

Review



Cite this article: Jiménez A, Lu Y, Jambhekar A, Lahav G. 2022 Principles, mechanisms and functions of entrainment in biological oscillators. *Interface Focus* **12**: 20210088. <https://doi.org/10.1098/rsfs.2021.0088>

Received: 17 December 2021

Accepted: 7 March 2022

One contribution of 5 to a theme issue 'Time-keeping and decision-making in living cells; Oscillations and Synchronization (Part I)'.

Subject Areas:

systems biology

Keywords:

entrainment, biological oscillators, synchrony, phase response curve, Arnold tongue

Author for correspondence:

Galit Lahav

e-mail: galit@hms.harvard.edu

Principles, mechanisms and functions of entrainment in biological oscillators

Alba Jiménez¹, Ying Lu¹, Ashwini Jambhekar^{1,2} and Galit Lahav^{1,2}

¹Department of Systems Biology, Blavatnik Institute at Harvard Medical School, Boston, MA 02115, USA

²Ludwig Center at Harvard, Boston, MA 02115, USA

AJ, 0000-0002-9014-5857; YL, 0000-0003-3516-7735; AJa, 0000-0003-1078-6601; GL, 0000-0003-4758-6427

Entrainment is a phenomenon in which two oscillators interact with each other, typically through physical or chemical means, to synchronize their oscillations. This phenomenon occurs in biology to coordinate processes from the molecular to organismal scale. Biological oscillators can be entrained within a single cell, between cells or to an external input. Using six illustrative examples of entrainable biological oscillators, we discuss the distinctions between entrainment and synchrony and explore features that contribute to a system's propensity to entrain. Entrainment can either enhance or reduce the heterogeneity of oscillations within a cell population, and we provide examples and mechanisms of each case. Finally, we discuss the known functions of entrainment and discuss potential functions from an evolutionary perspective.

1. Introduction

Oscillating systems can interact with each other in various ways. They can enhance or negate each other's effects (constructive and destructive interference, respectively) or synergize with each other to achieve amplitudes greater than the sum of the two systems (resonance). When two oscillating systems interact, one or both can experience an alteration in frequency to become phase-locked, meaning that the phase difference between the two oscillating systems remains constant in time and is robust to perturbations [1]. This situation is called entrainment.

Entrainment was originally described as two pendulum clocks coupled through a wooden structure [2] (figure 1*a*). Synchronization in this system was achieved via mechanical vibrations through the wooden coupling bar. Oscillations are also found in various biological systems and can operate at the molecular level (e.g. cardiac cell beating) or at the organismal level (e.g. sleep–wake cycles). Entrainment of these oscillations can occur through interactions between single cells, within a single cell or between a cell and its environment (figure 1*b,d*).

Biological oscillators can entrain in a variety of ways. Two biological oscillators in neighbouring cells can interact and influence each other through their extracellular environment (figure 1*b*). Entrainment between cells often occurs through secreted factors and therefore becomes apparent as cell density increases [3–6]. It allows coordination between cells in a tissue in order to perform a function: for example, cardiac cells synchronize their oscillations in order to provide a strong single voltage that leads to heart contraction [7,8]. Two biological clocks can also entrain *within* a single cell, as observed between the circadian and cell cycle oscillators (figure 1*c*). Entrainment of oscillators within a single cell allows for synchronizing the processes controlled by the two individual oscillators. Last, the frequency of a biological clock can entrain to an environmental periodic rhythm (figure 1*d*) that is itself unaffected by the biological oscillator. The most prevailing example of such unidirectional entrainment is the circadian clock, in which sleep–wake cycles entrain to

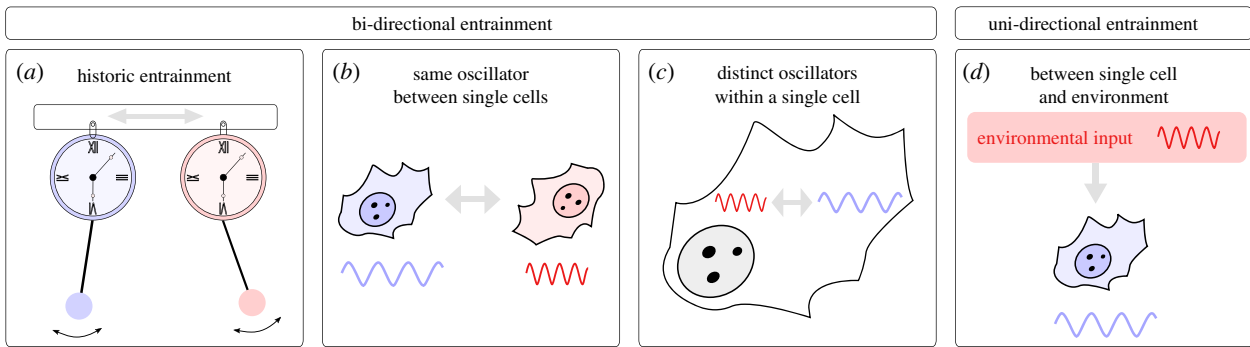
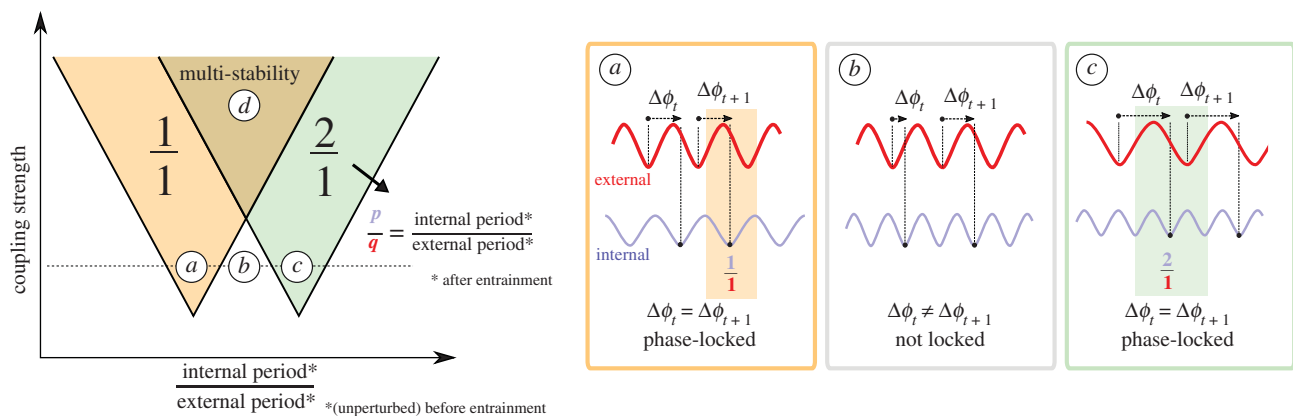


Figure 1. Entrainment types and their directionality. (a) Entrainment as originally described between two physically connected oscillating pendulums. (b) Entrainment of the same oscillator in two neighbouring single cells. (c) Entrainment of distinct oscillators within a single cell. (d) Entrainment of an oscillator within a single cell by an external periodic input.

Box 1. The coupling and uncoupling between the oscillator and external input can be summarized in an ‘Arnold tongue’ plot. The Arnold tongue plot can be interpreted in the following way: with a fixed coupling strength (y -axis), if the intrinsic frequency of one oscillator traverses horizontally across the Arnold tongue plot, the coupled system will either stay not locked (case b) or be locked into distinct frequency modes featuring fixed p/q ratios (cases a and c). Phase locking is defined by measuring $\phi(t)$ and $\Delta\phi(t)$, with $\phi(t)$ being the phase of an oscillator relative to the start of the cycle, expressed as a fraction of the period $\phi(t) \in [0, 2\pi]$, and $\Delta\phi(t)$ being the difference in phase between two periodic signals at a given time $\Delta\phi(t) = \phi_{\text{oscillator1}}(t) - \phi_{\text{oscillator2}}(t)$. When the phase difference $\Delta\phi(t)$ between two signals is constant in time, the two signals are considered to be phase-locked. Traversing vertically over the plot (increasing coupling strength) illustrates how the coupled system becomes more robust against fluctuations (broadening of Arnold tongues) or can lead to multi-stability (case d) or other irregular dynamics such as chaos. ‘Tongues’ associated with high-order entrainment modes ($5/4$, $3/2$, etc.) are usually smaller than that for the equal-frequency model ($1/1$) and therefore harder to observe experimentally (figure 2, *Other entrainment ratios*). The amplitude during entrainment remains unaltered as shown both theoretically [11] and experimentally [12].



light–dark cycles [9,10]. Entrainment between a cell and its environment allows organisms to keep their physiology in synchrony with their surrounding rhythms.

2. Key principles of entrainment

Entrainment depends on two basic conditions: (i) the coupling strength between the oscillator and external input and (ii) the similarity between the intrinsic frequencies of the internal oscillator and the external input in the absence of interaction [1]. Generally, a stronger coupling strength and closer intrinsic frequencies favour entrainment, though the exact requirement varies in different systems. The entrained state (or locked state) is represented by a rational number p/q ; after p periods of the internal oscillator and q periods of the external oscillator, the system returns to the same state. As the coupling strength increases, phase locking becomes possible at a wider range of external periods

(depicted by a broadening of Arnold tongues, see box 1), and the entrained mode is more robust against random fluctuations. Further increasing coupling strength may result in complex phenomena such as multi-stability, in which multiple entrainment modes coexist, and chaos. These are depicted by the overlap of different ‘tongues’. The transition between a robust locked state and a chaotic one has been observed in a classic example of periodically stimulated cardiac cells, in which a small variation of the period of the electrical stimuli caused a transition between normal and pathological behaviour of cardiac tissue (dysrhythmia) [13].

3. Examples of biological oscillators exhibiting entrainment

We will explore entrainment focusing on six biological examples of autonomous oscillators, which have been shown to entrain experimentally (figure 2). For each example,

	(a)	(b)	(c)	(d)	(e)	(f)
	circadian clock	cell cycle	mitotic exit (Cdc14)	cardiac pacemaker	glycolysis cycle	inflammatory response (NF- κ B)
network internal/external						
entrainment stimuli	light Plautz <i>et al.</i> , 1997 serum (cAMP, protein kinase C, Ca^{2+}) Balsalobre <i>et al.</i> , 1998 drugs (forskolin or dexamethason) Bieler, 2014; Feillet 2014	wee1 Matsuo <i>et al.</i> , 2003	Cyclin B (Cib2) Lu and Cross 2020	electric current Jalife, 1984	glucose and cyanide solution Bier <i>et al.</i> , 2000	TNF-alpha Kellogg and Tay, 2015
natural period	24 h	24 h	90 min	200 ms	50 s	90 min
range of 1/1 entrainment	8 h to 16 h Laranjeiro <i>et al.</i> , 2003	20 h to 24 h Goldbeter, 2012	40 to 100 min Lu & Cross 2020	180 to 240 ms Anumonwo <i>et al.</i> , 1991	40 s Gustavsson <i>et al.</i> , 2015	60 to 120 min Kellogg and Tay, 2015
other entrainment ratios	3/2 Bieler, 2014	1/2 (theoretical proof only) Gerard and Goldbeter, 2012a	\emptyset	2/1, 1/2, 3/2, 5/4 Anumonwo <i>et al.</i> , 1991	\emptyset	1/1, 2/1, 1/2, 3/1 Kellogg and Tay, 2015 bi-stability at inputs around 150 mins

Figure 2. Six biological cases of entrainment. For each oscillator, the internal minimal network (blue) and external nodes (red) are portrayed, along with the stimuli used for entrainment and the observed entrainment ratios. (a) The fly circadian clock is regulated at the levels of transcription, protein stability and post-translational modifications [14]. It responds to light and GFs, but it can oscillate freely in the dark [9]. (b) The mammalian cell cycle network contains four coupled modules each centred around one cycle/Cdk complex which promotes progression or transition into the ordered succession of the cell cycle phases G1, S, G2 and M. The cell cycle components Wee1, p21 and cyclin E are transcriptionally regulated by the circadian clock [15]. (c) The Cdc14 network module is a negative feedback loop controlling cycles of nucleolar sequestration and release of Cdc14, which is essential for mitotic exit in budding yeast [16]. Each component of this loop (Cdc14, Cdc5, Cdh1) is coupled to the cell cycle. (d) Cardiomyocytes of the sinoatrial node (SAN) autonomously oscillate through action potentials that result from the opening and closing of sodium, calcium and potassium channels in their membrane, creating depolarization and repolarization oscillations [17,18]. (e) Glycolysis consists of the step-by-step breakdown of glucose and storage of the released Gibbs energy in the form of ATP. Oscillations correspond to changes in the concentration of glycolytic metabolites nicotinamide adenine dinucleotide plus hydrogen (NADH) and ATP. The molecular mechanism for oscillations is based on the speed of enzymatic reactions [12]. Sustained glycolytic oscillations require both glucose and cyanide to be present in the medium [19,20]. (f) The transcription factor NF- κ B oscillates between the cytoplasm and nucleus in response to the inflammatory signal TNF-alpha [21,22]. TNF-alpha signalling induces the dissociation of the I κ B::NF- κ B complex in the cytoplasm, allowing NF- κ B to enter the nucleus and activate transcription of its inhibitor I κ B, which sequesters NF- κ B in the cytoplasm [23].

we present the simplest model that accounts for oscillatory behaviour, along with the node(s) receiving the stimuli for entrainment. Each example follows the basic principle of biochemical oscillators but differs in terms of its network architecture, the nature of the oscillations, their time-scale and the number of entrainment modes. All examples have a negative feedback loop within their core network (see blue networks in figure 2) with additional positive feedback loops providing robustness [24]. Other details of the networks vary with regard to the number of nodes, number of positive and negative interactions and number of points of coupling to external oscillators. In addition, the oscillating factors differ between the various systems. For example, in the circadian clock example, mRNA and protein levels oscillate [25], nuclear factor kappa B (NF- κ B) and Cdc14 oscillate in their nuclear-cytoplasmic localization [23,26] and the glycolysis network oscillates in the products of enzymatic reactions [27]. The time-scale of oscillations also varies between these systems, with transcriptionally regulated systems exhibiting longer time-scales (hours for NF- κ B and the circadian clock) and oscillators relying on enzymatic reactions operating on shorter time-scales (less than a minute for glycolytic oscillations) (figure 2, *Natural period*). In this review, we will not focus on systems that show irregular oscillations, such as bursting dynamics of calcium ions [28],

nuclear translocation of Msn2 [29], insulin secretion by B-cells [30] or neuron spiking [31].

The six biological examples covered here have been extensively modelled using ordinary differential equations (ODEs) to describe their regulatory networks [21,32–37]. Dynamical systems tools, such as ODEs, phase portraits and bifurcation diagrams, are keys to understand how various systems differ in their requirements for initiating and sustaining oscillations [38]. For example, a model of the cell cycle [39] shows self-sustained oscillations only in the presence of growth factor (GF), thus defining GFs as a trigger between quiescence (non-oscillatory state) and proliferation (oscillatory state). Glycolytic oscillations require both glucose and cyanide to be sustained [19,20], with glucose alone leading to dampened oscillations but the addition of cyanide leading to sustained oscillations. GF and cyanide are thus Hopf parameters that are responsible for a Hopf bifurcation [20,36,39], meaning that they lead the system to transition from steady state (non-oscillatory) to a limit cycle (self-sustained oscillations). In most cases, oscillation triggers (Hopf parameters) also serve as entrainment stimuli. For example, entrainment of glycolytic oscillations by cyanide [20], or entrainment of NF- κ B by tumour necrosis factor (TNF) [40], but that is not always the case, for example GFs only initiate but cannot entrain the cell cycle [39].

4. Distinguishing between entrainment and other mechanisms leading to synchrony

Synchrony is the empirical observation of two systems oscillating in phase, which can result from either entrainment or other mechanisms such as gating [41,42]. During entrainment, all phases of the follower oscillator must be affected by the leading oscillator—in other words, the oscillatory curve of the follower must be proportionately stretched out or compressed through all phases. By contrast, during gating, the leading oscillator defines windows of time in which different phases of the follower oscillator can occur. As opposed to entrainment, a gating mechanism follows these three principles: (i) arresting the lead oscillator at any constant level will arrest the follower oscillator; (ii) only 1 : 1 ratios will be observed; and (iii) the leading oscillator impacts only specific phases of the follower oscillator.

Distinguishing between gating and entrainment mechanisms has met with varying degrees of success. Strong evidence in favour of entrainment was obtained for the coordination between the cell cycle and Cdc14 nucleolar sequestration and release [43]. Blocking the cell cycle by maintaining cyclin B at constant physiological levels did not block Cdc14 oscillations, ruling out a gating mechanism. The mechanisms governing other synchronized systems, such as the synchronization between the cell cycle and circadian rhythm, have not reached consensus. Among the studies in favour of entrainment [15,33,44], Feillet *et al.* [44] reset the circadian clock using a glucocorticoid agonist and observed a variety of coupled states between the clock and the cell cycle (1 : 1, 1 : 2, 3 : 2), supporting an entrainment mechanism and aligned with computational studies [15]. Among the studies suggesting a gating mechanism [45–47], Laranjeiro *et al.* [45] manipulated light–dark cycles in zebrafish cells to vary the period of the circadian clock and observed an exclusive effect on the length of G1 with S/G2/M phases remaining relatively constant. As articulated above, impact over specific phases of the follower oscillator is characteristic of a gating mechanism.

Most studies in favour of entrainment between the circadian and cell cycle oscillators consider unidirectional entrainment with the circadian clock unidirectionally entraining the cell cycle (figure 1*d*). Circadian rhythms persisted in cells whose division was inhibited, initially suggesting unidirectional entrainment [48]. However, the possibility of bi-directional entrainment has not been ruled out [15]. It is plausible that altered cell cycle dependent changes in transcription or reduced protein concentrations after cell division may affect the circadian phase [49–51]. Future work using synthetic biology approaches to study isolated or minimally coupled oscillators could help elucidate both the mechanisms leading to synchrony in other systems (entrainment versus gating) and the directionality of entrainment (uni- versus bi-directionality).

5. Different biological oscillators vary in their propensity for entrainment

The study of entrainment can be greatly simplified by studying the response of an oscillator to a single pulse instead of to a periodic input. Such single perturbation is often shorter than the period of the oscillating system and can cause a shift in the original phase, either advancing or retarding the oscillations

depending on its start time relative to the phase of the natural oscillator. A common way to capture this dependency is through phase response curves (PRCs) [52,53]. The features of a PRC, such as its magnitude (amplitude in the *y*-axis), zero points (intercept of the *x*-axis) and discontinuities (i.e. phase singularities), impact the propensity for entrainment [1] (box 2).

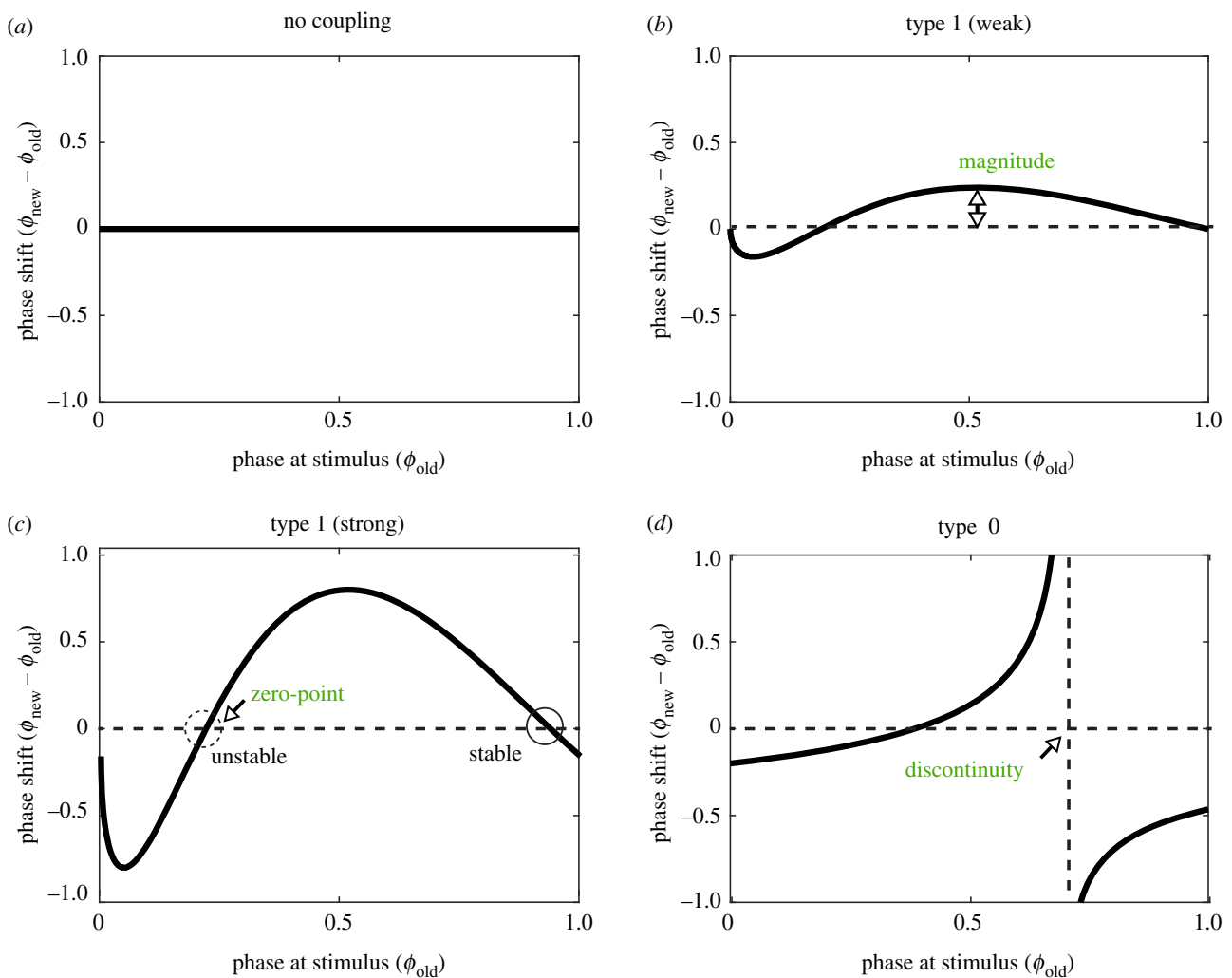
A system's PRCs can change by varying the amplitude or duration of the external pulse [54]. Stable entrainment of NF- κ B oscillators (figure 2*f*) requires a minimal duration and minimal concentration of the synchronizing TNF pulse [55]. The sensitivity of fly circadian clock has been tested by varying the duration of light pulses, which mainly affect the degradation of the clock gene *TIM* and can entrain the system in all tissues (figure 2*a*) (both neuronal and non-neuronal tissues in *Drosophila* are photoreceptive) [9,10]. Short light pulses lead to a PRC with a small magnitude and a continuous transition between phase advance and phase delay (called 'type 1' resetting) [56] (box 2*b*). As the duration of the light pulse increases, the PRC's magnitude increases (box 2*c*) and may show discontinuity between phase advance and phase delay regions (called 'type 0' resetting) (box 2*d*). Around this discontinuity, the new phase after perturbation is highly sensitive to the old phase and may lead to complex behaviours of the system, such as chaos [57,58]. Similarly, the PRC of the circadian clock of cyanobacteria is continuous, i.e. lacks phase singularities, under a short temperature pulse [59]. The phase shift increases with the increase of pulse duration, while the transition between phase advance and delay becomes sharper. Consequently, the PRC exhibits a singularity point above a certain pulse duration. This phase singularity may cause population-level arrhythmicity when certain perturbations cause stochastic phases of oscillations in individual cells. In the case of cardiac pacemaker, discontinuity of PRC has been suggested to lead to cardiac arrhythmias [1,60,61].

Absence of phase shift, i.e. flat curve (box 2*a*), indicates no possibility for entrainment. PRCs with low (box 2*b*) or high (box 2*c*) magnitude on phase shift indicate lower or higher propensity for entrainment. A PRC may have multiple zero points, with well-known examples in the circadian system [48,57,59,62], meaning that a perturbation administered when the system's phase is at these points will not cause phase change [1]. A PRC may have multiple zeros corresponding to distinct entrainment modes. The slope at a zero point of a PRC dictates the stability of this entrainment state: a negative or positive slope predicts stable or unstable (i.e. further from or closer to uncoupling regions) entrainment, respectively. Last, PRCs can exhibit phase singularities (marked by a vertical line in box 2*d*), at which the phase resetting is very sensitive to the phase at stimulus.

6. The impact of entrainment on heterogeneity between individual cells

During entrainment, each single-cell oscillator locks to the external input (figure 1*d*). If the population of cells is initially heterogeneous in its oscillations, phase locking results in a loss of heterogeneity. For example, the glycolytic oscillations of isolated yeast cells (figure 2*e*) display a broad distribution of frequencies around half a minute [63]. Periodic cyanide input can entrain this heterogeneous population through phase shifting (see section above). All cells' oscillations become synchronized after the first cyanide pulse [12]

Box 2. The inclination of a system to be entrained depends on its sensitivity to the perturbation and can be interpreted from the shape and properties of the phase response curves (PRCs). A PRC describes the magnitude of phase changes (also called phase shift) by plotting how much the oscillation is shifted in time (i.e. new phase ϕ_{new} minus unperturbed old phase ϕ_{old} on the y -axis) as a function of the phase at which it is received (x -axis).



reducing population heterogeneity. Furthermore, both robust and weak (or non-) oscillating cells entrain to the periodic input, further reducing population heterogeneity.

Entrainment through intercellular communication (figure 1c) can also decrease cell-to-cell variability. During glycolysis in yeast cultures, acetaldehyde secreted by cells induces synchronization of metabolic oscillations (even converting non-oscillating cells to an oscillatory state) [64]; this effect occurs only above a minimal cell density [3,19,63,65]. Similarly, dissociated cells of many organs show high heterogeneity of their oscillations. Isolated individual sinoatrial node cardiac pacemaker cells have varying periods [66–69], but at high density, they exhibit the stereotypical 80 beats per minute [61,70]. Dispersed cultures of suprachiasmatic nucleus (SCN) neurons behave as non-synchronous single-cell oscillators and fire with widely varying circadian periods distinct from 24 h [71,72]. When neurons are maintained at high density, either in explants or dispersals, their periods synchronize [5,73] to achieve tissue-level synchrony, in which all cells oscillate at the stereotypical 24 h period. The secreted factor synchronizing circadian oscillations of SCN neurons is less clear than that for glycolysis. Separation of the dorsal and ventral SCN resulted in a loss of synchrony of the neural rhythms of the dorsal (but not ventral) SCN,

suggesting that a neurotransmitter released by the ventral SCN maintains synchrony throughout the SCN [74]. Indeed, some of the candidate synchronizing factors (neurotransmitters γ -aminobutyric acid, vasoactive intestinal peptide and gastrin-releasing peptide) changed the firing rate of dorsal SCN neurons [73,75,76].

When an initially heterogeneous cell population entrains through intercellular communication to become more homogeneous, it is not clear which cells will dominate the final behaviour of the population. When two cell suspensions of yeast oscillating out of phase were mixed, synchronization was dominated by the culture whose NADH levels were decreasing [77,78]. A different mechanism operated when non-synchronized oscillating cardiomyocytes were placed into physical contact through a connected agarose micro-chamber [79,80] to synchronize their beating. The cells synchronized to the one showing smaller fluctuations in beating. Thus, it appears that different mechanisms can be employed to determine which of two functionally equivalent oscillators dominates during entrainment.

In some cases, entrainment can increase population heterogeneity, for example when it involves bi-stable responses. This phenomenon has been observed following periodic stimulation of the NF- κ B pathway by the cytokine

TNF- α (figure 2f) [40]. A single pulse of TNF- α leads to NF- κ B oscillations with a period of 90 min. When the TNF- α signal was provided in an oscillatory manner, cells entrained at multiple ratios for a given TNF periodic input. The multi-stability in entrainment ratios depended on the input frequency. When the stimulation period corresponded to the original unaltered period, 90 min, the population entrained nearly homogeneously with a 90 min phase-locked oscillation (1 : 1 ratio). By contrast, during a 150 min stimulation period cells showed a mixture of cellular responses including 150 min oscillation (1 : 1), 75 min oscillation (1 : 2), or without phase locking. Multi-stability rose from extrinsic noise (variation in signalling parameters between cells) that caused a significant broadening of the entrainment Arnold tongues regions (see box 1), revealing an important function of noise in allowing for a heterogeneous response to a periodic stimulus [40].

7. Plausible functions of entrainment

Entrainment is a ubiquitous phenomenon in biology, found across species and in diverse systems. In some cases, the function of entrainment is clear. For example, systems in which a population of cells synchronizes to achieve a specific coordinated task, such as the synchronization of SCN neurons to light–dark cycles provides further synchronization in downstream organs [81]. In the cardiac rhythms, synchronization of pacemaker cells provides blood circulation [61]. In systems in which cellular information is encoded in frequency, such as the frequency modulation of the transcription factor Crz1 by extracellular calcium concentration ensuring appropriate downstream expression [82] or frequency of motor protein-based oscillations in neurons is a read-out for axonal length [83]; a potential function for entrainment is to strengthen such a modality of signalling. However, in many other systems, the biological function for entrainment remains unclear. For example, despite its ubiquity, the physiological function of glycolytic oscillations and their entrainment are still uncertain [63]. In addition, while entrainment of NF- κ B was shown to coordinate the transcriptional response downstream of NF- κ B [40], entrainment in this system was achieved artificially through period stimulation by TNF- α , which is not known to oscillate *in vivo*.

Entrainment of biological clocks may also play an important role during evolution. As one example, the oscillation of cyclin-dependent kinase (CDK) activity drives other periodic events, such as DNA replication and chromosome separation, during the cell cycle. Interestingly, CDKs seem to have appeared late during evolution [84], raising the question as to how cells synchronize the series of events required for proliferation prior to CDK emergence. Recent studies in yeast identified several processes that show periodic behaviours even in the absence of CDK oscillator. These CDK-independent oscillators include budding, DNA replication, centrosome duplication, transcription and Cdc14 release

[85–89]. Intriguingly, their intrinsic periods are close to the normal cell cycle duration. It has been speculated that cell cycle processes may be intrinsically oscillatory before the emergence of CDK, and these oscillators entrain each other to create an aggregate rhythm [43]. The master CDK oscillator may have evolved to regulate other oscillators in order to yield a stable entrainment structure. This satisfies the evolutionary requirement of utility of intermediate forms [90]. Entrainment of autonomous oscillators could have been important in early cell cycle evolution, raising the possibility that it plays a role in promoting a stable cell cycle rhythm in modern eukaryotes.

8. Future perspectives

Many aspects of entrainment remain unexplored mainly due to the complex network interactions controlling and connecting oscillations in biology. One approach that can be useful in disentangling interconnected oscillatory systems is synthetic biology. Synthetic biology allows precise control of entrainment networks and has been used to study extremely complex systems such as a built-in circadian clock [81,91] or quorum sensing [4]. In the future, building synthetic oscillators that are heavily intertwined in nature (such as the cell cycle or the Cdc14 oscillators) could elucidate the mechanisms behind their coupling and avoid the use of genetic manipulation in their original natural systems. Finally, the potential of an oscillatory system to be entrained has not been explored in many networks, even in well studied oscillatory systems such as p53 or Msn2 [29,92,93], both having the potential to be entrained using distinct combinations of drugs.

Advances in technologies such as microfluidic devices, microscopy and optical traps allow precise spatial and temporal control of a cell's environment and facilitate single-cell measurements of oscillatory behaviours. Synthetic biology approaches along with technological advances will be essential to explore fundamental questions of entrainment such as the molecular determinants of the entrainment capability of a system and the functional consequences of entrainment.

Data accessibility. This article has no additional data.

Authors' contributions. A.Ji.: conceptualization, writing—original draft, writing—review and editing; Y.L.: conceptualization, writing—original draft, writing—review and editing; A.Ja.: conceptualization, supervision, writing—original draft, writing—review and editing; G.L.: conceptualization, supervision, writing—original draft, writing—review and editing.

All authors gave final approval for publication and agreed to be held accountable for the work performed therein.

Competing interests. We declare we have no competing interests.

Funding. Research in the Lahav Lab is supported by National Institutes of Health grant no. NIH R35 GM139572 and by the Ludwig Center at Harvard. Research in the Lu Lab is supported by National Institutes of Health grant no. NIH R01 GM134064-01.

Acknowledgements. We would like to thank Bill Jia and Adrian Granada for discussions.

References

1. Winfree A. 2001 *The geometry of the biological time*. New York, NY: Springer.
2. Huygens C. 1665 Correspondence 1664–1665. In *Oeuvres complètes de Christiaan Huygens vol. V (La societe hollandaise des sciences, 1893)*. The Hague, The Netherlands: M. Nijhoff.

3. De Monte S, D'Ovidio F, Danø S, Sørensen PG. 2007 Dynamical quorum sensing: population density encoded in cellular dynamics. *Proc. Natl Acad. Sci. USA* **104**, 18 377–18 381. (doi:10.1073/pnas.0706089104)
4. García-Ojalvo J, Elowitz MB, Strogatz SH. 2004 Modeling a synthetic multicellular clock: repressilators coupled by quorum sensing. *Proc. Natl Acad. Sci. USA* **101**, 10 955–10 960. (doi:10.1073/pnas.0307095101)
5. Nakamura W, Honma S, Shirakawa T, Honma KI. 2001 Regional pacemakers composed of multiple oscillator neurons in the rat suprachiasmatic nucleus. *Eur. J. Neurosci.* **14**, 666–674. (doi:10.1046/j.0953-816x.2001.01684.x)
6. Ypey DL, Clapham DE, DeHaan RL. 1979 Development of electrical coupling and action potential synchrony between paired aggregates of embryonic heart cells. *J. Membr. Biol.* **51**, 75–96. (doi:10.1007/BF01869344)
7. Jalife J. 1984 Mutual entrainment and electrical coupling as mechanisms for synchronous firing of rabbit sino-atrial pace-maker cells. *J. Physiol.* **356**, 221–243. (doi:10.1113/jphysiol.1984.sp015461)
8. Michaels DC, Matyas EP, Jalife J. 1986 Dynamic interactions and mutual synchronization of sinoatrial node pacemaker cells: a mathematical model. *Circ. Res.* **58**, 706–720. (doi:10.1161/01.RES.58.5.706)
9. Plautz JD, Kaneko M, Hall JC, Kay SA. 1997 Independent photoreceptive circadian clocks throughout *Drosophila*. *Science* **278**, 1632–1635. (doi:10.1126/science.278.5343.1632)
10. Zeng H, Qian Z, Myers MP, Rosbash M. 1996 A light-entrainment mechanism for the *Drosophila* circadian clock. *Nature* **380**, 129–135. (doi:10.1038/380129a0)
11. Strogatz SH. 2000 From Kuramoto to Crawford: exploring the onset of synchronization in populations of coupled oscillators. *Physica D* **143**, 1–20. (doi:10.1016/S0167-2789(00)00094-4)
12. Gustavsson AK, Adiels CB, Mehlig B, Goksör M. 2015 Entrainment of heterogeneous glycolytic oscillations in single cells. *Sci. Rep.* **5**, 9404. (doi:10.1038/srep09404)
13. Almendral J, Caulier-Cisterna R, Rojo-Álvarez JL. 2013 Resetting and entrainment of reentrant arrhythmias: part i: concepts, recognition, and protocol for evaluation: surface ECG versus intracardiac recordings. *Pacing Clin. Electrophysiol.* **36**, 508–532. (doi:10.1111/pace.12064)
14. Rosato E, Tauber E, Kyriacou CP. 2006 Molecular genetics of the fruit-fly circadian clock. *Eur. J. Hum. Genet.* **14**, 729–738. (doi:10.1038/sj.ejhg.5201547)
15. Yan J, Goldbeter A. 2019 Robust synchronization of the cell cycle and the circadian clock through bidirectional coupling. *J. R. Soc. Interface* **16**, 20190376. (doi:10.1098/rsif.2019.0376)
16. Azzam R, Chen SL, Shou W, Mah AS, Alexandru G, Nasmyth K, Annan RS, Carr SA, Deshaies RJ. 2004 Phosphorylation by cyclin B-Cdk underlies release of mitotic exit activator Cdc14 from the nucleolus. *Science* **305**, 516–519. (doi:10.1126/science.1099402)
17. Lakatta EG, Yaniv Y, Maltsev VA. 2014 Cardiac impulse is initiated by a coupled system of membrane ion channels and Ca²⁺ cycling proteins. In *Cardiac electrophysiology: from cell to bedside*, 6th edn (eds DP Zipes, J Jalife), pp. 243–252. Philadelphia, PA: Saunders/Elsevier.
18. Yaniv Y, Lakatta EG, Maltsev VA. 2015 From two competing oscillators to one coupled-clock pacemaker cell system. *Front. Physiol.* **6**, 28. (doi:10.3389/fphys.2015.00028)
19. Bier M, Bakker BM, Westerhoff HV. 2000 How yeast cells synchronize their glycolytic oscillations: a perturbation analytic treatment. *Biophys. J.* **78**, 1087–1093. (doi:10.1016/S0006-3495(00)76667-7)
20. Gustavsson AK, Van Niekerk DD, Adiels CB, Kooi B, Goksör M, Snoep JL. 2014 Allosteric regulation of phosphofruktokinase controls the emergence of glycolytic oscillations in isolated yeast cells. *FEBS J.* **281**, 2784–2793. (doi:10.1111/febs.12820)
21. Tay S, Hughey JJ, Lee TK, Lipniacki T, Quake SR, Covert MW. 2010 Single-cell NF- κ B dynamics reveal digital activation and analogue information processing. *Nature* **466**, 267–271. (doi:10.1038/nature09145)
22. Hoffmann A, Levchenko A, Scott ML, Baltimore D. 2002 The I κ B-NF- κ B signaling module: temporal control and selective gene activation. *Science* **298**, 1241–1245. (doi:10.1126/science.1071914)
23. Nelson DE *et al.* 2004 Oscillations in NF- κ B signaling control the dynamics of gene expression. *Science* **306**, 704–708. (doi:10.1126/science.1099962)
24. Tsai TYC, Yoon SC, Ma W, Pomerening JR, Tang C, Ferrell JE. 2008 Robust, tunable biological oscillations from interlinked positive and negative feedback loops. *Science* **321**, 126–139. (doi:10.1126/science.1156951)
25. Hardin PE, Hall JC, Rosbash M. 1990 Feedback of the *Drosophila* period gene product on circadian cycling of its messenger RNA levels. *Nature* **343**, 536–540. (doi:10.1038/343536a0)
26. Visintin R, Hwang ES, Amon A. 1999 Cfl1 prevents premature exit from mitosis by anchoring Cdc14 phosphatase in the nucleolus. *Nature* **398**, 818–823. (doi:10.1038/19775)
27. Duysens LN, Ames J. 1957 Fluorescence spectrophotometry of reduced phosphopyridine nucleotide in intact cells in the near-ultraviolet and visible region. *Biochim. Biophys. Acta* **24**, 19–26. (doi:10.1016/0006-3002(57)90141-5)
28. Toescu EC. 1995 Temporal and spatial heterogeneities of Ca²⁺ signaling: mechanisms and physiological roles. *Am. J. Physiol.* **269**, G173–G185. (doi:10.1152/ajpgi.1995.269.2.g173)
29. Hao N, O'Shea EK. 2012 Signal-dependent dynamics of transcription factor translocation controls gene expression. *Nat. Struct. Mol. Biol.* **19**, 31–40. (doi:10.1038/nsmb.2192)
30. Nunemaker CS, Satin LS. 2014 Episodic hormone secretion: a comparison of the basis of pulsatile secretion of insulin and GnRH. *Endocrine* **47**, 49–63. (doi:10.1007/s12020-014-0212-3)
31. Wright T, Gillespie LN, O'Leary SJ, Needham K. 2016 Firing frequency and entrainment maintained in primary auditory neurons in the presence of combined BDNF and NT3. *Sci. Rep.* **6**, 1. (doi:10.1038/s41598-016-0001-8)
32. Borghans JAM, Dupont G, Goldbeter A. 1997 Complex intracellular calcium oscillations. a theoretical exploration of possible mechanisms. *Biophys. Chem.* **66**, 25–41. (doi:10.1016/S0301-4622(97)00010-0)
33. Gérard C, Goldbeter A. 2012 Entrainment of the mammalian cell cycle by the circadian clock: modeling two coupled cellular rhythms. *PLoS Comput. Biol.* **8**, e1002516. (doi:10.1371/journal.pcbi.1002516)
34. Guevara MR, Glass L, Shrier A. 2011 Phase locking, period-doubling bifurcations, and irregular dynamics in periodically stimulated cardiac cells. *Science* **214**, 1350–1353. (doi:10.1126/science.7313693)
35. Hancioglu B, Tyson JJ. 2012 A mathematical model of mitotic exit in budding yeast: the role of Polo kinase. *PLoS ONE* **7**, e30810. (doi:10.1371/journal.pone.0030810)
36. Leloup JC, Goldbeter A. 2003 Toward a detailed computational model for the mammalian circadian clock. *Proc. Natl Acad. Sci. USA* **100**, 7051–7056. (doi:10.1073/pnas.1132112100)
37. Panda S, Hogenesch JB, Kay SA. 2002 Circadian rhythms from flies to human. *Nature* **417**, 329–335. (doi:10.1038/417329a)
38. Ballesta A, Innominato PF, Dallmann R, Rand DA, Lévi FA. 2017 Systems chronotherapeutics. *Pharmacol. Rev.* **69**, 161–199. (doi:10.1124/pr.116.013441)
39. Gérard C, Goldbeter A. 2009 Temporal self-organization of the cyclin/Cdk network driving the mammalian cell cycle. *Proc. Natl Acad. Sci. USA* **106**, 21 643–21 648. (doi:10.1073/pnas.0903827106)
40. Kellogg RA, Tay S. 2015 Noise facilitates transcriptional control under dynamic inputs. *Cell* **160**, 381–392. (doi:10.1016/j.cell.2015.01.013)
41. Edmunds LN. 1988 *Cellular and molecular bases of biological clocks: models and mechanisms for circadian timekeeping*. New York, NY: Springer.
42. Wille JJ, Ehret CF. 1968 Light synchronization of an endogenous circadian rhythm of cell division in tetrahymena. *J. Protozool.* **15**, 785–789. (doi:10.1111/j.1550-7408.1968.tb02214.x)
43. Lu Y, Cross FR. 2010 Periodic cyclin-cdk activity entrains an autonomous cdc14 release oscillator. *Cell* **141**, 268–279. (doi:10.1016/j.cell.2010.03.021)
44. Feillet C *et al.* 2014 Phase locking and multiple oscillating attractors for the coupled mammalian clock and cell cycle. *Proc. Natl Acad. Sci. USA* **111**, 9828–9833. (doi:10.1073/pnas.1320474111)
45. Laranjeiro R, Tamai TK, Letton W, Hamilton N, Whitmore D. 2018 Circadian clock synchronization of

- the cell cycle in zebrafish occurs through a gating mechanism rather than a period-phase locking process. *J. Biol. Rhythms* **33**, 137–150. (doi:10.1177/0748730418755583)
46. Mori T, Binder B, Johnson CH. 1996 Circadian gating of cell division in cyanobacteria growing with average doubling times of less than 24 hours. *Proc. Natl Acad. Sci. USA* **93**, 10 183–10 188. (doi:10.1073/pnas.93.19.10183)
 47. Yang Q, Pando BF, Dong G, Golden SS, Van Oudenaarden A. 2010 Circadian gating of the cell cycle revealed in single cyanobacterial cells. *Science* **327**, 1522–1526. (doi:10.1126/science.1181759)
 48. Matsuo T, Yamaguchi S, Mitsui S, Emi A, Shimoda F, Okamura H. 2003 Control mechanism of the circadian clock for timing of cell division in vivo. *Science* **302**, 255–259. (doi:10.1126/science.1086271)
 49. Bieler J, Cannavo R, Gustafson K, Gobet C, Gatfield D, Naef F. 2014 Robust synchronization of coupled circadian and cell cycle oscillators in single mammalian cells. *Mol. Syst. Biol.* **10**, 739. (doi:10.1525/msb.20145218)
 50. Nagoshi E, Saini C, Bauer C, Laroche T, Naef F, Schibler U. 2004 Circadian gene expression in individual fibroblasts: cell-autonomous and self-sustained oscillators pass time to daughter cells. *Cell* **119**, 693–705. (doi:10.1016/j.cell.2004.11.015)
 51. Zopf CJ, Quinn K, Zeidman J, Maheshri N. 2013 Cell-cycle dependence of transcription dominates noise in gene expression. *PLoS Comput. Biol.* **9**, e1003161. (doi:10.1371/journal.pcbi.1003161)
 52. Fei C, Cao Y, Ouyang Q, Tu Y. 2018 Design principles for enhancing phase sensitivity and suppressing phase fluctuations simultaneously in biochemical oscillatory systems. *Nat. Commun.* **9**, 1434. (doi:10.1038/s41467-018-03826-4)
 53. Granada A, Hennig RM, Ronacher B, Kramer A, Herzel H. 2009 Phase response curves: elucidating the dynamics of coupled oscillators. *Methods Enzymol.* **454**, 1–27. (doi:10.1016/S0076-6879(08)03801-9)
 54. Masuda K, Tokuda IT, Nakamichi N, Fukuda H. 2021 The singularity response reveals entrainment properties of the plant circadian clock. *Nat. Commun.* **12**, 864. (doi:10.1038/s41467-021-21167-7)
 55. Zambrano S, De Toma I, Piffer A, Bianchi ME, Agresti A. 2016 NF- κ B oscillations translate into functionally related patterns of gene expression. *eLife* **5**, e09100. (doi:10.7554/eLife.09100)
 56. Johnson CH. 1999 Forty years of PRCs: what have we learned? *Chronobiol. Int.* **16**, 711–743. (doi:10.3109/07420529909016940)
 57. Granada A, Herzel H. 2009 How to achieve fast entrainment? The timescale to synchronization. *PLoS ONE* **4**, e7057. (doi:10.1371/journal.pone.0007057)
 58. Heltberg ML, Krishna S, Kadanoff LP, Jensen MH. 2021 A tale of two rhythms: locked clocks and chaos in biology. *Cell Syst.* **12**, 291–303. (doi:10.1016/j.cels.2021.03.003)
 59. Gan S, O'Shea EK. 2017 An unstable singularity underlies stochastic phasing of the circadian clock in individual cyanobacterial cells. *Mol. Cell* **67**, 659–672. (doi:10.1016/j.molcel.2017.07.015)
 60. Abramovich-Sivan S, Akselrod S. 1998 A single pacemaker cell model based on the phase response curve. *Biol. Cybern.* **79**, 67–76. (doi:10.1007/s004220050459)
 61. Fenske S *et al.* 2020 cAMP-dependent regulation of HCN4 controls the tonic entrainment process in sinoatrial node pacemaker cells. *Nat. Commun.* **11**, 5555. (doi:10.1038/s41467-020-19304-9)
 62. Diekman CO, Bose A. 2016 Entrainment maps: a new tool for understanding properties of circadian oscillator models. *J. Biol. Rhythms* **31**, 598–616. (doi:10.1177/0748730416662965)
 63. Weber A, Zuschratter W, Hauser MJB. 2020 Partial synchronisation of glycolytic oscillations in yeast cell populations. *Sci. Rep.* **10**, 1. (doi:10.1038/s41598-019-56847-4)
 64. Richard P, Bakker BM, Teusink B, Van Dam K, Westerhoff HV. 1996 Acetaldehyde mediates the synchronization of sustained glycolytic oscillations in populations of yeast cells. *Eur. J. Biochem.* **235**, 238–241. (doi:10.1111/j.1432-1033.1996.00238.x)
 65. Aldridge J, Pye EK. 1976 Cell density dependence of oscillatory metabolism. *Nature* **259**, 670–671. (doi:10.1038/259670a0)
 66. Kim MS *et al.* 2018 Heterogeneity of calcium clock functions in dormant, dysrhythmically and rhythmically firing single pacemaker cells isolated from SA node. *Cell Calcium* **74**, 168–179. (doi:10.1016/j.ceca.2018.07.002)
 67. Lyashkov AE, Juhaszova M, Dobrzynski H, Vinogradova TM, Maltsev VA, Juhasz O, Spurgeon HA, Sollott SJ, Lakatta EG. 2007 Calcium cycling protein density and functional importance to automaticity of isolated sinoatrial nodal cells are independent of cell size. *Circ. Res.* **100**, 1723–1731. (doi:10.1161/CIRCRESAHA.107.153676)
 68. Manfredi O, Tsutsui K, Ziman B, Stern MD, Lakatta EG, Maltsev VA. 2018 Electrophysiological heterogeneity of pacemaker cells in the rabbit intercaval region, including the SA node: insights from recording multiple ion currents in each cell. *Am. J. Physiol. Heart Circ. Physiol.* **314**, H403–H414. (doi:10.1152/ajpheart.00253.2016)
 69. Wilders R, Jongsma HJ. 1993 Beating irregularity of single pacemaker cells isolated from the rabbit sinoatrial node. *Biophys. J.* **65**, 2601–2613. (doi:10.1016/S0006-3495(93)81289-X)
 70. Easterling M, Rossi S, Mazzella AJ, Bressan M. 2021 Assembly of the cardiac pacemaking complex: electrogenic principles of sinoatrial node morphogenesis. *J. Cardiovasc. Dev. Dis.* **8**, 40. (doi:10.3390/jcdd8040040)
 71. Herzog ED, Takahashi JS, Block GD. 1998 Clock controls circadian period in isolated suprachiasmatic nucleus neurons. *Nat. Neurosci.* **1**, 708–713. (doi:10.1038/3708)
 72. Welsh DK, Logothetis DE, Meister M, Reppert SM. 1995 Individual neurons dissociated from rat suprachiasmatic nucleus express independently phased circadian firing rhythms. *Neuron* **14**, 697–706. (doi:10.1016/0896-6273(95)90214-7)
 73. Aton SJ, Herzog ED. 2005 Come together, right... now: synchronization of rhythms in a mammalian circadian clock. *Neuron* **48**, 531–534. (doi:10.1016/j.neuron.2005.11.001)
 74. Yamaguchi S, Isejima H, Matsuo T, Okura R, Yagita K, Kobayashi M, Okamura H. 2003 Synchronization of cellular clocks in the suprachiasmatic nucleus. *Science* **302**, 1408–1412. (doi:10.1126/science.1089287)
 75. Liu C, Reppert SM. 2000 GABA synchronizes clock cells within the suprachiasmatic circadian clock. *Neuron* **25**, 123–128. (doi:10.1016/S0896-6273(00)80876-4)
 76. Piggins HD, Antle MC, Rusak B. 1995 Neuropeptides phase shift the mammalian circadian pacemaker. *J. Neurosci.* **15**, 5612–5622. (doi:10.1523/JNEUROSCI.15-08-05612.1995)
 77. Danø S, Hynne F, De Monte S, d'Ovidio F, Sørensen PG, Westerhoff H. 2001 Synchronization of glycolytic oscillations in a yeast cell population. *Faraday Discuss.* **120**, 261–276. (doi:10.1039/b103238k)
 78. Pye E. 1969 Biochemical mechanisms underlying the metabolic oscillations in yeast. *Can. J. Bot.* **2**, 271–285. (doi:10.1139/b69-040)
 79. Kojima K, Kaneko T, Yasuda K. 2006 Role of the community effect of cardiomyocyte in the entrainment and reestablishment of stable beating rhythms. *Biochem. Biophys. Res. Commun.* **351**, 209–215. (doi:10.1016/j.bbrc.2006.10.037)
 80. Yasuda K. 2020 Dominant rule of community effect in synchronized beating behavior of cardiomyocyte networks. *Biophys. Rev.* **12**, 481–501. (doi:10.1007/s12551-020-00688-3)
 81. Reppert SM, Weaver DR. 2002 Coordination of circadian timing in mammals. *Nature* **418**, 935–941. (doi:10.1038/nature00965)
 82. Cai L, Dalal CK, Elowitz MB. 2008 Frequency-modulated nuclear localization bursts coordinate gene regulation. *Nature* **455**, 485–490. (doi:10.1038/nature07292)
 83. Bressloff PC, Karamched BR. 2015 A frequency-dependent decoding mechanism for axonal length sensing. *Front. Cell. Neurosci.* **9**, 281. (doi:10.3389/fncel.2015.00281)
 84. Krylov DM, Nasmyth K, Koonin EV. 2003 Evolution of eukaryotic cell cycle regulation: stepwise addition of regulatory kinases and late advent of the CDKs. *Curr. Biol.* **13**, 173–177. (doi:10.1016/S0960-9822(03)00008-3)
 85. Gard DL, Hafezi S, Zhang T, Doxsey SJ. 1990 Centrosome duplication continues in cycloheximide-treated *Xenopus blastulae* in the absence of a detectable cell cycle. *J. Cell Biol.* **110**, 2033–2042. (doi:10.1083/jcb.110.6.2033)
 86. Haase SB, Winey M, Reed SI. 2001 Multi-step control of spindle pole body duplication by cyclin-dependent kinase. *Nat. Cell Biol.* **3**, 38–42. (doi:10.1038/35050543)
 87. Haase SB, Reed SI. 1999 Evidence that a free-running oscillator drives G1 events in the budding

- yeast cell cycle. *Nature* **401**, 394–397. (doi:10.1038/43927)
88. McClelland ML, O'Farrell PH. 2008 RNAi of mitotic cyclins in *Drosophila* uncouples the nuclear and centrosome cycle. *Curr. Biol.* **18**, 245–254. (doi:10.1016/j.cub.2008.01.041)
 89. Sluder G, Miller FJ, Cole R, Rieder CL. 1990 Protein synthesis and the cell cycle: centrosome reproduction in sea urchin eggs is not under translational control. *J. Cell Biol.* **110**, 2025–2032. (doi:10.1083/jcb.110.6.2025)
 90. Darwin C. 1859 *On the origin of species by means of natural selection*. London, UK: John Murray.
 91. Mondragón-Palomino O, Danino T, Selimkhanov J, Tsimring L, Hasty J. 2011 Entrainment of a population of synthetic genetic oscillators. *Science* **333**, 1315–1319. (doi:10.1126/science.1205369)
 92. Harton MD, Koh WS, Bunker AD, Singh A, Batchelor E. 2019 P53 pulse modulation differentially regulates target gene promoters to regulate cell fate decisions. *Mol. Syst. Biol.* **15**, e8685. (doi:10.15252/msb.20188685)
 93. Lahav G, Rosenfeld N, Sigal A, Geva-Zatorsky N, Levine AJ, Elowitz MB, Alon U. 2004 Dynamics of the p53-Mdm2 feedback loop in individual cells. *Nat. Genet.* **36**, 147–150. (doi:10.1038/ng1293)