

## Quantum time crystals with programmable disorder in higher dimensions

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We present fresh evidence for the presence of discrete quantum time crystals in two spatial dimensions. Discrete time crystals are intricate quantum systems that break discrete time translation symmetry in driven quantum many-body systems undergoing nonequilibrium dynamics. They are stabilized by many-body localization arising from disorder. We directly target the thermodynamic limit using instances of infinite tensor network states, and we implement disorder in a translationally invariant setting by introducing auxiliary systems at each site. We discuss how such disorder can be realized in programmable quantum simulators: This gives rise to the interesting situation in which a classical tensor network simulation can contribute to devising a blueprint of a quantum simulator featuring prethermal time crystalline dynamics, one that will ultimately have to be built in order to explore the stability of this phase of matter for long times.

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### I. INTRODUCTION

A *time crystal* is a periodic structure that not only repeats itself periodically in space, as regular crystals do, but also in time [1,2]. The idea of a crystal in space and time has been proposed by Wilczek [3] using superconducting rings and persistent currents. Not much later, it became clear that serious obstructions need to be overcome: Ground states of local Hamiltonians can provably not host a spontaneous breaking of time-translation symmetry [4,5]. So achieving a quantum time crystal becomes possible only in two special scenarios: (i) using a nonlocal Hamiltonian, or (ii) resorting to nonequilibrium dynamics. Indeed, one can construct Hamiltonian models whose ground states feature quantum time crystals, albeit at the price of an intricate interaction pattern that is presumably excessively challenging to achieve even in highly engineered quantum systems [6].

As an alternative route much more amenable to experimental realization, one can resort to settings of *nonequilibrium physics* [7–11]. Specifically, systems undergoing a time-periodic driving in one spatial dimension have been identified as suitable candidates to exhibit discrete time translation symmetry breaking [1,2,8,11–15], referred to as *discrete time crystals*, going back to the seminal work of Khemani, Lazarides, Moessner, and Sondhi [8]. Indeed, this idea has led to the experimental observation of time crystals, both in one-dimensional systems of trapped ions using a programmable potential [16], and in large settings of dipolar spin impurities in diamond at room temperature [17]. A further obstacle that arises along the way in such periodically driven physical systems can be largely overcome. To avoid heating due to the driving, disorder comes to the rescue, giving rise

to *many-body localization* [18,19], which prevents thermalization [17,20–22], either for long (as in prethermal) or all times [23,24]. That is to say, by suitably combining disorder and periodic driving, the realization of this intricate state of matter is conceivable in synthetic quantum devices. Indeed, experimental realizations of such discrete time crystals can be viewed as instances of *dynamical quantum simulations* [25–27] realizing quantum technological devices [28]. In this sense, they can be seen as physical systems that allow us to probe intriguing features of interacting quantum many-body systems under precisely controlled conditions.

This state of affairs, discussed in the previous paragraph, is most interesting in situations in which state-of-the-art classical simulation techniques can provide strong evidence of the existence of discrete time crystals. At the same time, a full simulation in all aspects is out of reach with present classical simulation tools (and in fact presumably for all classical simulation tools, as there are complexity-theoretic obstructions to this effect [29]). This points to the exciting direction of exploring discrete time crystals in higher dimensions, settings that are conceivable in *programmable quantum simulators* [21,30–36], but for which the best known classical codes can only keep track of the relevant features in time. Indeed, this state of affairs gives rise to the interesting situation that one can build trust in the functioning of a quantum simulator in relevant regimes. At the same time, there is scope for explorations outside the classically computationally accessible realm: after all, the full quantum simulation is barely beyond the boundary of what can be assessed with classical computers. And along these lines, e.g., we can discriminate prethermal behavior for long times from genuine infinite-time time crystals [1,2].

In this work, we provide evidence for the existence of discrete time crystals in two spatial dimensions. This is possible by resorting to tensor network techniques such as *projected entangled pair states (PEPS)* [37–41], here applied to and pushed further to a regime involving interactions, disorder, and a time-periodic drive in two spatial dimensions. Unlike other numerical techniques, these tools are built to precisely capture genuine quantum correlations [42], and they can directly target the thermodynamic limit (iPEPS) [37,39,43], thus overcoming finite-size effects that are often encountered in classical simulations of quantum many-body systems. Targeting the thermodynamic limit is quite important, especially in the context of diagnosing time crystals, as they are only meaningful in the infinite-size limit [2]. For these reasons, iPEPS have been successfully used in the past to calculate ground states of challenging condensed-matter problems [44–50], thermal states [51–54], as well as nonequilibrium steady states [55]. Tensor networks, in general, have the limitation that they cannot handle large entanglement buildup associated with studying time evolution, and therefore they cannot go to very long times as this would require exponentially large bond dimensions.

In fact, the challenge to reach longer times for tensor network algorithms is rooted in fundamental complexity theoretic obstructions, arising from considerations in quantum information theory: The upshot is that the development of efficient classical simulation methods for all time-evolving local Hamiltonians is implausible because such a time evolution turns out to be ultimately as powerful as a full quantum computer. In principle, any quantum algorithm could be run on a local Hamiltonian quantum system undergoing such a time evolution. In more technical terms, local Hamiltonian time evolution is bounded-error quantum polynomial (BQP) complete [29], and therefore a universal classical simulation of local Hamiltonian evolution is not expected to be possible for all times. Despite this, the application of tensor networks has been extended to the challenging realm of keeping track of the time evolution in two-dimensional interacting systems [56–60] to unprecedented times recently.

Equipped with this machinery, we take on the problem of one of the most intricate quantum phases of matter: quantum time crystals in two dimensions. To be concrete, we do so by building upon recent efforts to capture time evolution in many-body localized systems in two spatial dimensions [58,60]. Indeed, the stability of many-body localization in two spatial dimensions is an open and challenging topic due to the argument of an avalanche mechanism caused by the presence of rare ergodic bubbles [61,62], and with existing tensor network methods, we may never be able to predict the stability of such a phase at infinite times.

In fact, again it is clear that any such classical simulation must face obstructions when aiming at assessing the behavior of the system at long times. That said, by substantial computational effort and suitable algorithm development, on the practical side, we can push things to show the existence of the phenomenon and to make a strong case for many-body localization in two spatial dimensions, within the realm of finite times that can be assessed. In fact, recently, there

has been more evidence of many-body localization in two spatial dimensions [59,63–66] using a variety of numerical techniques. In Ref. [60], we have shown the existence of many-body localization in two spatial dimensions for the longest times available to us based on the particle imbalance and the growth of local Rényi entropies with programmable discrete disorder. We found that taking a sufficient number of disorder levels as well as disorder strength is required to achieve localization, albeit for the available times. Besides, as we have also found in extensive numerical work, as the instability has its origin in thermal bubbles, the probability measure underlying the disorder matters significantly, a situation that can be easily accommodated with programmable disorder, say, in programmable quantum simulators, as we have them in mind here. Our current work is built upon this setup. To achieve our goal, we employ the evolution under a time-dependent Floquet Hamiltonian within an iPEPS framework.

Our advances are basically threefold: (i) We establish (at least prethermal) the presence of time-crystalline behavior in higher than one spatial dimension in unprecedented regimes over other known methods, thereby (ii) also demonstrating that tensor networks are able to make meaningful predictions in this regime in which interactions, disorder, and time-dependent drives come together, and (iii) we signify the power of programmable disorder to stabilize such behavior at longer times, thereby providing further impetus for such quantum simulators from the perspective of quantum information science to assess the stability of intricate quantum phases of matter.

## II. MODEL

In what follows, we consider a piecewise constant time-dependent family of Floquet Hamiltonians  $t \mapsto H_F(t)$  as

$$H_F(t) = \begin{cases} H_{\text{MBL}} & \text{if } t \in [0, T/2), \\ H_S & \text{if } t \in [T/2, T). \end{cases} \quad (1)$$

Specifically, the time evolution operator advancing the system by one period is taken as  $U = e^{-iH_S T/2} e^{-iH_{\text{MBL}} T/2}$ , with  $H_{\text{MBL}}$  being a static many-body localizing Hamiltonian described below. This entails that the total Hamiltonian is time-periodic with period  $T$ , where within one period of  $t \in [0, T)$ , first only  $H_{\text{MBL}}$  is active for times  $t \in [0, T/2)$ , while for the second part of the period  $t \in [T/2, T)$  only  $H_S$  acts on the systems.  $H_S$  is a spin-flip operator that flips each spin in the system at suitable moments in time. We choose  $H_S$  to be a spin-flip operator with some periodicity  $T$  as in Ref. [15] as

$$H_S = (2\pi/T - 2\epsilon) \sum_i S_i^x. \quad (2)$$

The action of this Hamiltonian is to flip each spin in the system in the spin- $z$  basis  $\{|\uparrow\rangle, |\downarrow\rangle\}$  at suitable times.  $S^x$  is the spin-1/2 operator in the spin- $x$  basis acting on each site, working in units  $\hbar = 1$ . The frequency of this flipping depends on the time period  $T > 0$ . The real constant parameter  $\epsilon$  determines the deviation from a perfect flip ( $|\uparrow\rangle \mapsto |\downarrow\rangle$ ) and vice versa: The persistence of a time crystal for nonzero values reflects a macroscopic rigidity since there is a periodic response in a measurable observable even when the

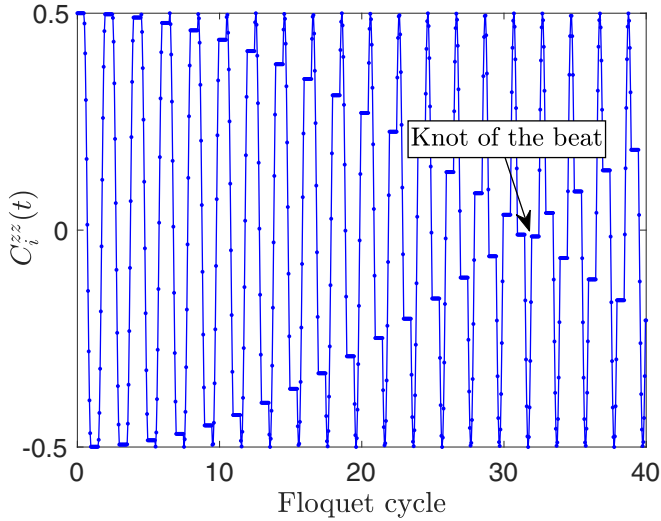


FIG. 1. Effect of imperfect flip ( $\epsilon = 0.5$ ) on a single uncoupled spin. One can see the emergence of a beating pattern along with the associated spikes leading to the formation of a knot at around Floquet cycle 32.

driving is not synchronized. A discrete time crystal is reflected by a response that is periodic in *integer multiples* of the driving period (here, period doubling given our specific setting). For the  $H_{\text{MBL}}$  contribution to the Hamiltonian, we choose a Heisenberg model on a square lattice with *discrete* disorder as

$$H_{\text{MBL}} = J \sum_{(i,j)} \vec{S}_i \cdot \vec{S}_j + \sum_i h_i S_i^z, \quad (3)$$

where  $h_i$  is drawn from a discrete uniform distribution with  $d_a$  levels between  $-h/2$  and  $h/2$  (the choice of naming the integer number of levels  $d_a$  will become apparent below). The above Hamiltonian is not only amenable to quantum simulation in programmable quantum simulators: it can at the same time be implemented in a perfectly translationally invariant fashion using iPEPS [60] by appending to every local physical spin-1/2 (denoted by subindex  $p$ ) an auxiliary one (denoted by subindex  $a$ ) with local Hilbert space dimension  $d_a$ . This *a posteriori* explains the above-mentioned choice of variable name. We have found evidence of localization based on the particle number very recently at sufficiently strong and enough levels of disorder. The disordered Heisenberg model is a paradigmatic model for many-body localization, a feature that is preserved under discrete disorder. In our tensor network simulation, briefly speaking, we start off from an initial state that is a tensor product of the physical initial state (local dimension  $d_p = 2$ ) and an auxiliary initial state (local dimension  $d_a$ ), i.e., a state vector  $|\Psi_0\rangle = |\psi_{p_0}\rangle \otimes |\psi_{a_0}\rangle$ , where  $|\psi_{a_0}\rangle = |\dots, +, +, +, +, \dots\rangle$  and  $|+\rangle = d_a^{-1/2}(\sum_{s=1}^{d_a} |s\rangle)$ ,  $s$  being the integer reflecting the allowed spin state determined by  $d_a$ .

The initial physical state vector  $|\psi_{p_0}\rangle$  can be freely chosen. In this work, we make use of two different initial state vectors:  $|\psi_{p_0}\rangle = |\dots, \uparrow, \downarrow, \uparrow, \downarrow, \dots\rangle$  (arranged in a checkerboard pattern, shown as a cartoon on the left side of the top panel of Fig. 2 and others) and  $|\psi_{p_0}\rangle = |\dots, \uparrow, \uparrow, \uparrow, \uparrow, \dots\rangle$ ,

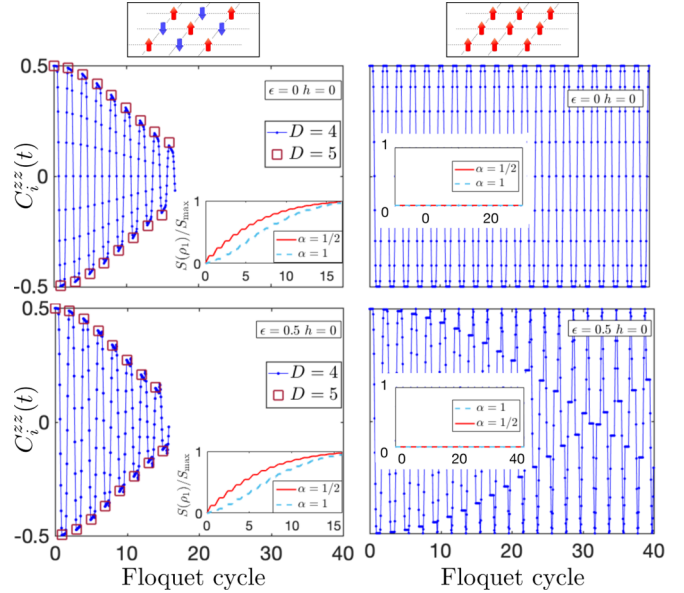


FIG. 2. Equal space correlator evaluated at different times for coupled spins without disorder starting from the Néel state (left) and the polarized state (right) for the perfect (top) and imperfect flips (bottom). Time crystalline behavior cannot survive for the Néel state due to thermalization, while there is no dynamics for the initial polarized state. Also shown are simulations with  $D = 5$  for stroboscopic times. Insets: growth of local Rényi entropies ( $\alpha = 1/2, 1$ ) for the blue and red curve, respectively.

referred to as the Néel and spin-polarized state vectors, respectively, in the following. These initial states are readily experimentally accessible and can be viewed as two limits of the entire state space of random product states. The dynamics of a given initial random product state, e.g., with respect to its decay (of the expectation values of the observables), will behave somewhere “in between” those two extreme configurations. Another reason for choosing these initial states is that because we are directly targeting the thermodynamic limit, we need to assume some translational invariance with a fixed unit cell. In our setting, we work with a two-site unit cell arranged in a checkerboard pattern (shown in Fig. 6). The two initial states can be written exactly as an iPEPS with bond dimension 1.

The dynamics generated by Hamiltonian (3) including the *disorder average* can be implemented by introducing  $S^z S^z$  coupling in the Hamiltonian between the physical and the auxiliary sites. This term ensures that the exact disorder average is recovered for all times and all disorder distributions. The modified form of (3) then takes the form

$$H_{\text{MBL}} = J \sum_{(i,j)} (S_{i_p}^x S_{j_p}^x + S_{i_p}^y S_{j_p}^y + S_{i_p}^z S_{j_p}^z) + h \sum_i S_{i_p}^z S_{i_a}^z, \quad (4)$$

where the subscripts  $p$  and  $a$  are used to denote the physical and the auxiliary sites. The disorder average of all possible configurations is already taken in one simulation while computing the expectation values of the observables (discussed in more detail in the Appendix A 3 of the paper). This is another advantage of our setting because we do not need to take multiple shots of disorder configuration and average them.

To monitor the time crystalline behavior, we will extend the notion of *long-ranged order* captured by order parameters from *equal-time correlations in space* to general correlations in space and time,

$$C_{i,i'}^{OO}(t,t') = \lim_{|L| \rightarrow \infty} \langle O(i,t)O(i',t') \rangle \neq 0, \quad (5)$$

where we put specific emphasis on *equal-space correlations in time*, abbreviated as

$$C_i^{OO}(t,t') = \lim_{|L| \rightarrow \infty} \langle O(i,t)O(i,t') \rangle = f(t,t'), \quad (6)$$

for times  $t, t' \geq 0$ . For our purposes, we call a system a time-crystal if, robust to perturbations,  $\epsilon \neq 0$  (to avoid trivial situations such as that considered in Ref. [4] being called a time crystal),  $f(t,t')$  shows a nontrivial (ordered) long time  $t \gg t'$  behavior that breaks the discrete time-translation symmetry of the underlying drive (here by period doubling or more generally taking integer values).<sup>1</sup> We will concentrate on  $O = S_i^z$  and  $t' = 0$  and write the corresponding correlation function as  $C_i^{S^z S^z}(t,0) = C_i^{zz}(t)$  in shorthand notation. The expectation values of these correlators are computed using an instance of a corner transfer matrix renormalization group (CTMRG) [67–70] (see also the Appendix A 3 for the techniques being used).

In our subsequent detailed discussion, we specifically consider the following different cases that seem particularly important and insightful: These are (i) uncoupled spins without disorder for  $J = 0, h = 0$ . Then, (ii) coupled spins without disorder,  $J = 1, h = 0$ . Finally, (iii) we investigate coupled spins with strong disorder,  $J = 1, h = 100$  for different levels of disorder, i.e.,  $d_a = 2$  and  $5$ , where  $d_a$  corresponds to the size of the local Hilbert space of the auxiliary systems.

### III. RESULTS

In all the cases discussed above, we choose  $T = 0.1$ , where  $T$  is the periodicity of the Floquet cycle. We consider a perfect flip  $\epsilon = 0$  and an imperfect flip  $\epsilon = 0.5$  in order to check the robustness of the time crystal in all the cases. To make sure that our results are not an artefact of the finite bond dimension, we make sure that the expectation values are consistent for the two best available bond dimensions we can afford, i.e.,  $D = 4, 5$ , as our stopping criterion in time, identifying the time when the iPEPS ansatz can no longer faithfully accommodate the entanglement growth (discussed in more detail in the Figs. 7–9 of the Appendix). We largely focus on these cases in order to understand the role of the various parts of the Hamiltonian and to establish which parameters of the Hamiltonian need to be tuned in actual quantum simulators in order to stabilize or realize the time crystal.

Case (i): Uncoupled spins ( $J = 0, h = 0$ ). This is the case when only the second part of the Hamiltonian (1) acts on the spins. Clearly, the spins stay uncoupled. While the system exhibits signatures such as period doubling as well as stability

<sup>1</sup>It is interesting to note that a *Magnus expansion* in terms of the periodicity  $T$  will lead to a perturbative series of Hamiltonians, each of which is captured by the no-go-theorem of Refs. [4,5], so that in such a description, all orders must be considered.

to infinite times, it cannot be called a time crystal because it is extremely sensitive to perturbations and is therefore not a well-defined phase of matter. This can be tested by introducing an imperfection in the spin flip denoted by  $\epsilon$ . This is shown in Fig. 1 for  $\epsilon = 0.5$ . Note that this regime is reminiscent of the setting of an equal-position correlation function in independent two-level systems spreading over space as considered in Ref. [4] as rather trivial and not time-crystalline behavior.

We plot the equal space correlator defined in Eq. (6) as a function of the Floquet cycle, and from this we can see the beating pattern emerging and the knot of the beat at around the Floquet cycle  $32$ . We also notice the spikes in  $C_i^{zz}(t)$  due to the effect of nonzero  $\epsilon$ . We resort to looking at identifying signatures of time crystal from the short time dynamics as different values of  $\epsilon$  would simply mean a change in the position of such knots along the  $X$  axis. In other words, for  $\epsilon = 0$ , the knot would completely disappear along with the spikes seen here.

Case (ii): Interacting spins in the absence of disorder ( $J = 1, h = 0$ ). This is the case when both parts of the Hamiltonian in (1) are active, however  $h = 0$  for  $H_{\text{MBL}}$ . We will show the plots for  $\epsilon = 0, 0.5$  starting from the Néel states as well as the polarized states. In the presence of many-body interactions without any disorder, our system is expected to thermalize very quickly [7,9]. An instance of a time evolution compatible with this expectation is shown by plotting the equal space correlator in time in the left panels of Fig. 2, where we are able to show time evolution only up to approximately 16 Floquet cycles. It is also evident from the growth of local Rényi entropies for one-site reduced quantum states shown in the insets of the left panels. The Rényi entropies ( $\alpha = 1/2, 1$ ) are rescaled to their maximal value of  $\ln(d_p)$ . The values of  $\alpha$  are chosen to provide insight into how well one would expect a low bond dimension tensor to approximate the quantum state at hand [71].

The right panels of this figure correspond to starting from a polarized initial state, and, other than the periodic flipping of the spins induced by  $H_S$ , there are no dynamics whatsoever. This is also revealed by the local Rényi entropies, which remain zero throughout the evolution (shown in the insets). This can be easily understood by the spin  $SU(2)$  symmetry of the Hamiltonian  $H_{\text{MBL}}$  at  $h = 0$ , which entails that even at nonzero  $\epsilon$ , the time evolution rotates all spins in the same manner during the application of  $H_S$ , while the spins remain inert during  $H_{\text{MBL}}$ . Ultimately, the fully spin-polarized state remains fully polarized (albeit with respect to a different polarization axis at  $\epsilon \neq 0$ ) after each flip, and the dynamics are the same as that of a single spin (compare Fig. 1).

Even in such a short timescale available (for the Néel initial state), one can still see important features such as period doubling and the absence of spikes due to finite  $\epsilon$ , as seen in Fig. 1. One can then speculate that the destruction of the time crystal behavior in this case is ultimately caused by thermalization for long times. We then resort to overcoming this effect by adding strong disorder  $h = 100$  to our system for different levels of disorder  $d_A = 2$  and  $5$ .

Case (iii): Interacting spins with strong disorder ( $J = 1, h = 100$ ). This is the case when both parts of the Hamiltonian are active and also the  $H_{\text{MBL}}$  features strong disorder. We start our discussion by fixing the number of levels of disorder to 2.

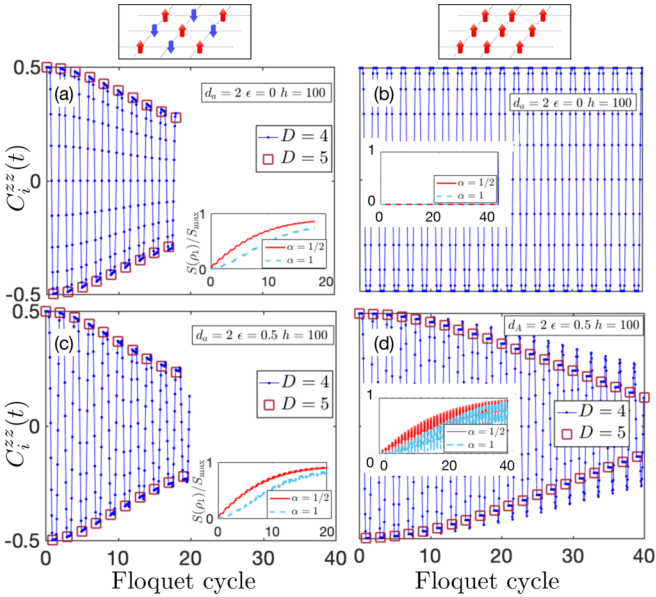


FIG. 3. Equal space correlator evaluated at different times for coupled spins with strong disorder of two levels for different initial states and  $\epsilon$ . Also shown are the simulations with  $D = 5$  for stroboscopic times. Insets: growth of local Rényi entropies ( $\alpha = 1/2, 1$ ) for the blue and red curve, respectively.

This means the disordered field can take only two different values. We see that, compared to case (ii), we have been able to delay thermalization only slightly for both  $\epsilon = 0$  and  $0.5$  for the Néel initial states (Fig. 3). For these cases, we see that despite a strongly disordered field, the two levels are not enough to sufficiently stabilize a time crystal. This is consistent with our previous findings on many-body localization in two spatial dimensions [60], where sufficient levels of disorder were required to achieve localization. However, if

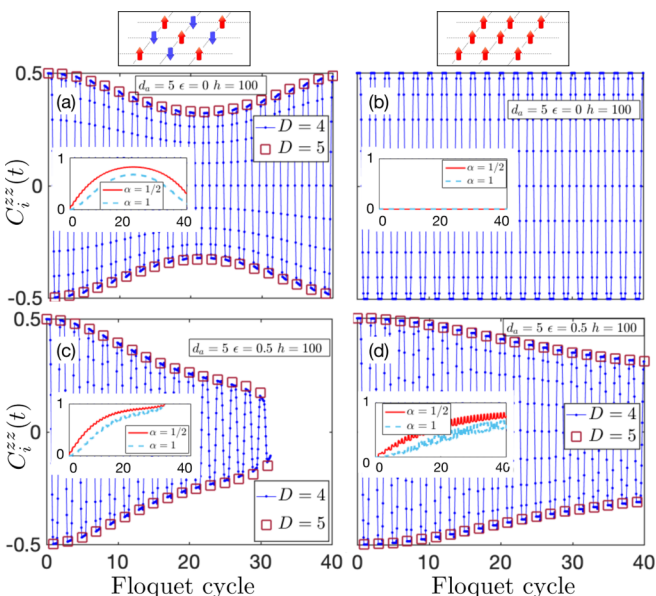


FIG. 4. Same as Fig. 3, but with five levels of disorder. The time crystal has stabilized due to the increasing number of disorder levels, thereby surviving much longer times for both of the initial states.

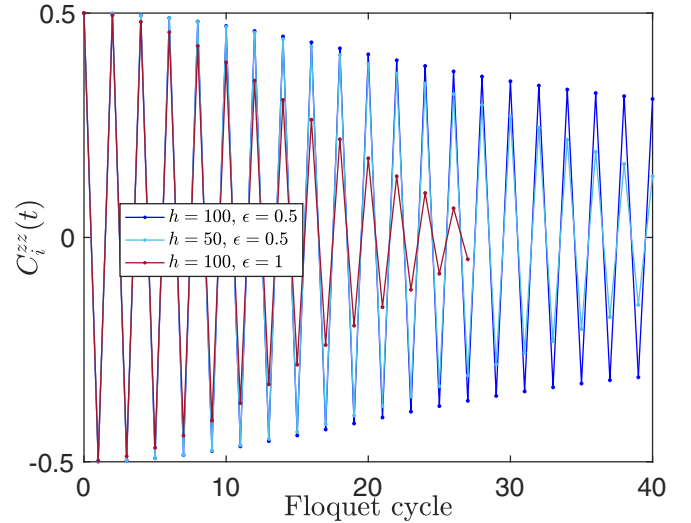


FIG. 5. Stability of the time crystal with respect to different disorder strength  $h$  and perturbation to the perfect flip  $\epsilon$ . The results are plotted only for stroboscopic times.

we start from the polarized state, we notice that the dynamics slow down considerably for the case of  $\epsilon = 0.5$  while there are none in the case of  $\epsilon = 0$ . The local Rényi entropies shown in the insets provide further substance to these results.

In a next step, we now increase the number of levels of disorder by taking  $d_a = 5$  such that our disordered field can take five different values. In all the cases, we see that the stability of the time crystal improves remarkably, thereby allowing us to go much longer times even beyond the currently shown 40 Floquet cycle.

The delay of thermalization is also consistent with the growth of local Rényi entropies (shown in the insets). Thus, we have managed to stabilize our two-dimensional time crystal (i.e., increase its lifespan) by increasing the number of levels of disorder at sufficiently large strength. We have also seen that the stability also depends on the initial state chosen, the one starting from the polarized state being the most favorable. It is worth pointing out that a resurgence is observed in the top left panel of Fig. 4 due to the strong disorder and the perfect flip at  $\epsilon = 0$ . In Fig. 5, we show the stability of our time crystal starting from the polarized state with respect to a smaller disorder ( $h = 50$ ) and a larger perturbation to the perfect flip ( $\epsilon = 1$ ). As expected, smaller disorder strength leads to a decrease in the lifespan of the time crystal. Unlike the case in Ref. [60], where only a modest disorder strength of  $h = 4$  (with at least four levels of disorder) was enough to localize our system, larger disorder strength is needed here in order to counter the extra heating effect of the drive. Similarly, a large value of  $\epsilon$  will also lead to the ultimate destruction of the time crystal [16]. This is also shown in Fig. 5. This behavior is consistent with the findings of Ref. [16].

#### IV. IMPLEMENTATIONS IN PROGRAMMABLE QUANTUM SIMULATORS

It is key to the setting laid out here that it is amenable to quantum simulation using programmable disorder.

Dichotomic disorder seems incapable of sufficiently stabilizing the time crystal, so that more intricate disorder is required. Superconducting systems seem to be specifically suitable for this kind of programmable disorder [30,36]. In such systems, two spatial dimensions with programmable disorder seem to be within reach, and also discrete disorder can be largely programmed. Other examples include individually controlled *Rydberg atoms* [35] and systems of *cold atoms in optical lattices* [32], in which the disorder is realized by imaging a two-dimensional random disorder potential to a single atomic plane in an optical lattice, controlled by a *digital mirror device* allowing for programming of discrete disorder, or possibly also by making use of another atomic species acting as a disorder potential to the other. Similarly, two-dimensional arrays of *trapped ions* [34] have programming capabilities of the kind required here.

The situation discussed here gives rise to a highly interesting state of affairs, linking notions of quantum many-body and quantum information theory, to make a point mentioned in the Introduction more precise and specific: For short times, state-of-the-art tensor network methods provide a reliable machinery to assess the prethermal time crystalline behavior. Based on such classically available data, and pushing the classically accessible regime with substantial effort, one can judge what phases of matter can reasonably be expected, with substantial predictive power.

Still, due to the linear growth of the entanglement entropy of subsystems in time [72], intermediate and long times will remain inaccessible using tensor network methods, even if folding methods [73] or the exploitation of mode transformations [74]—ideas that have not yet been developed for two spatial dimensions—may ultimately render data on somewhat longer times classically available.

This insight gives substantial impetus to the field of quantum simulation. It is perfectly compatible with complexity theoretic insights. When appropriately cast into the form of a decision problem, the problem of simulating the time evolution under general translationally invariant local Hamiltonians is in the complexity class BQP, the class of problems that can be solved by a quantum computer to bounded error in polynomial time. Interestingly, this problem is even complete for BQP [29], which implies that it is unreasonable to expect that there exists an efficient classical algorithm for the problem, as this would imply that  $BPP = BQP$ , which would in turn result in polynomial time classical algorithms for factoring, which is considered excessively implausible. Of course, physically “natural” problems may occur in principle outside the classically hard instances, but in light of these complexity theoretic obstructions, it is still highly unlikely that a classical efficient simulation method for interacting quantum many-body systems can ever be identified.

This comes along with advice concerning the necessity of implementing quantum simulators: Indeed, one needs programmable quantum simulators [21,30–36] to ultimately assess the stability of such phases of matter for long times [1,2]. This constitutes a most interesting situation and at the same time a valid and meaningful application for programmable quantum simulators as intermediate quantum technological devices between full quantum computers and analog quantum simulators: Here, the long-time behavior

only accessible for quantum but not classical devices is at the heart of the matter when discussing time-crystalline behavior.

## V. CONCLUSION AND OUTLOOK

In this work, we have provided fresh evidence of (prethermal) quantum time crystals in two spatial dimensions, using the infinite version of projected entangled pair states (iPEPS) that is able to address the thermodynamic limit directly. We have combined a tensor network machinery that includes a quantum dilation approach to capture strong disorder with a suitable stroboscopic Floquet Hamiltonian evolution that features a discrete time translation symmetry. Starting from different initial states, we have clearly encountered the breaking of time translation symmetry revealed by the equal space correlator in time that is robust to perturbations in the spin-flip Hamiltonian. The stability of such quantum time crystals can be increased by taking a sufficiently strong disorder featuring a large number of levels of disorder and suitable initial state. In future work, we will explore whether dissipation can be properly engineered to stabilize such phases of matter by avoiding thermalization through the use of *projected entangled pair operators* [55].

In addition to pushing the machinery of tensor networks to a new regime, it is our hope that this work provides significant further guidance for the use of programmable quantum simulators to explore intricate nonequilibrium quantum phases of matter, and ultimately to discriminate prethermal from genuine time crystals [2] as they cannot be classically discriminated. Quantum simulators will also help to answer more intricate questions such as that regarding broken translation symmetry in space as well as analyzing  $C_{i,i'}^{O,O}(t,t')$  at  $t' \neq 0$ , which is more challenging to compute in our classical simulation approach. This could allow us to obtain deeper insights into the mechanisms of spontaneous symmetry breaking in space and time [75].

Ultimately, phases such as discrete time crystals may have an impact beyond academic interests to devising technological applications in, say, quantum metrology. The present scheme adds to the portfolio of schemes for programmable quantum simulators to be explored that are feasible instances of quantum devices intermediate between analog quantum simulators and full-scale quantum computers. The present work, so we hope, also helps to delineate the delicate boundary between classically simulatable regimes of quantum many-body physics and those that ultimately have to be assessed with quantum devices.

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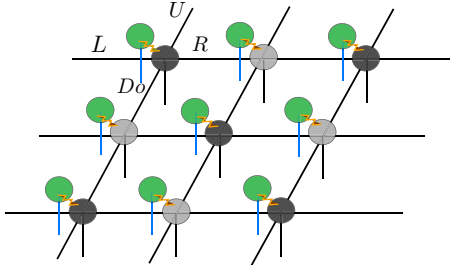


FIG. 6. Implementing disorder in a translationally invariant setting. We choose our initial global state to be a tensor product of the physical state (represented by gray tensors) and the auxiliary state (represented by green tensors).  $L$ ,  $U$ ,  $R$ , and  $Do$  correspond to the different links we need to update for a particular site.

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## APPENDIX

We make use of several tensor network methods here, developed to be applicable in new regimes, in particular a method rooted in *infinite projected entangled pair states* (iPEPS) [37,39,43], here pushed to the fresh regime of interactions, disorder, and time-dependent drive being present simultaneously. We implement disorder in such a translationally invariant setting using a quantum dilation technique [11]. We discuss briefly below the different numerical methods used and developed in this work.

### 1. Implementing disorder in a translationally invariant system

This approach was originally introduced in Ref. [11] for studying disordered infinite-sized one-dimensional systems [76,77]. Such ideas have only very recently been implemented in two-dimensional systems using iPEPS [58,60]. The procedure starts by appending an auxiliary site to each physical one in the system. The global initial state is thus a product state of the physical system and the auxiliary system represented by a state vector  $|\Psi_0\rangle = |\psi_{p_0}\rangle \otimes |\psi_{a_0}\rangle$ . The auxiliary state is fixed and is a tensor product of equal superposition states at each site as defined in the main text. The physical state depends on our choice and is also discussed in the main text. The original part of the Hamiltonian in Eq. (3) is then modified to the form in Eq. (4), which includes the classical interaction  $S_{i_p}^z S_{j_p}^z$  between the physical sites and the auxiliary sites. This is responsible for injecting the disorder into the physical system. This is shown in Fig. 6. We then perform a quench of the global initial state with the Floquet Hamiltonian defined in Eq. (1) to arrive at a time-dependent state vector

$$|\Psi(t)\rangle = e^{-iH_F t} |\Psi_0\rangle. \quad (\text{A1})$$

We note here that the different parts of the Hamiltonian act at different times of the evolution.

### 2. Time evolution

In this work, we exploit a variant of *time evolving block decimation* (TEBD) [78] in two spatial dimensions. We have specifically implemented an iPEPS algorithm with a two-site unit cell arranged in a checkerboard pattern. Such a configuration is enough to realize our two initial states discussed in the main text. The tensors are then optimized using the so-called *simple update* [79]. This scheme is more efficient and has also been found to be more accurate and stable compared to the *full update* procedure for real time evolution [57,58,60]. To describe this approach briefly, consider our two-dimensional Hamiltonian defined on a square lattice as in the setting used here. We split our Hamiltonian into different parts corresponding to the different links of the lattice, i.e.,

$$H = H^L + H^U + H^R + H^{Do}, \quad (\text{A2})$$

where

$$H^\alpha = \sum_{(i,j)} h_{i,j}^\alpha \quad (\text{A3})$$

for  $\alpha = L, U, R, Do$  correspond to the left, up, right, and down link of the square lattice. We then define the following two-body gate operators:  $e^{-i\delta t H} \approx e^{-i\delta t H^L} e^{-i\delta t H^U} e^{-i\delta t H^R} e^{-i\delta t H^{Do}}$ . After updating each link with the appropriate term, the bond dimension along that link is truncated. The procedure is performed while assuming a mean-field-like environment of the link. We use a first-order Trotter step of  $\delta t = 0.005$ .

We have performed such an optimization procedure with an iPEPS bond dimension of  $D = 4, 5$ . The large physical dimension of the local Hilbert space ( $d = d_p d_a = 4, 10$ ) restricts us from accessing very large bond dimensions  $D$  of the iPEPS. This also limits the timescales that we can access using tensor network approaches (reflecting the observation that time evolution under local Hamiltonians is a computationally hard problem in worst-case complexity). The results shown above are nonetheless converged and consistent with

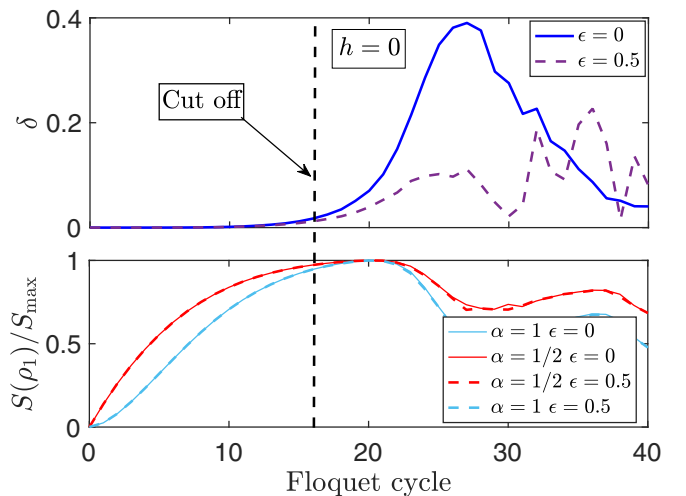


FIG. 7. Errors corresponding to Fig. 2 (left panels). Top:  $\delta \geq 0$  is the absolute value of the difference of the expectation values of the correlators corresponding to simulations with  $D = 4$  and 5. Bottom: growth of local Rényi entropies for one-site reduced density matrix for the full time evolution.

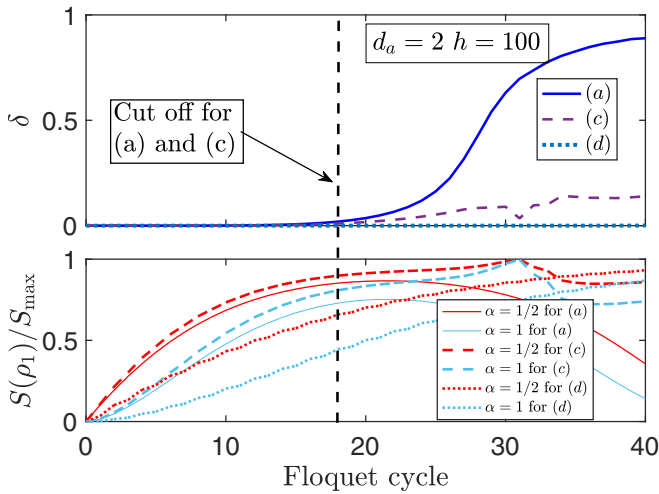


FIG. 8. Errors corresponding to Fig. 3. Top:  $\delta$  is the absolute value of the difference of the expectation values of the correlators corresponding to simulations with  $D = 4$  and 5. Bottom: growth of local Rényi entropies for one-site reduced density matrix for the full time evolution. (a), (c), and (d) refer to the different subplots of Fig. 3 in Sec. III.

the largest available bond dimensions used. Such an agreement between the highest available bond dimensions can be used to approximately determine the stopping criteria of our time evolution.

### 3. Disorder-averaged expectation values

Once the tensors have been obtained using the above procedure, we resort to computing the expectation values of the physical observables of interest. This involves contracting the entire two-dimensional tensor network in the thermodynamic limit. Unlike the situation for one-dimensional matrix product states, this step cannot be done efficiently and is known to be a computationally hard problem [80] for exact contraction even in average-case complexity [81]. Luckily, there are several approximation schemes available for this [43,67,69,82–84] (and there is good evidence that the contraction of PEPS is computationally feasible to a good approximation for physically meaningful PEPS describing gapped models [85]). The expectation values of the observables computed here are already the disorder-averaged expectation values of all the possible disorder configurations. This is easy to see from the expression

$$\begin{aligned} \langle\langle \hat{O}(t) \rangle\rangle &= \langle \Psi_0 | e^{iHt} \hat{O} e^{-iHt} | \Psi_0 \rangle \\ &= \langle {}_a \langle \psi_0 | \otimes {}_p \langle \psi_0 | e^{iHt} \hat{O} e^{-iHt} (| \psi_0 \rangle_p \otimes | \psi_0 \rangle_a). \end{aligned} \quad (\text{A4})$$

Thus, one of the advantages of our setting is having to avoid taking multiple shots of disorder and taking their averages. The programmable nature of the disorder employed here also

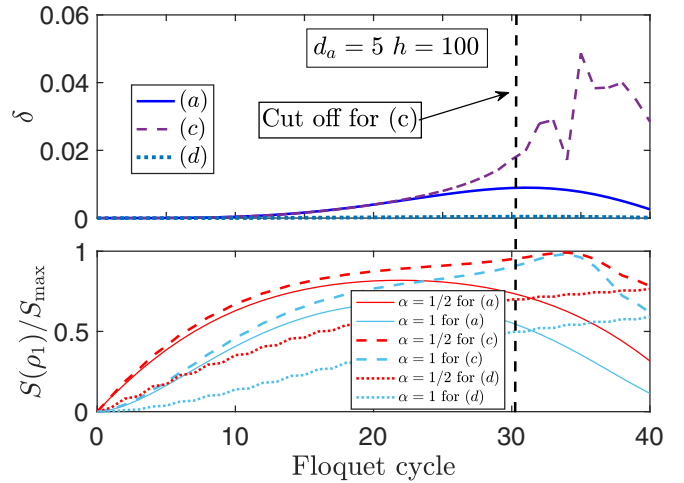


FIG. 9. Errors corresponding to Fig. 4. Top:  $\delta$  is the absolute value of the difference of the expectation values of the correlators corresponding to simulations with  $D = 4$  and 5. Bottom: growth of local Rényi entropies for one-site reduced density matrix for the full time evolution. (a), (c), and (d) refer to the different subplots of Fig. 4 in Sec. III.

has direct control over avoiding rare ergodic regions, which can lead to instability of MBL at long times [61,62].

In this work, we make use of the *corner transfer matrix renormalization group* method (CTMRG) [67–70] to approximate contraction, here freshly applied to classically keeping track of quantum Floquet dynamics. This involves computing a set of tensors known as the *environment* tensors approximating the fixed points of the “corners” of an infinite two-dimensional lattice. This infinite two-dimensional lattice to be contracted is actually composed of the double-layer norm tensors of bond dimension  $D^2$ . We use an environment bond dimension of up to  $\chi = D^2$  in our simulations, which is found to be sufficient for our purposes.

In Fig. 7 (top panel), we plot  $\delta \geq 0$ , defined as the absolute value of the difference between the expectation values of the correlators for bond dimensions  $D = 4$  and 5. In the bottom panel, we show the growth of local Rényi entropies for the extended time evolution. The time at which  $\delta$  becomes significant is used as the “cutoff” time for Fig. 2 (left panels). Incidentally, the local Rényi entropies start saturating to their maximal value near this point. For the right panels of Fig. 2, there is no dynamics and therefore no concomitant errors in the growth of entanglement. Similar error measures corresponding to Figs. 3 and 4 are also shown in Figs. 8 and 9, respectively. We see how the time evolution has been extended significantly depending on the choice of disorder levels and the initial states, directly corresponding to the stability of the time crystal.

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