis but can be tamed with insights gleaned from this method.

An implementation and example of TWIG analysis are available in Ref. [59].

ACKNOWLEDGMENTS

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APPENDIX A: FIM OF SADDLE-NODE BIFURCATIONS

The normal form of the saddle-node bifurcation is

$$\frac{dy}{dt} = r + y(t)^2 + \alpha_1 y(t)^3 + \alpha_2 y(t)^4 + \dots \quad (A1)$$

This differential equation can be solved locally when all parameters $\vec{\theta} = 0$, which happens to be the bifurcation point of the system. At that point:

$$\frac{dy}{dt} = y^2 \rightarrow \frac{dy}{y^2} = dt.$$

Integrating both sides yields

$$\begin{aligned} -\frac{1}{y} \bigg|_{y_0}^{y(t)} &= t \bigg|_0^t \\ \frac{1}{y_0} - \frac{1}{y(t)} &= t \\ y(t) &= \frac{y_0}{1 - y_0 t}. \end{aligned} \quad (\text{A2})$$

This implies there is a singularity at $t = 1/y_0$, so a proper coarse-graining procedure will involve taking data from $t = 0$ to some value near $1/y_0$, say, $0.99/y_0$. We avoided this singularity by using negative values for y_0 and were therefore able to run simulations to large values of t_{\max} . As noted in Eq. (6), to find the FIM of a system it is only necessary to find the Jacobian, so we need only find the first partial derivative of these data with respect to each parameter in the model.

1. Partial derivative of r

Let the α'_i 's = 0. The derivative of the normal form with respect to r becomes

$$\frac{\partial}{\partial r} \left(\frac{\partial y}{\partial t} = r + y^2 \right) \frac{\partial^2 y}{\partial r \partial t} = 1 + 2y \frac{\partial y}{\partial r}. \quad (\text{A3})$$

We let $w = \frac{\partial y}{\partial r}$, and this becomes $\frac{\partial w}{\partial t} = 1 + 2yw$, which requires the use of an integration factor to solve [60]. If $p_1 x' + p_0 x = q$, then

$$x = \frac{1}{\mu p_1} \left[C + \int \mu q dt \right] \text{ where } \mu = p_1^{-1} \exp \left(\int \frac{p_0}{p_1} dt \right). \quad (\text{A4})$$

Allowing $p_1 = 1$, $p_0 = -2y$, $q = 1$ implies that

$$\begin{aligned} \mu &= 1^{-1} \exp \left(\int \frac{-2y}{1} dt \right) \\ &= \exp \left(- \int \frac{2y_0 dt}{1 - y_0 t} \right) \\ &= \exp[2 \ln(1 - y_0 t)] \\ &= (1 - y_0 t)^2. \end{aligned}$$

Therefore,

$$\begin{aligned} w &= \frac{C + \int (1 - y_0 t)^2 dt}{(1 - y_0 t)^2} \\ &= \frac{C - \frac{(1 - y_0 t)^3}{3y_0} \bigg|_0^t}{(1 - y_0 t)^2} \\ &= \frac{C + \frac{1 - (1 - y_0 t)^3}{3y_0}}{(1 - y_0 t)^2}. \end{aligned}$$

Recall this function is being evaluated at the initial condition, where the partial derivative $w = \frac{\partial y}{\partial r} = 0$ (i.e., changes to r do not change y_0). This implies that $C = -\frac{1 - (1 - y_0 t)^3}{3y_0}$; when

$t = 0$ this further reduces to $C = 0$. Therefore,

$$\frac{\partial y}{\partial r} = \frac{1 - (1 - y_0 t)^3}{3y_0(1 - y_0 t)^2}. \quad (\text{A5})$$

2. Partial derivative of α_1

Using the same procedure as above,

$$\begin{aligned} \frac{\partial}{\partial \alpha_1} \left(\frac{\partial y}{\partial t} \right) &= \frac{\partial}{\partial \alpha_1} (y^2 + \alpha_1 y^3) \\ \frac{\partial^2 y}{\partial \alpha_1 \partial t} &= 2y \frac{\partial y}{\partial \alpha_1} + y^3 + \cancel{3y^2 \alpha_1 \frac{\partial y}{\partial \alpha_1}} \\ \frac{\partial w}{\partial t} &= 2yw + y^3. \end{aligned} \quad (\text{A6})$$

Note on the second line, we are able to cancel the third term because we are evaluating the slope where α_1 is zero. On the last line, note that p_0 and p_1 are the same as for r , so as above $\mu = (1 - y_0 t)^2$, but since now $q = y^3$:

$$\begin{aligned} w &= \frac{C + \int y^3 (1 - y_0 t)^2 dt}{(1 - y_0 t)^2} \\ &= \frac{C + \int \left(\frac{y_0}{1 - y_0 t} \right)^3 (1 - y_0 t)^2 dt}{(1 - y_0 t)^2} \\ &= \frac{C + \int \frac{y_0^3 dt}{1 - y_0 t}}{(1 - y_0 t)^2} \\ &= \frac{C - y_0^2 \log(1 - y_0 t) \big|_0^t}{(1 - y_0 t)^2} \\ &= \frac{C - y_0^2 \log(1 - y_0 t)}{(1 - y_0 t)^2}. \end{aligned}$$

Again, assuming $w = t = 0 \rightarrow C = 0$, so

$$\frac{\partial y}{\partial \alpha_1} = -\frac{y_0^2 \log(1 - y_0 t)}{(1 - y_0 t)^2}. \quad (\text{A7})$$

3. Partial derivatives of higher-order α' 's

Higher-order terms in the series are of the form $\alpha_n y^{n+2}$ and so

$$\begin{aligned} \frac{\partial}{\partial \alpha_n} \left(\frac{\partial y}{\partial t} = y^2 + \alpha_n y^{n+2} \right) \frac{\partial^2 y}{\partial \alpha_n \partial t} \\ &= 2y \frac{\partial y}{\partial \alpha_n} + y^{n+2} + \cancel{(n+2)y^{n+1} \alpha_n \frac{\partial y}{\partial \alpha_n} \frac{\partial w}{\partial t}} \\ &= 2yw + y^{n+2}. \end{aligned} \quad (\text{A8})$$

As above, we are able to cancel $(n+2)y^{n+1}\alpha_n \frac{\partial y}{\partial \alpha_n}$ because we are solving for slopes about the point $\alpha_n = 0$. With the same value of μ , we use integration factors to demonstrate

$$\begin{aligned} w &= \frac{C + \int \left(\frac{y_0}{1-y_0t}\right)^{n+2}(1-y_0t)^2 dt}{(1-y_0t)^2} \\ &= \frac{C + \int \frac{y_0^{n+2}dt}{(1-y_0t)^n}}{(1-y_0t)^2} \\ &= \frac{C - \frac{y_0^{n+1}}{1-n}(1-y_0t)^{1-n}|_0}{(1-y_0t)^2} \\ &= \frac{C + \frac{y_0^{n+1}}{n-1}[1 - (1-y_0t)^{1-n}]}{(1-y_0t)^2}, \end{aligned}$$

which again implies that $C = 0$ at the initial condition $t = 0$, and so for $n > 1$ we can say

$$\frac{\partial y}{\partial \alpha_n} = \frac{y_0^{n+1}[1 - (1-y_0t)^{1-n}]}{(1-n)(1-y_0t)^2}. \quad (\text{A9})$$

Recall that the Jacobian of our system is

$$J = \begin{bmatrix} \frac{\partial y_0}{\partial r} & \frac{\partial y_0}{\partial \alpha_1} & \frac{\partial y_0}{\partial \alpha_2} & \dots \\ \frac{\partial y_1}{\partial r} & \frac{\partial y_1}{\partial \alpha_1} & \frac{\partial y_1}{\partial \alpha_2} & \dots \\ \dots & \dots & \dots & \dots \end{bmatrix}. \quad (\text{A10})$$

Because the Fisher information matrix $\mathcal{I} = J^T J$, we can see that element $\mathcal{I}_{1,1} = (\frac{\partial y}{\partial r})^2$ will be $\mathcal{O}(t^2)$ because $\frac{\partial y}{\partial r}$ is $\mathcal{O}(t^1)$; all other elements will be a lower order of t . Thus, at long timescales, the FIM's element (1,1) will grow faster than all other elements, and therefore the most relevant parameter is clearly r .

In the case where \mathcal{I} is being derived from data [or from noise added to a (non-)normal form equation], the importance of r can be evaluated by increasing $\sigma^2 \propto y_0^{-3}$. Since, by the central limit theorem standard error $\sigma^2 \propto n$, then the number of time points sampled should decrease as $n \propto y_0^{-3}$.

APPENDIX B: FIM OF TRANSCRITICAL BIFURCATIONS

These have a similar normal form as the saddle-node bifurcations above:

$$\frac{dy}{dt} = ry(t) - y(t)^2 + \alpha_1 y(t)^3 + \alpha_2 y(t)^4 + \dots$$

However, the change of sign in the second term causes the solution to the differential equation to also have a changed sign:

$$\begin{aligned} \frac{dy}{dt} = -y^2 &\rightarrow -\frac{dy}{y^2} = dt \rightarrow \frac{1}{y} \Big|_{y_0}^{y(t)} = t \Big|_0^t \\ \frac{1}{y(t)} - \frac{1}{y_0} &= t \rightarrow y(t) = \frac{y_0}{1+y_0t}. \end{aligned} \quad (\text{B1})$$

Now the singularity occurs at $t = -\frac{1}{y_0}$, which generally only complicates the coarse-graining if initial conditions are negative.

1. Partial derivative of r

The full solution to the partial derivative of r is somewhat complicated because it depends on y :

$$\begin{aligned} \frac{\partial}{\partial r} \left(\frac{\partial y}{\partial t} = ry - y^2 \right) \frac{\partial^2 y}{\partial r \partial t} \\ = r \frac{\partial y}{\partial r} + y - 2y \frac{\partial y}{\partial r} \frac{\partial w}{\partial t} \\ = w(r - 2y) + y, \end{aligned} \quad (\text{B2})$$

where $w = \frac{\partial y}{\partial r}$. Recall that the derivative is being evaluated where $r = 0$, and so we can argue that

$$\begin{aligned} \frac{\partial w}{\partial t} + 2yw &= y \rightarrow \\ \mu &= \exp \left(\int \frac{2y_0 dt}{1+ty_0} \right) \\ &= \exp[2 \log(1+ty_0)] \\ &= (1+ty_0)^2. \end{aligned} \quad (\text{B3})$$

Using our integration factors, we see

$$\begin{aligned} w &= \frac{C + \int (1+ty_0)^2 \frac{y_0}{1+ty_0} dt}{(1+ty_0)^2} \\ &= \frac{C + y_0 t (1 + \frac{y_0 t}{2})}{(1+ty_0)^2} \rightarrow C = 0 \\ &= \frac{y_0 t (2 + y_0 t)}{2(1+ty_0)^2} = \frac{\partial y}{\partial r}. \end{aligned} \quad (\text{B4})$$

Note that in the limit that $t \rightarrow \infty$, this expression is order 0 for t ; therefore, unlike the other bifurcation classes, transcriticals are expected to have a *relevant*, rather than a hyperrelevant, leading eigenvalue. This was confirmed with simulations (see Fig. 7).

2. Partial derivative of α_1

The derivative can be set up as:

$$\begin{aligned} \frac{\partial}{\partial \alpha_1} \left(\frac{\partial y}{\partial t} = -y^2 + \alpha_1 y^3 \right) \frac{\partial^2 y}{\partial \alpha_1 \partial t} \\ = -2y \frac{\partial y}{\partial \alpha_1} + 3\alpha_1 y^2 \frac{\partial y}{\partial \alpha_1} + y^3 \frac{\partial w}{\partial t} \\ = -2yw + y^3. \end{aligned} \quad (\text{B5})$$

Since we already know that $\mu = (1+ty_0)^2$, it follows that

$$\begin{aligned} w &= \frac{C + \int (1+ty_0)^2 \left(\frac{y_0}{1+ty_0}\right)^3}{(1+ty_0)^2} \\ &= \frac{C + y_0^2 \int \frac{y_0}{1+ty_0}}{(1+ty_0)^2} \\ &= \frac{C + y_0^2 \log(1+ty_0)}{(1+ty_0)^2} \rightarrow C = 0 \\ \frac{\partial y}{\partial \alpha_1} &= \frac{y_0^2 \log(1+ty_0)}{(1+ty_0)^2}. \end{aligned} \quad (\text{B6})$$

3. Partial derivative of higher-order α 's

Using similar arguments, we arrive at the conclusion that for α_n where $n > 1$

$$\frac{\partial y}{\partial \alpha_n} = \frac{y_0^{n+1}[(1+ty_0)^{1-n} - 1]}{(1-n)(1+ty_0)^2}. \quad (\text{B7})$$

Plots of the sensitivities suggest that r is the dominant parameter for values of $y_0 < 1$, though exactly where this transition occurs is probably worth investigating.

The top-left entry in the FIM is

$$\begin{aligned} \mathcal{I}_{1,1} &= \left(\frac{\partial y}{\partial r} \right)^2 \\ &= \left[\frac{y_0 t (y_0 t + 2)}{2(y_0 t + 1)^2} \right]^2 \\ &= \frac{y_0^2 t^2 (y_0 t + 1)^2}{4(y_0 t + 1)^4}. \end{aligned} \quad (\text{B8})$$

In the limit $t \rightarrow \infty$, this approaches $\frac{t^4}{r^4}$ which is order $\mathcal{O}(t^0)$. This implies that the leading eigenvector of transcritical bifurcations will be relevant, not hyperrelevant like for all other forms of bifurcations considered here. It is tempting to speculate that the topological interpretation of this quirk in the algebra stems from the unique flow field around transcritical bifurcations. For $r < 0$, the vector field has a negative-positive-negative pattern; for $r > 0$ this negative-positive-negative pattern is duplicated, just with an unstable equilibrium at $y = 0$ which had been stable before. Only at the critical value itself ($r = 0$) is there a topological inhomogeneity. The other bifurcations have fundamentally different flow fields on either side of the critical value, and thus, perhaps, their bifurcation parameters acquire hyperrelevance rather than simply relevance. Further study is needed to prove this conjecture.

Because $\frac{\partial y}{\partial \alpha_i} \rightarrow \mathcal{O}[\log(t) - 2]$ and $\frac{\partial y}{\partial \alpha_n} \rightarrow \mathcal{O}(t^{-1-n})$, simple multiplication shows that all the other entries in the FIM will be of lower order than the top-left.

APPENDIX C: FIM OF PITCHFORK BIFURCATIONS

In the supercritical case, the normal form is

$$\frac{dy}{dt} = ry(t) - y(t)^3 + \alpha_1 y(t)^4 + \alpha_2 y(t)^5 + \dots \quad (\text{C1})$$

and the subcritical case is the same except the sign on the cubic term changes. At the critical value of $\theta_i = 0$, the system reduces to

$$\begin{aligned} \frac{dy}{dt} &= -y^3 \rightarrow -\frac{dy}{y^3} = dt \rightarrow \frac{1}{2y^2} \bigg|_{y_0}^{y(t)} = t \bigg|_0 \\ \frac{1}{y(t)^2} - \frac{1}{y_0^2} &= 2t \rightarrow \frac{1}{y(t)^2} = 2t + \frac{1}{y_0^2} \\ \rightarrow y(t) &= \frac{y_0}{\sqrt{1+2ty_0^2}}. \end{aligned} \quad (\text{C2})$$

Following the same logic, the formula for the subcritical case is

$$y(t) = \frac{y_0}{\sqrt{1-2ty_0^2}}. \quad (\text{C3})$$

Note that this creates a potentially problematic singularity at $t = \frac{1}{2y_0^2}$.

1. Partial derivative of r

Let the α_i 's = 0. The derivative of the normal form with respect to r becomes

$$\begin{aligned} \frac{\partial}{\partial r} \left(\frac{\partial y}{\partial t} = ry - y^3 \right) \frac{\partial^2 y}{\partial r \partial t} \\ = \cancel{r \frac{\partial y}{\partial r}} + y - 3y^2 \frac{\partial y}{\partial r} \frac{\partial w}{\partial t} \\ = y - 3y^2 w, \end{aligned} \quad (\text{C4})$$

where $w = \frac{\partial y}{\partial r}$. Using integration factors $p_1 = 1$, $p_0 = 3y^2$, $q = y$, we see that

$$\begin{aligned} \mu &= \exp \left(\int -3y^2 dt \right) \\ &= \exp \left(- \int \frac{3y_0^2 dt}{1+2y_0^2 t} \right) \\ &= \exp \left[\frac{3}{2} \ln(1+2y_0^2 t) \right] \\ &= (1+2y_0^2 t)^{3/2}. \end{aligned}$$

Therefore,

$$\begin{aligned} w &= \frac{C + \int \mu y(t) dt}{\mu} \\ &= \frac{\mathcal{C} + \int \frac{y_0}{\sqrt{1+2ty_0^2}} (1+2y_0^2 t)^{3/2} dt}{(1+2y_0^2 t)^{3/2}} \\ \frac{\partial y}{\partial r} &= \frac{y_0 t (1+2y_0^2 t)}{(1+2y_0^2 t)^{3/2}}. \end{aligned} \quad (\text{C5})$$

Following the same logic for the subcritical case eventually brings us to

$$\frac{\partial y}{\partial r} = \frac{ty_0(1-ty_0^2)}{(1-2ty_0^2)^{3/2}}. \quad (\text{C6})$$

2. Partial derivative of α 's

When $r = 0$, and all $\alpha_{i \neq n} = 0$, then the normal form reduces to

$$\frac{dy}{dt} = -y(t)^3 + \alpha_n y(t)^{n+3}, \quad (\text{C7})$$

which conveniently allows us to use the same μ integration factor as above. Using the integration scheme outlined there, after many steps we reach the conclusion that

$$\frac{\partial y}{\partial \alpha_n} = \frac{y_0^{n+1}}{2-n} \frac{(1+2ty_0^2)^{1-n/2} - 1}{\mu}. \quad (\text{C8})$$

This produces an obvious problem when $n = 2$, but in that case the integration step simplifies and we find that

$$\frac{\partial y}{\partial \alpha_2} = \frac{y_0^3 \ln(1+2ty_0^2)}{2\mu}. \quad (\text{C9})$$

All this indicates that in the FIM, the entry corresponding to $(\partial y / \partial r)^2$ is $\mathcal{O}(t^1)$, while all other entries are lower order, so r will be the only hyperrelevant direction.

APPENDIX D: FIM OF HOPF BIFURCATIONS

Analysis of the Hopf bifurcation in either the complex or Cartesian formulation is complicated, because the introduction of nuisance parameters to the normal form equations tends to alter the period of limit cycles. This means standard trigonometric functions would also need to be altered with time-dependent terms to dilate or expand the period for a closed-form solution of the trajectories $z(t)$ or $x(t), y(t)$, respectively.

However, reparameterizing the equation into polar coordinate form simplifies matters greatly. The system $\dot{r} = r(\mu - r^2)$; $\dot{\theta} = -1$ should look familiar, as the equation for r is simply the normal form for a supercritical pitchfork bifurcation. Therefore, deriving the elements of its Fisher information matrix has already been performed in Appendix C, albeit with different variable and parameter names.

[1] P. Hartman, *Ordinary Differential Equations* (SIAM, Philadelphia, PA, 1982), p. 632.

[2] J. D. Crawford, Introduction to bifurcation theory, *Rev. Mod. Phys.* **63**, 991 (1991).

[3] E. Brown, J. Gao, P. Holmes, R. Bogacz, M. Gilzenrat, and J. D. Cohen, Simple neural networks that optimize decisions, *Int. J. Bifurcat. Chaos* **15**, 803 (2005).

[4] S. Feng, P. Holmes, A. Rorie, and W. T. Newsome, Can monkeys choose optimally when faced with noisy stimuli and unequal rewards?, *PLoS Comput. Biol.* **5**, e1000284 (2009).

[5] I. M. Sobol', Sensitivity estimates for nonlinear mathematical models, *Math. Model. Civil Eng.* **1**, 407 (1993).

[6] B. Bettonvil and J. P. Kleijnen, Searching for important factors in simulation models with many factors: Sequential bifurcation, *Eur. J. Oper. Res.* **96**, 180 (1997).

[7] T. Homma and A. Saltelli, Importance measures in global sensitivity analysis of nonlinear models, *Reliabil. Eng. Syst. Safety* **52**, 1 (1996).

[8] R. Gul and S. Bernhard, Parametric uncertainty and global sensitivity analysis in a model of the carotid bifurcation: Identification and ranking of most sensitive model parameters, *Math. Biosci.* **269**, 104 (2015).

[9] A. Berezhkovskii and A. Szabo, One-dimensional reaction coordinates for diffusive activated rate processes in many dimensions, *J. Chem. Phys.* **122**, 10.1063/1.1818091 014503 (2005).

[10] M. J. Feigenbaum, Quantitative universality for a class of nonlinear transformations, *J. Stat. Phys.* **19**, 25 (1978).

[11] M. J. Feigenbaum, The universal metric properties of nonlinear transformations, *J. Stat. Phys.* **21**, 669 (1979).

[12] M. Widom and L. P. Kadanoff, Renormalization group analysis of bifurcations in area-preserving maps, *Physica D* **5**, 287 (1982).

[13] B. Hu and J. Rudnick, Exact solutions to the Feigenbaum renormalization-group equations for intermittency, *Phys. Rev. Lett.* **48**, 1645 (1982).

[14] J. Hu, Renormalization, Rigidity, and Universality in Bifurcation Theory, Ph.D. thesis, City University of New York, New York (1995).

[15] D. Hathcock and J. P. Sethna, Reaction rates and the noisy saddle-node bifurcation: Renormalization group for barrier crossing, *Phys. Rev. Res.* **3**, 013156 (2021).

[16] L.-N. Chang and N.-P. Chang, Bifurcation and dynamical symmetry breaking in a renormalization-group-improved field theory, *Phys. Rev. Lett.* **54**, 2407 (1985).

[17] A. Raju, C. B. Clement, L. X. Hayden, J. P. Kent-Dobias, D. B. Liarte, D. Z. Rocklin, and J. P. Sethna, Normal form for renormalization groups, *Phys. Rev. X* **9**, 021014 (2019).

[18] R. E. DeVille, A. Harkin, M. Holzer, K. Josić, and T. J. Kaper, Analysis of a renormalization group method and normal form theory for perturbed ordinary differential equations, *Physica D* **237**, 1029 (2008).

[19] M. K. Transtrum, B. B. Machta, K. S. Brown, B. C. Daniels, C. R. Myers, and J. P. Sethna, Perspective: Sloppiness and emergent theories in physics, biology, and beyond, *J. Chem. Phys.* **143**, 010901 (2015).

[20] M. K. Transtrum, B. B. Machta, and J. P. Sethna, Why are nonlinear fits to data so challenging? *Phys. Rev. Lett.* **104**, 060201 (2010).

[21] M. K. Transtrum and P. Qiu, Bridging mechanistic and phenomenological models of complex biological systems *PLoS Comput. Biol.* **12**, e1004915 (2016).

[22] B. B. Machta, R. Chachra, M. Transtrum, and J. P. Sethna, Parameter space compression underlies emergent theories and predictive models, *Science* **342**, 604 (2013).

[23] K. S. Brown, C. C. Hill, G. A. Calero, C. R. Myers, K. H. Lee, J. P. Sethna, and R. A. Cerione, The statistical mechanics of complex signaling networks: Nerve growth factor signaling, *Phys. Biol.* **1**, 184 (2004).

[24] M. K. Transtrum, B. B. Machta, and J. P. Sethna, Geometry of nonlinear least squares with applications to sloppy models and optimization, *Phys. Rev. E* **83**, 036701 (2011).

[25] K. N. Quinn, M. C. Abbott, M. K. Transtrum, B. B. Machta, and J. P. Sethna, Information geometry for multiparameter models: New perspectives on the origin of simplicity, *Rep. Prog. Phys.* **86**, 035901 (2023).

[26] M. K. Transtrum and P. Qiu, Model reduction by manifold boundaries, *Phys. Rev. Lett.* **113**, 098701 (2014).

[27] H. H. Mattingly, M. K. Transtrum, M. C. Abbott, and B. B. Machta, Maximizing the information learned from finite data selects a simple model, *Proc. Natl. Acad. Sci. USA* **115**, 1760 (2018).

[28] A. Raju, B. B. Machta, and J. P. Sethna, Information loss under coarse graining: A geometric approach, *Phys. Rev. E* **98**, 052112 (2018).

[29] J. E. Jeong, Q. Zhuang, M. K. Transtrum, E. Zhou, and P. Qiu, Experimental design and model reduction in systems biology, *Quant. Biol.* **6**, 287 (2018).

[30] E. Roesch and M. P. Stumpf, Parameter inference in dynamical systems with co-dimension 1 bifurcations, *R. Soc. Open Sci.* **6**, 10.1098/rsos.190747, 190747 (2019).

[31] S. H. Strogatz, *Nonlinear Dynamics and Chaos: With Applications to Physics, Biology, Chemistry, and Engineering*, 2nd ed. (Westview Press, Boulder, CO, 2015), p. 513.

[32] O. Jiménez-Ramírez, E. J. Cruz-Domínguez, M. A. Quiroz-Juárez, J. L. Aragón, and R. Vázquez-Medina, Experimental detection of Hopf bifurcation in two-dimensional dynamical systems, *Chaos Solitons Fractals: X* **6**, 100058 (2021).

[33] J. P. C. Kleijnen, B. Bettonvil, and F. Persson, Screening for the important factors in large discrete-event simulation models: Sequential bifurcation and its applications, *Screening: Methods Exp. Industr. Drug Discov. Genet.* **287** (2006).

[34] S. Waldherr and F. Allgower, A feedback approach to bifurcation analysis in biochemical networks with many parameters, *Proceedings of the 2nd Foundations of Systems Biology in Engineering (FOSBE) 2007* (Fraunhofer IRB Verlag, Stuttgart, 2007), pp. 479–484.

[35] A. F. Brouwer and M. C. Eisenberg, The underlying connections between identifiability, active subspaces, and parameter space dimension reduction, [arXiv:1802.05641](https://arxiv.org/abs/1802.05641).

[36] X. Song, B. A. Bryan, A. C. Almeida, K. I. Paul, G. Zhao, and Y. Ren, Time-dependent sensitivity of a process-based ecological model, *Ecol. Model.* **265**, 114 (2013).

[37] A. Alexanderian, P. A. Gremaud, and R. C. Smith, Variance-based sensitivity analysis for time-dependent processes, *Reliabil. Eng. Syst. Safety* **196**, 106722 (2020).

[38] T. Sumner, E. Shephard, and I. D. Bogle, A methodology for global-sensitivity analysis of time-dependent outputs in systems biology modelling, *J. R. Soc. Interface* **9**, 2156 (2012).

[39] I. Bendixson, Sur les courbes définies par des équations différentielles, *Acta Math* **24**, 1 (1901).

[40] B. L. Francis and M. K. Transtrum, Unwinding the model manifold: Choosing similarity measures to remove local minima in sloppy dynamical systems, *Phys. Rev. E* **100**, 012206 (2019).

[41] I. J. Pérez-Arriaga, G. C. Verghese, and F. C. Schweppe, Selective modal analysis with applications to electric power systems, part I: Heuristic introduction, *IEEE Trans. Power Apparatus. Syst. PAS-101*, 3117 (1982).

[42] F. Garofalo, L. Iannelli, and F. Vasca, Participation factors and their connections to residues and relative gain array, *IFAC Proc. Vol.* **35**, 125 (2002).

[43] E. E. Sel'kov, Self-oscillations in glycolysis 1. A simple kinetic model, *Eur. J. Biochem.* **4**, 79 (1968).

[44] A. N. Tikhonov, On the dependence of the solutions of differential equations on a small parameter, *Mat. Sbornik. Nov. Ser.* **22**, 193 (1948).

[45] B. Chance, G. Williamson, I. Lee, L. Mela, D. DeVault, A. Ghosh, and E. Pye, Synchronization phenomena in oscillations of yeast cells and isolated mitochondria, in *Biological and Biochemical Oscillators* (Academic Press, New York, 1973), pp. 285–300.

[46] J. Briggs and F. D. Peat, *Turbulent Mirror* (Harper & Row, New York, 1989), p. 222.

[47] P. Holmes, Poincaré, celestial mechanics, dynamical-systems theory and “chaos,” *Phys. Rep.* **193**, 137 (1990).

[48] G. Sugihara and R. M. May, Nonlinear forecasting as a way of distinguishing chaos from measurement error in time series, *Nature (Lond.)* **344**, 734 (1990).

[49] C.-H. Hsieh, C. Anderson, and G. Sugihara, Extending nonlinear analysis to short ecological time series, *Am. Natur.* **171**, 71 (2008).

[50] D. L. Hitzl and F. Zele, An exploration of the hénón quadratic map, *Physica D* **14**, 305 (1985).

[51] O. E. Rössler, An equation for continuous chaos, *Phys. Lett. A* **57**, 397 (1976).

[52] S. Bleher, E. Ott, and C. Grebogi, Routes to chaotic scattering, *Phys. Rev. Lett.* **63**, 919 (1989).

[53] Y.-C. Lai, Abrupt bifurcation to chaotic scattering with discontinuous change in fractal dimension, *Phys. Rev. E* **60**, R6283 (1999).

[54] J. Guckenheimer and P. Holmes, *Nonlinear Oscillations Dynamical Systems, and Bifurcations of Vector Fields* (Springer, New York, 1983) p. 497.

[55] R. Chachra, M. K. Transtrum, and J. P. Sethna, Structural susceptibility and separation of time scales in the van der Pol oscillator, *Phys. Rev. E* **86**, 026712 (2012).

[56] S. N. Rasband, *Chaotic Dynamics of Nonlinear Systems*, 2nd ed. (Wiley, New York, 1990) p. 230.

[57] D. A. Rand, A. Raju, M. Sáez, F. Corson, and E. D. Siggia, Geometry of gene regulatory dynamics *Proc. National Acad. Sci.* **118**, e2109729118 (2021).

[58] P. D. Kirk, T. Toni, and M. P. Stumpf, Parameter inference for biochemical systems that undergo a Hopf bifurcation, *Biophys. J.* **95**, 540 (2008).

[59] <https://github.com/oceanchaos/TWIG>.

[60] S. Hassani, *Mathematical Physics: A Modern Introduction to Its Foundations*, 2nd ed. (Springer, Heidelberg, 2013), p. 1205.