

RESEARCH ARTICLE

Are circadian amplitudes and periods correlated? A new twist in the story [version 1; peer review: 3 approved]

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V1 First published: 31 Aug 2023, **12**:1077

https://doi.org/10.12688/f1000research.135533.1

Latest published: 31 Aug 2023, 12:1077 https://doi.org/10.12688/f1000research.135533.1

Abstract

Three parameters are important to characterize a circadian and in general any biological clock: period, phase and amplitude. While circadian periods have been shown to correlate with entrainment phases, and clock amplitude influences the phase response of an oscillator to pulse-like zeitgeber signals, the co-modulations of amplitude and periods, which we term twist, have not been studied in detail. In this paper we define two concepts: parametric twist refers to amplitude-period correlations arising in ensembles of self-sustained clocks in the absence of external inputs, and phase space twist refers to the co-modulation of an individual clock's amplitude and period in response to external zeitgebers. Our findings show that twist influences the interaction of oscillators with the environment, facilitating entrainment, fastening recovery to pulse-like perturbations or modifying the response of an individual clock to coupling. This theoretical framework might be applied to understand the emerging properties of other oscillating systems.

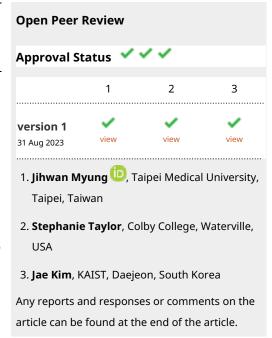
Keywords

circadian clocks, mathematical modeling, amplitudes, periods, twist, heterogeneity, entrainment, coupling



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This article is included in the Circadian Clocks in

Health and Disease collection.

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Competing interests: No competing interests were disclosed.

Grant information: This study was supported by Deutsche Forschungsgemeinschaft (DFG, German Research Foundation) Project-ID 278001972 – TRR 186 to H.H., A.K., C.G. and M.dO.; SCHM 3362/2-1 as well as SCHM 3362/4-1, project-IDs 414704559 and 511886499 to C.S.; RTG2424 CompCancer to S.G.

The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

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How to cite this article: del Olmo M, Schmal C, Mizaikoff C *et al.* Are circadian amplitudes and periods correlated? A new twist in the story [version 1; peer review: 3 approved] F1000Research 2023, 12:1077 https://doi.org/10.12688/f1000research.135533.1

First published: 31 Aug 2023, 12:1077 https://doi.org/10.12688/f1000research.135533.1

Introduction

Oscillations are happening all around us, from the vibrating atoms constituting matter to the beating of the animal heart or to circadian clocks present in all kingdoms of life. Circadian clocks are autonomous clocks that tick in the absence of external timing cues with a period of about 24 h and regulate our behavior, physiology and metabolism. A fundamental property of circadian clocks is that their phase and periodicity can be adjusted to external timing signals (zeitgebers) in a process known as entrainment. It is believed that natural selection has acted on the phase relationship between biological rhythms and the environmental cycle, and thus, this phase of entrainment is of central importance for the fitness of the organism, allowing it to anticipate to changes in the external world.

Three key properties of a circadian rhythm are its period, amplitude and phase. Chronobiological studies have usually focused on period because there are established tools that allow their direct measurement, including running wheels for mice or race tubes for fungi. Phase, which refers to the position of a point on the oscillation cycle relative to a reference point, can be measured similarly. Circadian rhythms can be entrained to various zeitgeber periods T as reviewed in, but under the natural conditions of T = 24 h, variations of intrinsic periods lead to different phases of entrainment that are the basis of chronotypes: faster running clocks (shorter endogenous periods) lead to early phases ('morning larks') and slower clocks (longer periods) correspond to later phases ('night owls'). 3-7

Measuring amplitudes, however, is less straightforward. Some studies have considered activity recordings^{8,9} or conidiation in race tubes, ¹⁰ but one might argue that these measures do not represent the clockwork's amplitude, as they reflect outputs of the circadian system. Other studies have quantified gene expression profiles after careful normalization, ¹¹ but from the ~20 core clock genes that constitute the mammalian circadian oscillator, ¹² it is not immediately evident which gene or protein is best representing the core clock amplitude. Actually, reporter signals monitoring expression of different clock genes and proteins have been used to quantify amplitudes. ^{4,13,14} Others have approached the amplitude challenge indirectly by measuring the response of an oscillator to zeitgeber pulses. ^{15–18} While small-amplitude clocks exhibit larger pulse-induced phase shifts and are easier to phase-reset, larger amplitude rhythms display smaller phase shifts, ^{17–22} with evident consequences in the size of the phase response curve⁵ or in jet lag duration. ²³ Amplitudes, together with periods, also govern entrainment ^{23,24} and seasonality. ^{22,23,25} There have been various theoretical and experimental studies showing, for example, how clocks with larger amplitudes display narrower ranges of entrainment than rhythms of lower amplitude, ²⁰ and how the phase of entrainment is modulated by oscillator amplitude. ^{22,24}

Taken together, these observations indicate, firstly, that the phase of entrainment is correlated with the intrinsic period, and secondly, that both phase of entrainment and phase changes in response to perturbations also correlate with oscillator amplitude. This leads to the question of whether amplitudes and periods are also co-modulated and what insights these interdependencies provide about the underlying oscillator. These questions are the focus of this paper. Do faster-running clocks have larger or smaller amplitudes than slower clocks? What are the implications? Experimental observations have provided evidence for both: in a human osteosarcoma cell line in culture, clocks with longer periods display larger amplitudes; but in cells from the choroid plexus, the major producer of cerebrospinal fluid of the central nervous system, clocks with shorter periods are associated with larger amplitudes (scheme in Figure 1). This dependence between periods and amplitudes is what we here refer to as *twist*, also known as *shear* in the literature. This dependence between twist describes oscillators in which amplitude increases are accompanied by a decreasing period (also termed hard oscillators) and *vice versa* for positive twist (soft oscillators).

In this paper we provide definitions for two important concepts: parametric twist and phase space twist. We show, firstly, that nonlinearities can introduce amplitude-period correlations in simple oscillator models. Moreover, clock models of different complexity can reproduce the experimentally-observed positive²⁶ and negative¹⁴ twist effects, with the type of correlation depending on the model, on parameters, as well as on the variable being measured, illustrating the complexity in defining circadian amplitudes. Lastly, we show how twist effects can speed up or slow down zeitgeber-induced amplitude changes in a simple oscillator model. This helps the clock phase adapt and modulate entrainment of oscillators to a periodic signal or their response to coupling. Our results support the use of oscillator theory as a framework to understand emerging properties of circadian clocks with different twist. Moreover, they provide insights into how temporal or spatial phase patterning might arise in coupled networks, as well as how amplitude changes could help in stabilizing the circadian period in the face of temperature changes. Although we focus on circadian clocks, the presented theory can be applied to any other oscillatory system such as cardiac rhythms, flashing fireflies or voice production.

Materials and Methods

Conservative linear and non-linear oscillators: Harmonic vs. Duffing oscillators

In classical mechanics, a mass-spring harmonic oscillator is a system of mass m that, when displaced from its equilibrium position, experiences a restoring force F proportional to the displacement x, namely F = -kx, where k is the spring

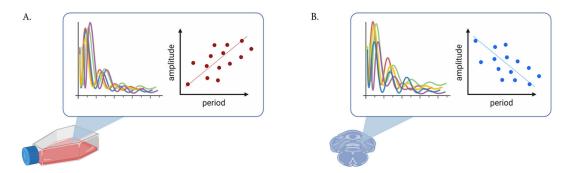


Figure 1. Schematic of experimental observations of circadian amplitude-period correlations. (A) Positive correlations (i.e., positive twist) have been observed in U-2 OS cells kept in culture²⁶; (B) negative correlations have been observed in cells from the choroid plexus in the mouse brain.¹⁴

constant. When F is the only restoring force, the system undergoes harmonic motion (sinusoidal oscillations) around its equilibrium point. In the absence of damping terms, the harmonic motion can be mathematically described by the following linear second order ordinary differential equation (ODE)

$$\frac{\mathrm{d}^2 x}{\mathrm{d}t^2} + \frac{k}{m}x = 0. \tag{1}$$

The solution to this differential equation is given by the function $x(t) = A\cos\left(\frac{2\pi}{\tau} + \phi\right)$, $\frac{29,30}{\tau}$ where A represents the amplitude; ϕ , the phase; and τ represents the period of the motion, $\tau = 2\pi\sqrt{\frac{m}{k}}$. Thus the oscillatory period is determined only by the mass m and the spring constant k. The amplitude A, on the other hand, is determined solely by the starting conditions (by both initial displacement x and velocity $v = \dot{x}$).

Introducing a non-linear term in the restoring force such that $F = -kx - \beta x^3$ allows the conversion of the simple harmonic oscillator into a Duffing oscillator. The equation of β , the coefficient that determines the strength of the non-linear term, the spring is termed *hard* or *soft* oscillator. The equation of the Duffing oscillator, in the absence of damping terms, reads

$$\frac{d^2x}{dt^2} + \frac{k}{m}x + \frac{\beta}{m}x^3 = 0. {2}$$

Due to the non-linear term introduced in equation 2, it is helpful to write this system in a form that can be easily treated via numerical integration. Considering the following change of variable $v = \dot{x}$, the equation can be reformulated as a system of two first order ODEs:

$$\frac{\mathrm{d}x}{\mathrm{d}t} = v$$

$$\frac{\mathrm{d}v}{\mathrm{d}t} = -\frac{k}{m}x - \frac{\beta}{m}x^{3}.$$
(3)

Goodwin-like models

The three-variable Goodwin model is a minimal model based on a single negative feedback loop that can describe the emergence of oscillations in simple biochemical systems. All synthesis and degradation terms are linear, with the exception of the repression that z exerts on x which is modeled with a sigmoidal Hill curve. The equations that describe the dynamics read

$$\frac{\mathrm{d}x}{\mathrm{d}t} = k_1 \frac{K_1^n}{K_1^n + z^n} - k_2 x$$

$$\frac{\mathrm{d}y}{\mathrm{d}t} = k_3 x - k_4 y$$

$$\frac{\mathrm{d}z}{\mathrm{d}t} = k_5 y - k_6 z,$$
(4)

where k_1 , k_3 and k_5 represent the rates of synthesis of x, y and z, respectively; k_2 , k_4 and k_6 , the degradation rates; and n, the Hill exponent.

The Goodwin model, however, requires a very large Hill exponent (n > 8) to produce self-sustained oscillations,³² which biologists and modelers have often considered unrealistic. Gonze³³ and others have shown that, by introducing additional nonlinearities in the system, the need for such high value of the Hill exponent can be reduced. In contrast to the linear degradation of variables of the original Goodwin model, the Gonze model³³ describes degradation processes with Michaelis-Menten kinetics as follows

$$\frac{dx}{dt} = k_1 \frac{K_1^n}{K_1^n + z^n} - k_2 \frac{x}{K_2 + x}
\frac{dy}{dt} = k_3 x - k_4 \frac{y}{K_4 + y}
\frac{dz}{dt} = k_5 y - k_6 \frac{z}{K_6 + z}.$$
(5)

To mimic clock heterogeneity and evaluate the amplitude-period correlations among ensembles of Gonze or Goodwin oscillators, degradation rates were varied around $\pm 10\%$ their default value (Table 1).

Almeida model

The Almeida model³⁶ is a protein model of the mammalian clockwork that includes 7 core clock proteins along with the PER:CRY complex and the regulation (activatory or repressive) that these exert on DNA binding sites known as clock-controlled elements to regulate circadian gene expression. The regulation at these clock-controlled elements, namely E-boxes, D-boxes and RORE elements, is described by the following terms:

$$\begin{split} \mathbf{E}_{\text{box}} &= V_E \frac{\text{BMAL1}}{\text{BMAL1} + k_E + (k_{Er} \, \text{BMAL1} \, \text{CRY})} \\ \text{RORE} &= V_R \frac{\text{ROR}}{k_R + \text{ROR}} \frac{k_{Rr}^2}{k_{Rr}^2 + \text{REV}^2} \\ \mathbf{D}_{\text{box}} &= V_D \frac{\text{DBP}}{\text{DBP} + k_D} \frac{k_{Dr}}{k_{Dr} + \text{E4BP4}}. \end{split}$$

The system of ODEs that describes the dynamics of the clock proteins in the Almeida model reads

Table 1. Default parameters of the Goodwin and Gonze models used for our simulations. Their biological interpretation as well as the default values from publications are shown.

Parameter	Biological interpretation	Value in Goodwin model ^{34,35}	Value in Gonze model ³³
<i>k</i> ₁	transcription rate of core clock gene <i>x</i>	1 nM	0.7 nM h ⁻¹
k ₂	degradation rate of <i>x</i>	0.2 h ⁻¹	0.35 nM h ⁻¹
k ₃	translation rate of core clock protein <i>y</i>	1 h ⁻¹	$0.7 h^{-1}$
k ₄	degradation rate of <i>y</i>	0.15 h ⁻¹ *	0.35 nM h ⁻¹
k ₅	nuclear import rate of the repressor z	1 h ⁻¹	$0.7 h^{-1}$
<i>k</i> ₆	degradation rate of z	$0.1 h^{-1}$	0.35 nM h ⁻¹
<i>K</i> ₁	Michaelis constant of <i>z</i> -mediated repression	1 nM	1 nM
K ₂	Michaelis constant of <i>x</i> degradation	-	1 nM
K ₄	Michaelis constant of y degradation	-	1 nM
K ₆	Michaelis constant of z degradation	-	1 nM
n	Hill exponent of cooperative inhibition of z on x	9.5 *	4

^{*}Parameters have been adapted to obtain limit cycle oscillations, since the default degradation parameters from³⁵ produced weakly damped rhythms.

$$\frac{dBMAL1}{dt} = RORE - \gamma_{BP} BMAL1 PERCRY$$

$$\frac{dROR}{dt} = E_{box} + RORE - \gamma_{Ror} ROR$$

$$\frac{dREV}{dt} = 2 E_{box} + D_{box} - \gamma_{Rev} REV$$

$$\frac{dDBP}{dt} = E_{box} - \gamma_{Db} DBP$$

$$\frac{dE4BP4}{dt} = 2 R_{box} - \gamma_{E4} E4BP4$$

$$\frac{dCRY}{dt} = E_{box} + 2 R_{box} - \gamma_{PC} PERCRY + \gamma_{CP} PERCRY - \gamma_{C} CRY$$

$$\frac{dPER}{dt} = E_{box} + D_{box} - \gamma_{PC} PERCRY + \gamma_{CP} PERCRY - \gamma_{P} PERCRY$$

All parameter descriptions along with their default values are given in Table 2.

To analyze the twist effects that arise from a population of heterogeneous clocks (i.e., parametric twist), all 18 model parameters were randomly varied around $\pm 20\%$ their default value (Table 2) one at a time. Since this model can lead to period-doubling effects upon changes of certain parameters, ³⁷ those oscillations whose period change resulted in oscillations with period-doubling were removed from the analysis. Moreover, if changing the default parameter in the ensemble resulted in a range of ratio of amplitude variation relative to the default amplitude < 0.1, then we considered that ensemble to have no twist for that particular control parameter. Periods and amplitudes were determined from the peaks and troughs of oscillations using the continuation software XPP-AUTO.

Poincaré model

The intrinsic dynamical properties from single oscillators and their interaction to external stimuli can be very conveniently described by means of a Poincaré model.³⁸ We here propose a modification of its generic formulation that explicitly takes into account twist effects through the phase space twist parameter ϵ . The modified Poincaré model with twist reads

$$\frac{\mathrm{d}r}{\mathrm{d}t} = \lambda r(A - r)$$

$$\frac{\mathrm{d}\phi}{\mathrm{d}t} = \omega + \epsilon (A - r).$$
(7)

Table 2. List of parameters of the Almeida model. Time units are given in hours and concentration units as arbitrary units. The default values are taken from.³⁶

Parameter	Biological interpretation	Value	Parameter	Biological interpretation	Value
V_R	rate of RORE activation	44.4 h ⁻¹	γ_{Ror}	ROR degradation rate	2.55 h ⁻¹
k _R	strength of RORE activation	3.54	γ _{Rev}	REV degradation rate	0.241 h ⁻¹
k _{Rr}	strength of RORE inhibition	80.1	γР	PER degradation rate	0.844 h ⁻¹
V_E	rate of E box activation	30.3 h ⁻¹	γ _C	CRY degradation rate	2.34 h ⁻¹
k _E	strength of E box activation	214	γ_{DB}	DBP degradation rate	0.156 h ⁻¹
k _{Er}	strength of E box inhibition	1.24	γ _E 4	E4BP4 degradation rate	0.295 h ⁻¹
V_D	rate of D box activation	202 h ⁻¹	ΫРС	PER:CRY degradation rate	0.191 h ⁻¹
k_D	strength of D box activation	5.32	ΫСР	PER:CRY formation rate	0.141 h ⁻¹
k _{Dr}	strength of D box inhibition	94.7	γ_{BP}	BMAL1 nuclear export rate	2.58 h ⁻¹

The first equation describes the rate of change of the radial coordinate r(t) (i.e., the time-dependent distance from the origin), whereas the second equation determines the rate of change of the angular coordinate $\phi(t)$, where $\omega = \frac{2\pi}{\tau}$. The parameters τ, A, λ and ϵ denote the free-running period (in units of time), amplitude (arbitrary units), amplitude relaxation rate (in units of time⁻¹) and phase space twist of the oscillator (in units of time⁻¹), respectively. In the absence of twist, namely $\epsilon = 0$, the phase changes constantly along the limit cycle at a rate $\frac{2\pi}{\tau}$, independently of the radius. In the case of $\epsilon \neq 0$, the phase changes at a constant rate only when r = A; if any perturbation is to modify r such that $r \neq A$, then the phase change will be accelerated or decelerated depending on the sign of ϵ and on whether r > A or r < A. The model parameters, unless otherwise specified in the figures or captions, are the following: A = 1 a.u., $\lambda = 0.05$ h⁻¹, $\tau = 24$ h and ϵ values of 0 or ± 0.1 h⁻¹.

To study the effects of twist on the oscillator's response to pulse-like perturbations pert(t), periodic zeitgeber input Z(t) or mean-field coupling M, the individual Poincaré oscillators i were converted into Cartesian coordinates and the respective terms were added in the equations of the x_i variable as follows:

$$\frac{\mathrm{d}x_i}{\mathrm{d}t} = \lambda x_i (A - r_i) - y_i (\omega + \epsilon (A - r_i)) + Z(t) + M + \mathrm{pert}(t)
\frac{\mathrm{d}y_i}{\mathrm{d}t} = \lambda y_i (A - r_i) + x_i (\omega + \epsilon (A - r_i)), \tag{8}$$

where $r_i = \sqrt{x_i^2 + y_i^2}$. The zeitgeber Z(t) is given by:

$$Z(t) = F_Z \cos\left(\frac{2\pi}{T}t + \frac{\pi}{2}\right),\,$$

where T represents the zeitgeber period and F_Z the strength (amplitude) of the zeitgeber input. The mean-field M coupling is given by:

$$M = \frac{K}{N} \sum_{i=1}^{N} x_i(t),$$

where K represents the coupling strength and N is the number of oscillators in the coupled ensemble. Lastly, the square-like perturbation pert(t) is defined as follows:

$$pert(t) = \begin{cases} 0, & \text{if } t_{\text{start}} \le t \le t_{\text{start}} + 1 \\ F_P & \text{otherwise,} \end{cases}$$

where F_P is the strength of the perturbation and t_{start} is the time at which the perturbation starts. The perturbation lasts 1 h and is set to $F_P = 0.7$ a.u. in all our simulations.

Analytical derivation of isochrones

Arthur T. Winfree introduced the concept of isochrones as any set of dynamical states which oscillate with the same phase when they reach the limit cycle at time $t \to \infty$. ³⁹ That asymptotic phase at $t \to \infty$ is what Winfree termed latent phase Φ . Since the dynamical flow of the Poincaré model has polar symmetry, the isochrones must also have polar symmetry such that

$$\Phi = g(\phi, r) \stackrel{!}{=} \phi - f(r). \tag{9}$$

By definition, the latent phase velocity $\dot{\Phi}$ necessarily increases at units of the angular velocity $\omega = \frac{2\pi}{\tau}$ as the oscillator follows its kinetic equation:

$$\frac{\mathrm{d}\Phi}{\mathrm{d}t} \stackrel{!}{=} \omega \tag{10}$$

Combining the two previous equations and with the use of the chain rule, we can calculate the radius dependency of the latent phase Φ :

$$\frac{\mathrm{d}\Phi}{\mathrm{d}t} \stackrel{9}{=} \frac{\mathrm{d}\phi}{\mathrm{d}t} - \frac{\mathrm{d}f(r)}{\mathrm{d}r} \frac{\mathrm{d}r}{\mathrm{d}t} \stackrel{10}{=} \omega$$

The terms $\frac{d\phi}{dt}$ and $\frac{dr}{dt}$ are defined in the Poincaré model (equation 7), so that the equation above can be rewritten as:

$$\omega + \epsilon (A - r) - \frac{\mathrm{d}f(r)}{\mathrm{d}r} \lambda r (A - r) \stackrel{10}{=} \omega$$

Solving for $\frac{df(r)}{dr}$:

$$\frac{\mathrm{d}f(r)}{\mathrm{d}r} = \frac{\epsilon(A-r)}{\lambda r(A-r)} = \frac{\epsilon}{\lambda r}$$

Next, the solution for f(r) can be found through integration:

$$f(r) = \frac{\epsilon}{\lambda} \int_{r=A}^{r} \frac{1}{r} dr = \frac{\epsilon}{\lambda} \ln r - \frac{\epsilon}{\lambda} \ln A$$

If A = 1 (like in all our simulations), the last term can be neglected such that

$$f(r) = \frac{\epsilon}{\lambda} \ln r$$

Finally, the solution of f(r) can be inserted in equation 9 to end up with the equation for isochrones as a function of the radius:

$$\Phi = \phi - \frac{\epsilon}{\lambda} \ln r \tag{11}$$

Isochrones are thus loci of polar coordinates (ϕ, r) in phase space with the same latent phase Φ . Equation 11 shows how the shape of the isochrone for the Poincaré model in equation 7 depends on the ratio of twist ϵ to relaxation rate λ . Specifically, higher ϵ values (in modulus) and lower relaxation rates λ lead to more curved isochrones.

To plot the isochrones, we simply reorder equation 11 to plot ϕ as a function of r at isochrones with fixed values of latent phases $\Phi = \left\{0, \frac{\pi}{4}, \frac{\pi}{7}, \frac{3\pi}{4}, \pi, -\frac{3\pi}{4}, -\frac{\pi}{2}, -\frac{\pi}{4}\right\}$:

$$\phi = \frac{\epsilon}{\lambda} \ln r - \Phi$$

Computer simulations and data analysis

All numerical simulations were performed and analyzed in Python with the numpy, scipy, pandas and astropy libraries. The function odeint from scipy was used to numerically solve all ordinary differential equations. Bifurcation analyses were computed in XPP-AUTO⁴⁰ using the parameters $N_{tst} = 150, N_{max} = 20000, Ds_{min} = 0.0001$ and $Ds_{max} = 0.0002$.

Throughout our analyses, periods were determined by (i) normalizing the solutions to their mean, (ii) centering them around 0 (by subtracting one unit from the normalized solution), and (iii) by then computing the zeroes of the normalized rhythms. The period was defined as the distance between two consecutive zeros with a negative slope. Amplitudes were determined as the average peak-to-trough distance of the last (normalized) oscillations after removing transients.

Results

Nonlinearities can result in amplitude-period correlations across oscillators

The simple harmonic oscillator (Figure 2A) is a classical model of a system that oscillates with a restoring force proportional to its displacement, namely F = -kx. The ordinary differential equation that describes the motion of a mass attached on a spring is linear (equation 1 in Materials and Methods) and the solution can be found analytically. The period is determined by the size of the mass m and the force constant k (see Materials and Methods), while the amplitude and phase are determined by the starting position and the velocity. Thus, an ensemble of harmonic oscillators with different initial conditions will produce results that differ in amplitudes but whose periods are the same (Figure 2B, C). When plotting amplitudes against periods, no correlation or *twist* is observed: the period of a simple harmonic oscillator is independent of its amplitude (Figure 2D).

The simple harmonic oscillator can be converted into a Duffing oscillator³¹ by including a cubic nonlinearity in its equation. In the Duffing oscillator (equations 2, 3 in Materials and Methods), the restoring force is no longer linear (see deformed springs in Figure 2) but instead described by $F = -kx - \beta x^3$, where β represents the coefficient of non-linear elasticity. These classical conservative oscillators instead are known to have an amplitude-dependent period. A network of Duffing clocks with different starting conditions will produce oscillations of different amplitudes as well as periods. Duffing oscillators with a negative cubic term have been termed soft oscillators and display periods that grow with amplitudes (Figure 2E–H), similar to Kepler's Third Law of planetary motion, ⁴² where planets with larger distances to the Sun run at slower periods than those that are closer. On the other hand, Duffing oscillators with a positive β term are known as hard oscillators and show negative twist (amplitude-period correlations) (Figure 2I–L). ^{43,44}

In short, nonlinearities in oscillator models can introduce twist effects among ensembles of oscillators with slight differences in their properties (initial conditions, parameters, etc.). Thus, models for the circadian clock, which are based on nonlinearities, are expected to show amplitude-period correlations.

Oscillator heterogeneity produces parametric twist effects in limit cycle clock models

Most circadian clock models generate stable limit cycle oscillations. Limit cycles are isolated closed periodic orbits with a given amplitude and period, where neighboring trajectories (e.g. perturbations applied to the cycle) spiral either towards or out of the limit cycle. Stable limit cycles are examples of attractors: they imply self-sustained oscillations.

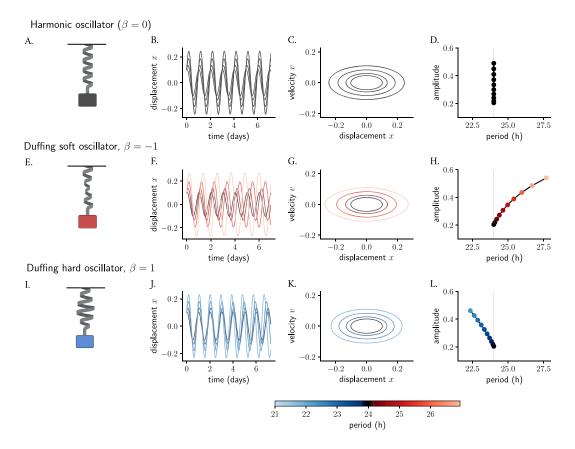


Figure 2. Nonlinearities introduce amplitude-period correlations in oscillator models. The mass-spring harmonic oscillator (A) is a linear oscillator that, in the absence of friction, produces persistent undamped oscillations (B, in time series and C, in phase space). Different starting conditions produce oscillations that differ in amplitudes but whose period is independent of amplitude (D). The harmonic oscillator is converted to a Duffing oscillator (E, I) by introducing nonlinearities in the restoring force of the spring (the non-linear behavior is represented with deformations of the spring). The new restoring force results in slight changes in the initial conditions producing oscillations whose amplitude and period change (F, G, J, K) and become co-dependent (H, L) and where the correlations depend on the sign of the non-linear term. The terms soft and hard oscillators refer to, by convention, those with positive and negative twist, respectively. Period values in (F–H, J–L) are color-coded.

The closed trajectory describes the perfect periodic behavior of the system, and any small perturbation from this trajectory causes the system to return to it, to be *attracted back* to it.

Limit cycles are inherently non-linear phenomena and they cannot occur in a linear system (i.e., a system in the form of $\vec{x} = A\vec{x}$, like the harmonic oscillator). From the previous section we have learned that non-linear terms can introduce amplitude-period co-dependencies. In this section we show that kinetic limit cycle models of the circadian clock also show twist effects. Instead of studying the amplitude-period correlation of an oscillator model with fixed parameters and changing initial conditions as in Figure 2 (as all initial conditions would be attracted to the same limit cycle), we study here the correlations that arise among different uncoupled oscillators due to oscillator heterogeneity (i.e., differences in biochemical parameters), as found experimentally. We refer to amplitude-period correlations that arise due to oscillator heterogeneity as parametric twist.

Single negative feedback loop models

The Goodwin model is a simple kinetic oscillator model 46 that is based on a delayed negative feedback loop, where the final product of a 3-step chain of reactions inhibits the production of the first component (equation 4 in Materials and Methods). In the context of circadian rhythms, 34,35 the model can be interpreted as a clock activator x that produces a clock protein y that, in turn, activates a transcriptional inhibitor z that represses x (Figure 3A). The Goodwin model has been extensively studied and fine-tuned by Gonze and others 33 to study fundamental properties of circadian clocks $^{47-50}$ or synchronization and entrainment. 33,51,52 The Gonze model 33 (equation 5) includes additional nonlinearities, where the degradation of all 3 variables is modeled with non-linear Michaelis Menten kinetics, to reduce the need of very large Hill exponents (n > 8), required in the original Goodwin model to generate self-sustained oscillations, 32 that have been questioned to be biologically meaningful. These Michaelian degradation processes can be interpreted as positive feedback loops which aid in the generation of oscillations 53 (Figure 3B).

To study whether twist effects are present in an ensemble of 100 uncoupled Goodwin and Gonze oscillators with different parameter values, we randomly varied the degradation parameters of x, y or z (k_2 , k_4 and k_6 , respectively) in each oscillator

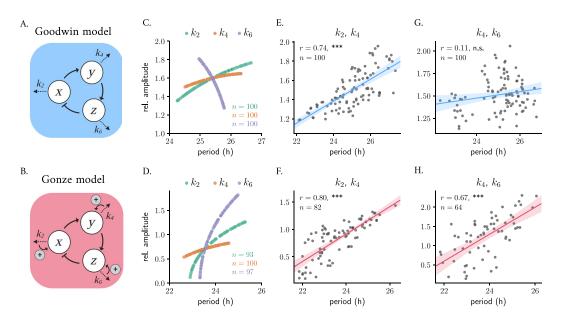
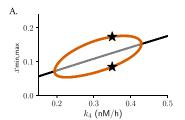
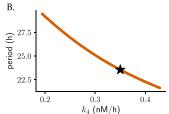


Figure 3. Parametric twist in an ensemble of 100 uncoupled Goodwin-like models: correlations are model- and parameter-dependent. Scheme of the Goodwin (A) and Gonze (B) oscillator models. Changes in degradation parameters result in twist effects when varied randomly around $\pm 10\%$ their default parameter value, mimicking oscillator heterogeneity of the Goodwin (C) and Gonze (D) models. Simultaneous co-variation of degradation parameters results in parametric twist: changes of k_2 and k_4 produce significant positive amplitude-period correlations in both models (E, F); co-variations of k_6 and k_4 result in amplitude-period correlations that are not significant in the Goodwin model (G), but significant positive parametric twist in the Gonze model (H). Shown in all panels is Spearman's R and the significance of the correlation (n. s.: not significant; ***: p value<0.001). Default model parameters are summarized in Table 1. Relative amplitudes are computed as the peak-to-trough distance of the x variable. Oscillators from the ensemble whose amplitude was < 0.1 have been removed from the plots; the total number of oscillators (n) is indicated in the plots.





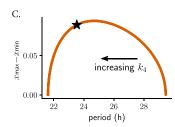


Figure 4. The type of parametric twist depends on the parameter location within the Hopf bubble. Bifurcation analysis of Gonze oscillators with changing k_4 , simulating heterogeneous oscillators: (A) Hopf bubble representing the peaks (maxima) and troughs (minima) of variable x as a function of changes in the degradation rate of y (k_4); (B) monotonic period decrease for increasing k_4 . Represented in (C) is the combination of (A) and (B), namely how the difference between the maxima and minima of x changes with the period: a non-monotonic behavior of parametric twist is observed, which depends on the values of k_4 . Note that, since k_4 results in a decrease in period length, the k_4 dimension is included implicitly in (C) and k_4 increases from the right to the left of the plot. Stars indicate the default parameter value, period or amplitude.

around $\pm 10\%$ their default parameter value (given in Table 1) and analyzed the resulting amplitude-period correlation. The resulting parametric twist depends on the model and on the parameter being changed: in the Goodwin model, variations in the degradation rate of the transcriptional activator (k_2) or of the clock protein (k_4) result in positive twist effects (i.e., soft twist-control), while changes in the transcriptional repressor's degradation rate (k_6) result in a hard twist-control, i.e., negative amplitude-period correlation (Figure 3C). In the Gonze model, k_2 , k_4 and k_6 are all soft twist-control parameters, as individual changes of any of them all produce positive parametric twist effects (Figure 3D).

To mimic cell-to-cell variability in a more realistic manner, we introduced heterogeneity by changing combinations of the degradation parameters simultaneously around $\pm 10\%$ their default parameter value and analyzing the resulting periods and amplitudes. We observed that the overall twist behavior depends on the particular influence that each parameter has individually. Random co-variations of k_2 and k_4 produce significant positive parametric twist effects in ensembles of both Goodwin (Figure 3E) or Gonze clocks (Figure 3F), consistent with the positive correlations when either of the parameters is changed individually (Figure 3C, D). Nevertheless, when k_6 , which has a negative twist effect in the Goodwin model, is changed at the same time as k_4 (or k_2 , data not shown) the correlation is not significant (Figure 3G). Random co-variation of k_4 and k_6 in an ensemble of Gonze clocks results in significant positive parametric twist effects (Figure 3H).

Changes in parameter values can result in significant alterations to a system's long-term behavior, which can include differences in the number of steady-states, limit cycles, or their stability properties. Such qualitative changes in non-linear dynamics are known as bifurcations, with the corresponding parameter values at which they occur being referred to as bifurcation points. In oscillatory systems, Hopf bifurcations are an important type of bifurcation point. They occur when a limit cycle arises from a stable steady-state that loses its stability. A 1-dimensional bifurcation diagram illustrates how changes in a mathematical model's control parameter affect its final states, for example the period or amplitude of oscillations. The Hopf bubble refers to the region in parameter space where the limit cycle exists, and it is commonly represented with the peaks and troughs of a measured variable in the y axis, with the control parameter plotted on the x axis (Figure 4A). The term "bubble" is used because of the shape of the curve, that resembles a bubble that grows or shrinks as the parameter is changed. Such bifurcation analyses can be used to predict the type of twist that a system shows: the amplitude will increase with parameter changes if the default parameter value is close to the opening of the Hopf bifurcation, or will decrease if the value is near the closing of the bubble.

We observed that Gonze oscillators display self-sustained oscillations for values of k_4 between 0.2 and 0.43 (Figure 4A) and that, for increasing k_4 , periods decrease monotonically (Figure 4B). For oscillators with increasing k_4 , the parametric twist effects from the ensemble are first negative (amplitudes increase and periods decrease); however, when Gonze clocks have k_4 values that are part of the region of the bubble where amplitudes decrease, positive amplitude-period correlations appear in the ensemble (Figure 4C). In summary, the type of parametric twist depends on both the model and the parameter being studied but also on where the parameter is located within the Hopf bubble. Other more complex kinetic models of the mammalian circadian clockwork have shown that changes in some of the degradation parameters produce non-monotonic period changes,⁵⁴ and thus the twist picture is expected to become even more complex.

Model with synergistic feedback loops

The Almeida model³⁶ is a more detailed model of the core clock network in mammals that includes seven core clock proteins that exert their regulation at E-boxes, D-boxes and ROR binding elements (RORE) through multiple positive and

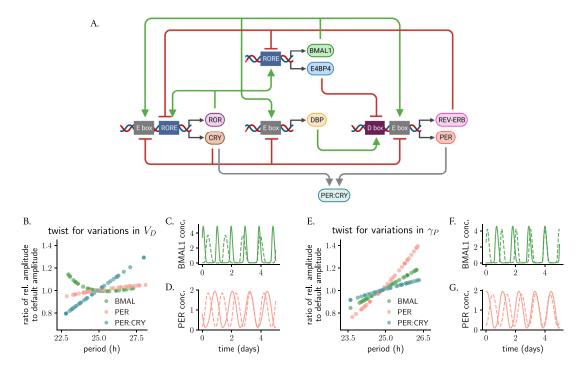


Figure 5. The type of parametric twist in models of the circadian clockwork with synergistic feedback loops depends on the parameter being studied and on the variable that is measured. (A) Scheme of the Almeida model, ³⁶ illustrating activation (green arrows) and inhibition (red arrows) of different clock-controlled elements (at E-boxes, D-boxes and ROR elements) by the respective core clock proteins. Parametric twist was studied by modeling a heterogeneous uncoupled ensemble where model parameters were randomly and individually varied around $\pm 20\%$ of their default value (shown in Table 2) and assessing the period-amplitude correlation of the ensemble. (B–D) Almeida oscillators (n=40) with heterogeneous values in the activation rate of D-boxes (V_D): the twist is negative from the perspective of BMAL1, positive from the perspective of PER:CRY, but the amplitude of PER is not greatly affected by changing V_D . (E–G) Almeida oscillators (n=40) with heterogeneous values in the degradation rate of PER (γ_P) show positive parametric twist effects for BMAL1, PER and PER:CRY. The twist in panels (B, E) is evaluated by comparing the relative amplitude of the respective protein/protein complex (computed as the average peak-to-trough distance) after parameter change to the default amplitude. Shown in (C, D, F, G) are representative oscillations with long and short periods: dashed lines represent rhythms of smallest amplitudes.

negative feedback loops (Figure 5A, equation 6 in Materials and Methods). To study parametric twist in an ensemble of uncoupled Almeida oscillators, we randomly changed all 18 parameters individually around $\pm 20\%$ their default value (Table 2 in Materials and Methods) and computed the amplitude-period correlations. In this case, we calculated the ratio of amplitude variation after the parameter change relative to the default amplitude.

We found, interestingly, that the overall twist effects depend not only on the parameter being studied, but also on the variable which is measured. For example, changes in the rate of D-box activation parameter V_D result in negative twist effects for BMAL1 (i.e., lower amplitude BMAL1 rhythms run slower than oscillators with higher amplitude BMAL1 rhythms), positive parametric twist for the PER:CRY complex but almost no parametric twist from the perspective of PER (Figure 5B–D). Changes in PER degradation γ_P produce positive parametric twist for BMAL1, PER and PER:CRY but of different magnitudes (Figure 5E–G). Supplementary Table S1 (in⁵⁷) summarizes the parametric twist effects for the additional parameters from the Almeida model, highlighting the complexity that arises with synergies of feedback loops and bringing us again to the question of defining what the relevant amplitude of a complex oscillator is.

Phase space twist in single oscillators: On amplitude-phase models, isochrones and perturbed trajectories

Up until now, our results have focused on studying amplitude-correlations among ensembles of self-sustained oscillators with differences in their intrinsic properties (e.g. biochemical parameters) in the absence of external cues. However, when a clock is exposed to external stimuli, its amplitude and period undergo adaptation. We refer to these amplitude-period correlations in individual clocks as they return to their steady-state oscillation after a stimulus as *phase space twist*.

In contrast to *parametric twist*, which emerged as a result of a *heterogenous population* of oscillators, here we focus on *individual* clocks.

To further explore the correlation and interdependence between the frequency of an oscillator and its amplitude upon an external stimulus, more generalized models can be of use. The Poincaré oscillator model (equation 7 in Materials and Methods) is a simple conceptual oscillator model with only two variables, amplitude and phase, that has been widely used in chronobiology research. 20,22,24,38,39,55 This amplitude-phase model, regardless of molecular details, can capture the dynamics of an oscillating system and what happens when perturbations push the system away from the limit cycle. When a pulse is applied and an oscillating system is 'kicked out' from the limit cycle, the perturbed trajectory is attracted back to it at a rate λ . Within the limit cycle, the dynamics are strictly periodic: if one takes a point in phase space and observes where the system returns to after exactly one period, the answer is trivial: to exactly the same spot (Figure 6, red dots). Nevertheless, outside the limit cycle, during the transient relaxation time (i.e., time between the perturbation and the moment that the trajectory reaches the limit cycle), the time between two consecutive peaks might be shorter or longer than the period of the limit cycle orbit. We will come back to this point in the next section.

Arthur T. Winfree introduced a practical term, isochrones, to conceptualize timing relations in oscillators perturbed off their attracting cycles, ³⁹ which becomes important in physiological applications because often, biological oscillators are not on their attracting limit cycles. To understand what isochrones are, Winfree proposes in ³⁹ a simple experiment where a pulse-like perturbation applied in a system produces a 'bump' in the limit cycle. As the perturbation relaxes and spirals towards the attracting limit cycle, one records the position of the oscillator in time steps which equal the period of the unperturbed limit cycle. The sequence of points left as a footprint will converge to a fixed point on the cycle and will outline an isochrone (Figure 6). Isochrones are related to oscillator twist, and whereas in a Poincaré oscillator with no twist, the isochrones are straight and radial (Figure 6A), when twist is present, the isochrones become bent or skewed (Figure 6B, C).

In an individual oscillator with positive twist, a perturbation that increases the oscillator's instantaneous amplitude will relax back and intersect the isochrone at an angle $< 360^{\circ}$ (Figure 6B) than an oscillator with no twist in the time of one period, whereas an oscillator with negative twist will cover an angle $> 360^{\circ}$ in the time of one period (Figure 6C). Thus, isochrones and consequently twist illustrate how a perturbation away from the limit cycle is decelerated or accelerated during the course of relaxation.

Mathematically, the 'skewness' of isochrones in an amplitude-phase oscillator is directly related to the twist parameter ϵ , that adds a radius dependency on the phase dynamics. The phase changes at a constant rate $\omega = \frac{2\pi}{\tau}$ in the limit cycle, but in conditions outside the limit cycle, the phase is made to be modulated by the radius through the twist parameter ϵ until

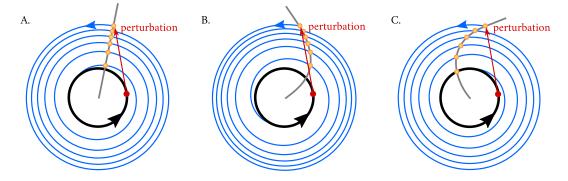


Figure 6. Schematic of an isochrone in a Poincaré oscillator with (A) no twist, (B) positive twist or (C) negative twist. To understand the concept of isochrones, a simple experiment is performed: if one considers a point in phase space (shown in red) within a limit cycle (black) and observes where the system returns to after exactly one period, the answer is trivial: to the same spot. If, however, now a perturbation is applied (red arrow) such that the system starts in a section of the state space which is outside of the stable periodic orbit, and one maps the points that the system 'leaves as a footprint' after exactly one period as it relaxes back to the limit cycle (blue trajectories), these points (shown in yellow) will outline an isochrone (grey line). The twist parameter ϵ affects the curvature of the isochrone, with implications in the response of the oscillating system to the perturbation. The oscillator with positive twist (B) arrives at a later phase than that with no twist (A), resulting in a phase delay with respect the clock with no twist, whereas the clock with negative twist (C) arrives to the limit cycle at an earlier phase (i.e., advanced with respect to the clock with no twist).

r = A (equation 7 in Materials and Methods). This *phase space twist* parameter ϵ now represents the amplitude-period correlations (acceleration/deceleration) as an *individual* oscillator returns to its steady-state oscillation.

Phase space twist in single oscillators affects their interaction with the environment

We have seen how, in an individual oscillator, the twist parameter ϵ acts to slow down (in case of positive twist) or to speed up (in case of negative twist) trajectories further away from the limit cycle compared to those closer to it. As a result, ϵ has a direct consequence on how the individual oscillator responds to pulse-like perturbations or periodic zeitgebers coming from the environment.

To analyze how phase space twist affects the response of an oscillator to a zeitgeber pulse, we applied a pulse-like perturbation to individual Poincaré oscillators with different phase space twist values ϵ . The pulse was applied at CT3 and was made to increase the instantaneous amplitude (such that r > A). If the oscillator has no twist ($\epsilon = 0$, Figure 7A), the isochrones are straight and radial, but for a soft or a hard oscillator with positive or negative ϵ values, respectively, isochrones get skewed (Figure 7B, C, also Figure 6). In the case of a soft oscillator with positive ϵ (Figure 7B), the pulse at CT3 gets decelerated during the course of its relaxation, and as a consequence, the oscillator arrives back to the limit cycle at an *later* phase than the oscillator with $\epsilon = 0$ (Figure 7D, E). The same can also be inferred mathematically: for this particular perturbation where r > A, A - r < 0 holds and hence the phase velocity outside the limit cycle is smaller ($\phi < \omega$) than at the limit cycle for the oscillator with positive twist, since $\phi = \omega + \epsilon(A - r)$ (equation 7). The opposite holds for the oscillator with negative twist: this clock arrives at an *earlier* phase to the limit cycle (Figure 7C, F).

The positive and negative amplitude-period correlations characteristic of twist effects are evident in how, upon the pulse-like perturbation, the peak-to-peak distance changes (compared to the limit cycle period) as the perturbation returns to the periodic orbit. For a clock with $\epsilon = 0$, the peak-to-peak distance outside the limit cycle after a perturbation coincides with the 24 h peak-to-peak distance within the periodic orbit (Figure 7G). In the case of the Poincaré oscillator with positive twist, the peak-to-peak distance of the oscillator after the perturbation that increases the instantaneous amplitude is *longer* than the period of the steady-state rhythm (24 h), since $\dot{\phi} < \omega$. As the perturbed amplitude decreases and is attracted back to the stable cycle, the peak-to-peak distance approaches 24 hours (Figure 7H, consistent with the formulation 'positive' twist, which implies positive amplitude-period correlations). Conversely, the clock with negative twist exhibits a *shorter* peak-to-peak distance than the 24 h rhythm during the relaxation time ($\dot{\phi} > \omega$), which subsequently increases as the system returns to the stable periodic orbit (Figure 7I). When considering the findings from the last two paragraphs as a whole, it becomes clear how the extent of the phase shift between a perturbed and an unperturbed oscillator (red and black dashed lines in Figure 7D–F, respectively) depends on ϵ and thus the shape of the phase response curve (PRC) and magnitude of the phase shift depend on this twist parameter (Figure 7J–L).

We then evaluated how twist affects the response of an individual oscillator to a periodic zeitgeber input. We observed how the range of entrainment becomes larger with larger values of absolute value ϵ , as seen by the wider Arnold tongues in Figure 7M–O. It is widely known that, when the frequency of an applied periodic force is equal or close to the natural frequency of the system on which it acts, the amplitude of the oscillator on which the driving force (zeitgeber) acts on increases due to resonance effects. Interestingly, also the resonance curve was affected by twist: for the oscillator with no twist, the maximum amplitude occurred as expected for a zeitgeber period of 24 h, matching the intrinsic oscillator's frequency (Figure 7P). In oscillators with positive or negative twist, however, the maximum amplitude after zeitgeber forcing increases and decreases with zeitgeber period, respectively (Figure 7Q, R), resulting in skewed resonance curves. Interestingly, we observed that twist also affects the entrainment of an individual clock, and oscillators with high absolute values of twist cannot entrain to a periodic signal (Supplementary Figure S1 in 57), a phenomenon that has previously been described as shear-induced chaos 27,58,59 and that arises because the isochrones become so skewed, that the trajectory of the relaxation gets stretched and folded.

Relaxation rate affects the response of oscillators to perturbations

Not only the twist parameter ϵ , but also the amplitude relaxation rate λ affects the response of oscillators to perturbations and consequently the entrainment range. We have seen how, in the Poincaré model (equation 7), the twist parameter ϵ dictates how much trajectories are 'slowed down' or 'sped up' dependent on their distance from the limit cycle. The amplitude relaxation rate λ describes the rate of attraction back to the limit cycle, which is independent of ϵ . Together, these parameters both dictate how 'skewed' the isochrones are in phase space (see analytical expression in Materials and Methods, equation 11). For Poincaré oscillators with twist, the isochrones become more radial and straight as the amplitude relaxation rate increases (Figure 8A–C), with the implications that this has on PRCs and entrainment that have been mentioned before. A more 'plastic' clock (with a lower λ value) responds to an external pulse with larger phase shifts than a more rigid clock (with higher λ), resulting in phase response curves of larger amplitude (Figure 8D–F). This is also

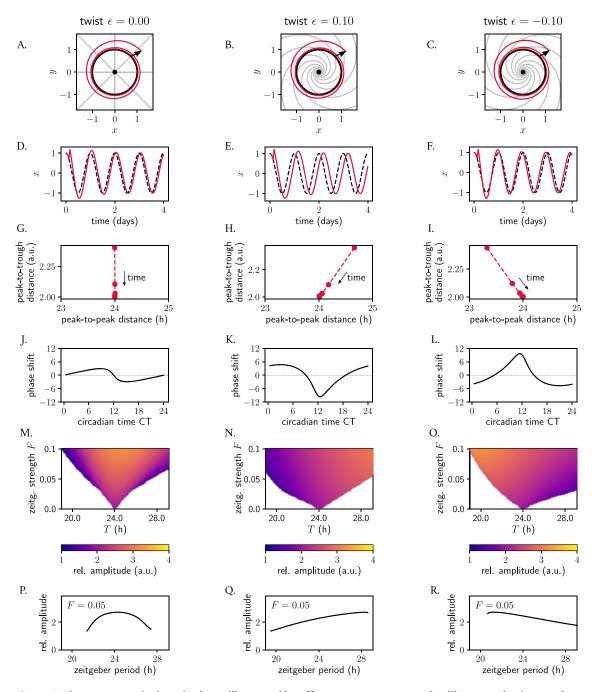


Figure 7. Phase space twist in a single oscillator and its effect on responses to pulse-like perturbations and to entrainment to periodic zeitgebers. (A–C) Poincaré oscillator models for different twist values ϵ , shown in phase space. The limit cycle is shown in black; perturbed trajectories are shown in red; isochrones are depicted in grey. (D–F) Time series corresponding to panels (A–C): the unperturbed limit cycle oscillations are shown with a dashed black line; perturbed trajectories are shown in red. (G–I) Peak-to-trough distance as a function of the peak-to-peak distance from the perturbed time series from (D–F) relaxing back to the limit cycle (the implicit time dimension is indicated in the panels). (J–L) Phase response curves: the shape of the PRC and the extent of the phase shifts (in hours) depends on the twist ϵ . (M–O) Arnold tongues illustrating entrainment ranges of an individual oscillator with different phase space twist values to a periodic sinusoidal zeitgeber input of different periods τ . The amplitude of the entrained clock is color-coded, with yellow colors corresponding to larger amplitudes. (P–R) The amplitude of an individual oscillator driven by a sinusoidal zeitgeber input of strength τ and varying periods τ . Amplitudes are defined as the average peak-to-trough distances of the entrained signal.

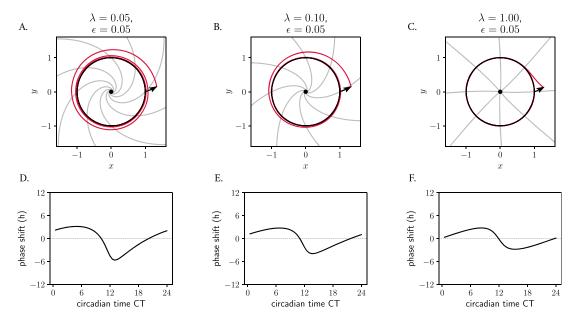


Figure 8. Role of amplitude relaxation rate on the skewing of isochrones and consequently on the response of oscillators to perturbations. (A–C) Poincaré oscillator models with different amplitude relaxation values λ , but otherwise identical (free running period $\tau = 24$ h, amplitude A = 1 and twist $\epsilon = 0.05$ h⁻¹), shown in phase space. The limit cycle is shown in black; perturbed trajectories are shown in red; isochrones are depicted in grey. (D–F) Phase response curves: the shape of the PRC and the extent of the phase shifts depends on the relaxation rate λ . See Materials and Methods for details on the analytical derivation of the equation of isochrones and roles of λ and ϵ .

intuitive, as it is clear that for larger values of λ , the twist has shorter effective time to act upon perturbed trajectories because the perturbation spends less time outside the limit cycle.

Single oscillator phase space twist affects coupled networks

Biological oscillators are rarely alone and uncoupled. Coupled oscillators, instead, are at the heart of a wide spectrum of living things: pacemaker cells in the heart, ³⁸ insulin- and glucagon-secreting cells in the pancreas ⁶⁰ or neural networks in the brain and spinal cord that control rhythmic behavior as breathing, running and chewing. ⁶¹ A number of studies have shown that networks of coupled oscillators behave in a fundamentally different way than 'plain' uncoupled oscillators. ^{33,62,63} This suggests that single oscillator twist might also affect how coupling synchronizes a network of oscillators.

To analyze the effect of twist on coupled oscillators, we study systems of Poincaré oscillators coupled through a mean-field (equation 8). We start by coupling two *identical* oscillators and analyzing the effect that different twist values have on the behavior of the coupled network. The numerical simulations show that increasing coupling strengths affect the period of the coupled system: oscillators with positive twist show longer periods as the coupling strength increases, whereas oscillators with negative twist show period-shortening (Figure 9A and Supplementary Figure S2A–C in⁵⁷). Coupling results in an increase in amplitude but, for our default relaxation rate value ($\lambda = 0.05 \text{ h}^{-1}$), no significant differences across oscillators with different twist values were found (Supplementary Figure S2 in⁵⁷). Increasing relaxation rates (oscillators that are attracted faster back to the limit cycle upon a perturbation) nevertheless resulted in less coupling-induced period or amplitude changes (Supplementary Figure S3 in⁵⁷).

We then analyzed what happens when oscillators with *different* phase space twist values are coupled through their mean-field. For this purpose, we took a system of 3 oscillators (with twist values $\epsilon=0$, $\epsilon<0$ and $\epsilon>0$) and turned on mean-field coupling after a certain time. We observed that all three oscillators synchronize to the mean-field for low values of the absolute value of twist (Figure 9B) although, due to the coupling-induced period differences, the relative phases of the individual oscillators to the mean-field depend on the specific twist values of the individual clocks: those with negative twist oscillate 'ahead' of the mean-field (with a phase advance) but clocks with positive twist cycle with a phase delay (Figure 9B). Very large absolute values of twist, interestingly, produce again complex chaotic dynamics (Supplementary Figure S4 in 57). The oscillator with a large value of ϵ does not synchronize to the mean-field and, as a result of the interoscillator coupling, this desynchronized oscillator 'pulls' to desynchronization the other two oscillators which otherwise would have remained in sync.

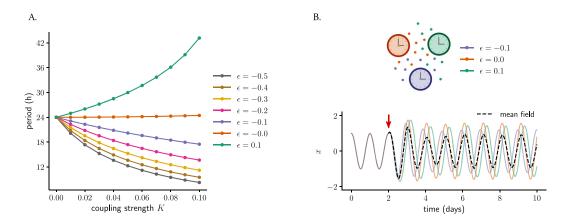


Figure 9. Phase space twist ϵ affects the response of a network to coupling. (A) Mean-field coupling of two identical Poincaré oscillators (sharing the same value of twist ϵ) and the effect on the period of the coupled system: whereas coupling among oscillators with positive twist results in longer periods of the coupled network, mean-field coupling among oscillators with negative twist results in the network running at a faster pace (period shortening). (B) Mean-field coupling can synchronize a network of oscillators with different twist values. Shown are three oscillators that only differ in their twist parameters (but else identical: amplitude, amplitude relaxation rate and period are the same for the three oscillators). All oscillations overlap during the first part of the time series, since twist has no effect within the limit cycle. At a certain point (indicated by the red arrow), mean-field coupling is turned on and it is observed how the oscillators respond differently to that mean-field coupling, ending up with different individual amplitudes as well as relative phases to the mean-field (depicted in dashed black line).

Discussion

This study aimed at characterizing period-amplitude correlations across circadian clock models of different complexity. Body clocks have to cope with cellular heterogeneity, what results in cellular clocks being variable across networks and tissues. We addressed whether the amplitude-period correlations that have been observed experimentally can be explained through heterogeneity in the cellular clocks, and what design principles are needed to produce such twist effects that we term *parametric twist*. Moreover, clocks live in constantly changing environments, that requires them to get adapted in the face of external changes. With the concept of *phase space twist* we study how this parameter, included explicitly in models, tunes the oscillator's response to coupling, entrainment and pulse-like perturbations. We also retrieve the 'old' terminology of hard *versus* soft oscillators to refer to oscillations with negative and positive amplitude-period correlations, respectively.

It is important to note that a limit cycle oscillates with a characteristic period and amplitude regardless of the initial conditions. In contrast, conservative oscillations, like the Duffing oscillator (Figure 2), have a period and amplitude defined by the initial conditions. For this reason we have not termed *twist*, in the context of conservative oscillations, *parametric* or *phase space*. We refer to *parametric twist* as that type of amplitude-period correlations that appears when studying an *ensemble* of limit cycle oscillators with differences in their intrinsic properties (i.e., biochemical parameters) and *phase space twist*, as that type of transient amplitude-period correlation that an *individual* oscillator experiences when encountering an external stimulus that results in an adaptation of its period and amplitude.

Parametric twist effects appear when analyzing populations of cells and require nonlinearities in oscillator models. The feedback loops needed to generate oscillations, which are commonly modeled with non-linear terms, ^{33,36,47,54} result in variations of parameters producing oscillations of different but correlated amplitudes and periods. We found that the type of parametric twist (positive or negative) depends on a number of factors: (i) on the biochemical parameter being affected, since different model parameters control the oscillation properties (amplitude and period) differently (Figures 3 and 5); (ii) on the region within the Hopf bubble where the clock's parameter set is at (Figure 4); (iii) on the type of model and variable being measured: simple models with single negative feedback loops show the same parametric twist effects for all variables because the parameter of interest has the same effect on the amplitude of all variables. In complex models with synergies of loops, a change in one parameter might increase the amplitude of one variable but decrease the amplitude of a second variable (Figure 5 and Supplementary Table S1 in⁵⁷). This highlights the challenge of defining circadian amplitude (and twist effects) to find a metric for the whole oscillating system. Our findings explain why some experimental studies have found positive twist, ²⁶ others negative, ¹⁴ whereas some other works found very little correlations. ⁶⁴

We also addressed the concept of *phase space twist*, a parameter that we (and others 14,38,39) introduced in a generic Poincaré oscillator. This twist parameter ϵ refers to period-amplitude correlations within a single oscillator, and it is a measure of the modulation of both instantaneous amplitude and period of the clock as the perturbed clock returns to the steady-state rhythm (i.e., limit cycle). Zeitgeber pulses, recurring zeitgeber inputs or coupling can all be regarded as 'perturbations', as any of these inputs modifies the natural clock's limit cycle in phase space. Given this, it is not surprising that phase space twist, which characterizes the adaptation to a perturbation, has significant implications for phase response curves (PRCs), entrainment and coupling.

We start by discussing the role of twist in PRCs. The twist parameter affects the extent of the phase shift after a zeitgeber pulse (Figure 7D–F): increasing the twist parameter in absolute value results in alterations in the phase response curves and consequently in the resetting properties of oscillators. In particular, increasing $|\epsilon|$ increases the amplitude of the PRC (Figure 7J–L) until the PRC is converted from a type 1 PRC to a type 0 PRC. ^{19,65} Consistently, amplitudes ^{17,18,20,21,66} and periods ¹⁸ of oscillators have also been shown to modulate their resetting properties. In these examples, clocks with short periods ¹⁸ and small amplitudes (in mathematical models ^{17,18,20} or in experiments with reduced coupling ^{20,21} or mutations ⁶⁶ that disrupt the normal rhythmicity) are easier to reset.

One can easily extrapolate these findings to responses to jet lag. If twist modulates the extent of a phase shift in response to a perturbation, it will also be critical in the adaptation to jet lag. This has indeed been found by Ananthasubramaniam *et al.*²³ Interestingly, the authors found that the instantaneous amplitude effects induced by jet lag are also modulated by twist. In particular, positive twist aided recovery to jet lag in simple Goodwin-like models: reduced amplitudes were accompanied by faster clocks (i.e., shorter periods) upon a phase advance (e.g. when traveling eastwards), but larger amplitudes coincided with longer periods when the phase had to be delayed (e.g. when traveling westwards). Consistent with these theoretical observations are experiments performed in mice, where compromising coupling in the SCN (and thus decreasing amplitudes) reduce jet lag drastically, since resetting signals are much more efficient.²¹

Transient amplitude-period correlations also affect the entrainment properties of an oscillator. Twist increases the range of zeitgeber periods to which the Poincaré oscillator can entrain (resulting in larger entrainment ranges, i.e., wider Arnold tongues, Figure 7M–O) and skews the resonance curves (Figure 7P–R). Some studies have even found coexisting limit cycles for driven oscillators with twist. ^{31,67} Of note, however, is that the period of the Poincaré oscillator in all our phase space twist simulations was set to 24 h. Other theoretical studies have shown that clocks with different intrinsic periods show different amplitude responses to entrainment. ^{23,68} thus combining phase space and parametric twist effects.

Single cells harbor self-sustained clocks, but they are coupled (see⁶⁹ for a review on coupling) to produce a coherent rhythm at the level of tissues and organisms. Coupled networks show different rhythmic properties than clocks in isolation: phases synchronize and ensemble amplitudes increase for over-critical values of coupling strengths.⁶² Our simulations provide an additional level of regulation, by showing that twist influences period length of individual clocks due to coupling. Positive twist results in periods lengthening upon mean-field coupling (Figure 9). The presence of coupling-induced changes in the oscillation period or amplitude can provide insights into the underlying oscillator type and the presence of twist. For instance, the longer period that has been observed in dispersed (and presumably uncoupled) U-2 OS cells compared to high-density cultures, ⁷⁰ but not in dispersed SCN neurons in culture, ^{71,72} could be explained by different implicit oscillator twist and amplitude relaxation rate values in different tissues.

In addition, the twist-induced modulation of period also affects the phase relation with which the clock oscillates respect to the mean-field: clocks with positive twist that tend to period-lengthen in response to coupling show later phases than those with negative twist, which tend to run quicker, consistent with chronotypes. It should be noted, however, that different phase relationships to the mean-field can also be obtained without twist, in networks of coupled oscillators with different intrinsic periods. But here again, the slower running clocks will tend to be phase-delayed in comparison to the faster-running clocks which will be phase-advanced. This suggests that twist is critical in how synchronization arises in network of coupled oscillators.

Our simulations assume that the coupling strength is constant across all oscillators. This assumption however might be questioned, since oscillators might 'talk' with different strengths to each other, especially because individual clocks are spatially organized within tissues, and have been shown to produce waves of oscillations for example in the SCN. Thus, a more plausible scenario may involve local coupling, where clocks couple to neighbor clocks with a strength proportional to the oscillator distance. In fact, spatial gradient in nearest neighbor coupling can lead to a robust phase patterning. Thus twist might not only be critical in how temporal synchronization but also spatial patterning arises in coupled ensembles.

The focus of this paper has been predominantly on twist, but it is important to remark that also the relaxation rate λ (i.e., how rigid/plastic an oscillator is) affects the response of oscillators to perturbations, consistent with previous computational work. 20,55 It is in fact the ratio of ϵ to λ what determines the skewing of isochrones (see the analytical derivation in Materials and Methods) and the oscillator's response to zeitgebers. Larger values of λ (more 'rigid' oscillators) imply that any perturbation is attracted back to the limit cycle at a higher rate and thus, perturbations 'spend less time' outside the limit cycle. Consequently, the twist parameter ϵ has a shorter effective time to act on the perturbed trajectory, and the isochrones become more straight and radial. Relaxation rate also has implications on coupling and entrainment. Rigid oscillators with $\epsilon \neq 0$ display less coupling-induced amplitude expansions (Supplementary Figure S3B in, ⁵⁷ also ²⁰) and less period variations (Supplementary Figure S3A in ⁵⁷) in response to coupling than more 'plastic' oscillators with lower λ values. Moreover, rigid oscillators have smaller ranges of entrainment and narrower Arnold tongues. ²⁰ Again, our results imply that any observation of resonant behavior and/or period changes might provide information on whether the underlying oscillator is a rigid *versus* plastic and hard *versus* soft clock.

We finish with an open question. Throughout our work we have claimed that period-amplitude correlations are widespread in both *in vivo* and in *in silico* clocks, and how they are critical to define how oscillators function in their environment. But, how does one integrate the circadian clock's property of temperature compensation within this framework? The circadian clockwork is temperature-compensated, ^{18,49,76–78} which means that increasing temperatures do not speed up significantly the clock, which still runs at approximately 24 h. A proposed hypothesis for temperature compensation suggests that temperature-sensitivity of oscillation amplitude could stabilize the period. ^{18,78} In support of this supposition is evidence from *Neurospora* and *Gonyaulax* suggesting that amplitude increases in response to temperature, but decreases in *Drosophila*. What are the twist effects behind this adaptation? Twist might play a role in how fast that temperature variation is sensed in the clockwork and modulate the adaptation to find the new steady-state rhythm of larger or smaller amplitude at the new temperature.

Conclusions

Despite the complexities in quantifying amplitude, our models stress the important role of circadian amplitudes and their correlation with oscillator period. Although amplitudes are known to be regulated by a number of internal and external cellular factors, including cellular biochemical rates, light conditions, coupling, equation ageing, our findings stress that twist effects (i.e., co-modulations of amplitudes and periods) also feed back and affect the interaction of oscillators with the environment, facilitating entrainment, fastening response to pulse-like perturbations or modifying the response of a system to coupling. The theory of our conceptual models can also be applied to other oscillating system such as cardiac rhythms, somite formation, central pattern generators or voice production.

Data and code availability

Extended data

GitHub: Simulated (numerical) data generated for the manuscript. https://github.com/olmom/twist

GitHub: Supplementary Material. https://github.com/olmom/twist

Analysis code

GitHub: Source code to generate and analyze the data and to reproduce all figures. https://github.com/olmom/twist

Data and codes are available under the terms of the Creative Commons Attribution 4.0 International license (CC-BY 4.0).

Acknowledgements

The authors thank Gianmarco Ducci, Bharath Ananthasubramaniam and Adrián Granada for stimulating and helpful comments as well as constructive criticism. The authors would also like to acknowledge the work done by the Bachelor and Master students Pia Rose, Julia K. Schlichting, Barbara Zita Peters Couto and Rebekka Trenkle.

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Reviewer Report 29 September 2023

https://doi.org/10.5256/f1000research.148657.r202730

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- 1. The authors have conducted a study on the relationship between amplitude and period in various circadian clock models. However, it is crucial to address the novelty and distinctions of this study compared to prior work, specifically the study by Bokka et al. in 2018 (https://doi.org/10.1049/iet-syb.2018.0015). Additionally, while the models in this manuscript are based on Hill-type repression (such as the Goodwin and Gonze models), it is important to acknowledge recent research findings that reveal different repression mechanisms in the Drosophila (https://doi.org/10.1073/pnas.2113403119) and mammalian circadian clocks (https://doi.org/10.1098/rsfs.2021.0084). These studies demonstrate that the use of multiple repression mechanisms, including protein sequestration, can substantially alter the oscillator dynamics (https://doi.org/10.1049/iet-syb.2015.0090). Given these discoveries, it would be valuable to explore or discuss how different repression mechanisms impact the relationship between period and amplitude, as demonstrated by Bokka et al. in 2018.
- 2. In the Conclusions section on page 19, please provide a reference for 'cellular biochemical rates' as a factor influencing circadian amplitudes. For instance, it was demonstrated that faster production and degradation rates of proteins can generate circadian rhythms with higher amplitudes in the Drosophila circadian clock (https://doi.org/10.1073/pnas.2113403119).

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Is the work clearly and accurately presented and does it cite the current literature? Partly

Is the study design appropriate and is the work technically sound? Yes

Are sufficient details of methods and analysis provided to allow replication by others? Yes

If applicable, is the statistical analysis and its interpretation appropriate?

Are all the source data underlying the results available to ensure full reproducibility? Yes

Are the conclusions drawn adequately supported by the results? Yes

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: Circadian rhythms, mathematical modeling

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

Reviewer Report 18 September 2023

https://doi.org/10.5256/f1000research.148657.r202731

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In this article, the authors describe the phenomenon of twist (period-amplitude inter-dependence) using a range of differential equation oscillator models, ultimately with a view to understanding circadian clocks. The simplest models (harmonic and Duffing) are used to demonstrate the non-linearity introduces a relationship between amplitude and period. The authors then switch to limit-cycle oscillators, because limit-cycle oscillators are more appropriate for studying circadian clocks.

They use two approaches to studying twist. The first, referred to as "parametric twist", computes the periods and amplitudes of oscillators with varied parameter values, creating graphs that relates period to amplitude (and thus demonstrate twist - if the correlation is positive, there is positive twist, and negative correlation indicates negative twist). This parametric twist analysis is performed using two limit cycle models, one with one feedback loop and one with more (synergistic) loops. The second, referred to as "phase space twist", examines the response of a single oscillator to a temporary change in one parameter, mimicking signaling. A separate limitcycle model (a Poincare oscillator) is used for this analysis. Important to the discussion of phase space twist is the concept of isochrons, which are useful for understanding the effects of perturbations on the phase and speed of the clock (does it speed up or slow down the clock?) and it is easy to see how amplitude and phase response are related, when we look at how curved the isochrons are. The authors relate these analyses to phase response curves, recovery from jetlag, the range of entrainment, and temperature compensation and conclude, that although measuring amplitude and understanding relationship between amplitude and period are highly complicated, it is nonetheless important to be mindful of the role of amplitude and its correlation to period when studying clocks.

On the one hand, I really appreciated that the authors chose models that best illustrated the points and that the paper is organized to go from simple ideas to more complex ideas. It was helpful to have the initial explanation of twist use duffing oscillator (which has non-linearity) as a contrast to a harmonic oscillator (which doesn't). It provided a nice introduction to the cause of twist and helped me to develop an intuition. Then, it made sense to use a very simple (3-ODE Goodwin oscillator) as the first example of a limit cycle oscillator. Since it has so few parameters, it was possible to perturb them one at a time and in pairs to illustrate a good portion of the possibilities for how one might observe heterogeneity in the system. Then, moving to Almeida's oscillator demonstrated how quickly the analysis can become very complex. All of this provides a nice progression of complexity. Then, we pivot a bit to the Poincare oscillator, to study twist in a more dynamical way - instead of permanently choosing different parameters, the authors model light signaling (at least, initially) and study the effects of the model returning to its limit cycle after the perturbation is over. This model is particularly appropriate for studying the dynamics of recovery because it is a 2-ode model (so we can examine it in "phase space", which is helpful for visualizing the clock) and because there is a parameter that controls how related period and amplitude are. It is also straight-forward to adjust how curved the isochrons are, which makes illustrating twist very clear. Then, after studying the response of a single oscillator to light, they use the same model to study the effects of intercellular signaling and the effects of different twist on the properties of the ensemble oscillation (e.g. how long the ensemble period is in contrast to that of individual oscillators).

On the other hand, I would have appreciated a more explicit connection between the discussion of "parametric twist" and "phase space twist". They are both due to changes in parameter values causing changes in period and amplitude. In the one case, we measure the period and amplitude of oscillators with different (constant) parameter values. In the second case, we measure instantaneous changes in period and amplitude due to a temporary changes in parameter values (I am referring to Z(t) as a parameter here). In both cases, the shape of the isochrons in the unperturbed condition should help us to predict what will happen in the perturbed condition. There are two challenges with making that point clear with the Goodwin and Almeida models - 1) the isochrons aren't lines, they are multi-dimensional manifolds, and 2) the isochrons in the unperturbed and perturbed systems are different. So I am asking for too much to relate the two

discussions by using images of isochrons, but a more explicit comparison would help the reader understand how the analyses are related to each other.

In the analysis of the modified Poincare oscillator, the term "twist" is used differently than it is in the "parametric twist" analysis. In the parametric analysis, the twist captures the full effects of the perturbations (we plot the measured periods and amplitudes across the heterogeneous set of oscillators). In the phase space analysis, "twist" refers to a particular parameter \epsilon that, when non-zero, introduces a dependence of the speed and amplitude on each other. However, that twist parameter (\epsilon) is not the only parameter that affects the skewness of the isochrons. The relaxation rate \lambda also affects the skewness of the isochrons. And the skewness of the isochrons affects the changes in speed and amplitude when the oscillator is signaled (i.e. a parameter is temporarily perturbed). So, if we were to measure the effects on period and amplitude (as is done with figures 7G, H, and I), that reflects that values of both \lambda and \epsilon. When I was thinking about how to relate the "parametric twist" and "phase twist" analysis, I was looking for aspects of figures 3 (Gonze model), 5 (Almeida model), and 7 (poincare model) to relate and it seemed to me that that the subfigures that related the amplitude of the oscillator to its period (3C, 3D, 3E, 3F, 3G, 3H, 5B, 5E, 7G, 7H, 7I) should be the figures that I can connect. They are labeled as twist in figures 3 and 5. In Figure 7, they show an approximation of instantaneous amplitude and period (rather than steady-state amplitude and period), so it isn't a perfect 1-1 mapping, but it is still the closest that makes sense to me. And in all of these cases, the output could be labeled twist if \epsilon didn't get to claim that label. I think it would make the concept of twist clearer to the reader if it were consistent across both analyses that twist is the "effect" we can measure. My reading of the text as it is currently written is that twist is a phenomenon we can measure in "parametric twist" analyses and that twist is a parameter that causes one change in the isochrons (but not all of the changes that can happen) that ultimately leads to something we can measure. Also, labeling the parameter as twist in the Poincare oscillator makes it seem like we need to use the Poincare oscillator to perform a "phase space twist" analysis. But we aren't limited to the Poincare oscillator for such analyses.

Specific Points

- It would be helpful to have a little more framing text that indicates the different models are used to make different points.
- I think of the "phase space twist" analysis as analysis of recovery from temporary changes to parameters and "parametric twist" as analysis of permanent differences in parameters. If that is the way you intend readers to think about it, then I would suggest changing the names to reflect those ideas more directly. Right now, the term "phase space twist" stands out as a highly specific analysis the model most be 2D so you can draw the isochrons. But I think that is limiting. I totally embrace idea that there is both heterogeneity among oscillators and temporary changes due to intercellular signaling, so I would love to see it characterized that way more explicitly. What about "twist due to heterogeneity" and "twist due to signaling"? They aren't quite zippy enough and rely on my characterization of twist as a measurable phenomenon, so I don't push those terms exactly.
- On page 8, when you are introducing isochrons, it would be helpful to have a figure to refer to. The most natural figure is Figure 7, but that would force you to make an unnatural reordering of the figures. So maybe introduce a new figure?

- On Page 11, (analyses shown in Figure 3), how did you choose which parameters to pair together?
- On Page 11, there is a discussion of the Hopf bifurcation that is interesting (and highlights how sensitivity twist is to our choice of parameters). It occurs to me that this has implications for modeling SCN oscillators as they recover from TTX treatment. The individual oscillator amplitude crashes and the oscillators lose synchrony during TTX and both recover after TTX is washed out. Is there anything you could add to the discussion about this?
- On Page 12, in the paragraph about how overall twist effects depend not only on the parameter being studies, but also on the variable that is measured, I was reminded of early modeling studies that showed not-immediately intuitive results. In his 1995 fly model paper (A model of circadian oscillations in the Drosophila period protein, Proc. R. Soc. Lond. B, 1995), Goldbeter showed that the rate of (doubly-phosphorylated) PER degradation affected both period and amplitude and that increasing this rate led to PER amplitude increasing (rather than decreasing, as one might expect). This can be reasoned through, but it is another illustration of the complexity of the non-linear feedback and the relationship between period and amplitude. Is this worth bringing up here or on the discussion?
- Figure 6. Why are there no yellow dots on the limit cycle for A and B? Also, maybe make the
 dot color on the limit cycle different from yellow (e.g. to blue), so that you can describe what
 is happening over time with even more clarity (i.e. that the simulation eventually reaches
 the "blue" dot after enough cycles).
- Figure 8 and text referring to it. I appreciate that you relate twist to entrainment and PRCs. I think this is interesting and shows why twist is important to understand.
- Page 16 first paragraph. "This suggests that single oscillator twist might also affect how coupling synchronizes a network of oscillators" is too weak. Given that coupling (M) and the perturbation used in Figure 7 (Z(t)) both affect the model the same way (they both are added to the rate equation for the x variable), Figure 7 shows that the PRCs are affected by twist. Not only are they affected, they are affected dramatically Figures 7K and 7L look like they are opposite each other (i.e. flipped sign). This means we are guaranteed that the oscillators will respond differently to intercellular signaling and that that the ensemble oscillation will be different.
- Page 16, in the discussion of the ensemble period, we see that for this model that positive twist leads to longer periods (and negative twist to shorter). What is missing from this discussion is that the relative phase of the signal and of the PRC of the receiver determine the period of the ensemble. Positive twist leads to longer periods because the mean field is determined by the value of the x variable. If it were determined by a variable with the opposite phase, then we would have the opposite effect. This affects the related paragraph in the Discussion as well.
- Page 17. The meaning of "For this reason we have not termed twist, in the context of conservative oscillations, parametric or phase space" wasn't clear to me. I think you mean you can't study twist in non-limit-cycle oscillators in the same way you can for limit cycle oscillators. Is that correct?

Is the work clearly and accurately presented and does it cite the current literature? Yes

Is the study design appropriate and is the work technically sound?

Yes

Are sufficient details of methods and analysis provided to allow replication by others? Yes

If applicable, is the statistical analysis and its interpretation appropriate?

Are all the source data underlying the results available to ensure full reproducibility? Yes

Are the conclusions drawn adequately supported by the results? Yes

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: I study the response of limit cycle oscillators to signals and sensitivity analysis involving isochrons.

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

Reviewer Report 18 September 2023

https://doi.org/10.5256/f1000research.148657.r202732

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In this manuscript, the authors make a comprehensive exploration of the "twist" (amplitude-period correlation) in nonlinear oscillators, especially in the context of circadian rhythms. The Introduction insightfully summarizes the background and offers a clear explanation of the topic. While some sections exhibit a degree of coarseness in their explanations, the central theme remains consistently and clearly highlighted. Overall, this manuscript is a noteworthy addition to the field, and I recommend its acceptance for indexing with minor revisions.

Minor comments:

- 1. Abstract: I suggest changing "..., fastening recovery..." to either "speeding up recovery" or "accelerating recovery".
- 2. p.4, subsection "Goodwin-like models": I recommend changing "sigmoidal Hill curve" to "Hill curve", as "sigmoidal" often refers to forms involving exponential functions.
- 3. p.4, subsection "Goodwin-like models": Add biological interpretations of the variables x, y, and z (mRNA, ...).
- 4. p.5. II.4-5: While the Gonze model[33] describes degradation processes with Michaelis-Menten dynamics, it may not provide sufficient historical context. I recommend citing Kurosawa & Iwasa (2002)¹ for a more comprehensive historical perspective.
- 5. p.6, Table 2: The reference [36] should be spelled out in "The default values are taken from [36].
- 6. p.11. Figure 4: It is unclear if the graphs are figurative cartoons or based on actual numerical data from Gonze oscillators.
- 7. pp.11-12: The justification for the use of the Almeida model can be described. For example, why not use the detailed Gonze model or Becker-Weimann model?
- 8. p.13, ll.1-2: It is challenging to assert that the "parametric twist" emerges solely due to a heterogeneous population. While it describes the average oscillator behavior in a population (often assumed to be normally distributed), it essentially characterizes the behavior of a single representative oscillator.
- 9. p.14, "... 24 h, matching the intrinsic oscillator's frequency (Figure 7P).": I suggest to rephrase it to "... 24 h, matching the oscillator's intrinsic period (Figure 7P)." The change from "frequency" to "intrinsic period" would provide more specificity and clarity.
- 10. p.18 "the longer period that has been observed in dispersed..., but not in dispersed SCN neurons in culture": In light of the findings in Figure 5, it's noteworthy that when Bmal1 oscillation is observed (as opposed to Per2), the SCN period lengthens under pharmacological decoupling or physical sub-sectioning of the SCN (Myung *et al.*, 2012²),.
- 11. p.18 "Thus, a more plausible scenario may involve local coupling...": Especially in the SCN, there are papers that highlight the role of specific coupling among subregions, which can reproduce the wave-like spatial patterns and seasonal encoding of circadian time. These can be discussed.

References

- 1. Kurosawa G, Iwasa Y: Saturation of enzyme kinetics in circadian clock models. *J Biol Rhythms*. 2002; **17** (6): 568-77 PubMed Abstract | Publisher Full Text
- 2. Myung J, Hong S, Hatanaka F, Nakajima Y, et al.: Period coding of Bmal1 oscillators in the suprachiasmatic nucleus. *J Neurosci.* 2012; **32** (26): 8900-18 PubMed Abstract | Publisher Full Text

Is the work clearly and accurately presented and does it cite the current literature?

Partly

Is the study design appropriate and is the work technically sound?

Yes

Are sufficient details of methods and analysis provided to allow replication by others? Yes

If applicable, is the statistical analysis and its interpretation appropriate? Yes

Are all the source data underlying the results available to ensure full reproducibility? Yes

Are the conclusions drawn adequately supported by the results? γ_{PS}

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: Circadian rhythms, computational modeling of circadian clocks, suprachiasmatic nucleus (SCN), bioluminescence reporter imaging

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

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