

Are circadian amplitudes and periods correlated?

A new *twist* in the story

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Supplementary Material

Table S1. Summary of parametric twist effects upon changes of model parameters in the Almeida model. Model parameters were changed individually around $\pm 20\%$ their default parameter value (see Table 2 in main text) to simulate oscillator heterogeneity and study amplitude-period correlations from all variables of the ensemble. Due to the synergies of feedback loops present in the Almeida model [1], different parametric twist effects can appear for variations of one particular parameter depending on the measured variable. If the default parameter variation resulted in an amplitude-period correlation where the range of ratio of amplitude variation (compared to the default amplitude) was < 0.1 , that ensemble was considered to have no twist (twist = 0) for that particular control parameter. + and – signs refer to the sign of the correlations.

effect on:	$\pm 10\%$ parameter change in											
	V_R	k_R	k_{Rr}	V_E	V_D	γ_{Ror}	γ_{Rev}	γ_P	γ_C	γ_{PC}	γ_{CP}	γ_{BP}
BMAL1	+	+	+	–	–	0	0	+	+	+	0	0
ROR	0	0	–	0	0	0	–	0	0	0	0	0
REV	+	+	0	+	0	0	+	+	+	0	0	0
DBP	+	+	+	0	0	+	+	+	0	+	0	0
E4BP4	0	0	0	0	+	+	0	0	0	0	0	0
CRY	0	0	0	0	0	0	0	0	0	0	0	0
PER	+	+	+	+	0	+	+	+	+	+	0	0
PERCRY	+	+	+	+	0	+	+	+	+	+	0	0

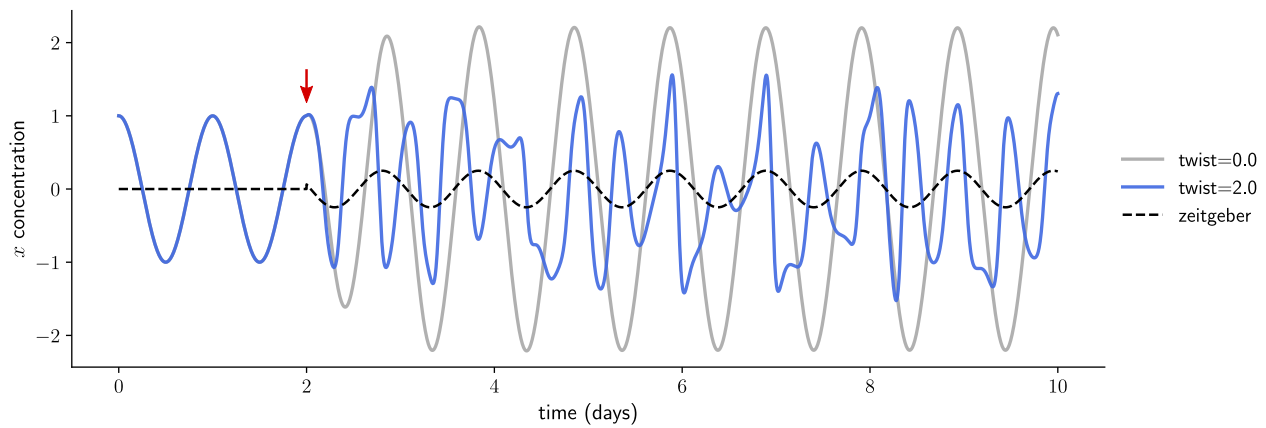


Figure S1. Large values of oscillator twist ϵ affect sync of individual oscillators to external zeitgeber inputs. An oscillator with no twist (shown in grey) entrains to the zeitgeber (indicated with a dashed line) when the periodic input is turned on (red arrow, $T = 24.5$ h, $F = 0.25$), but the oscillator with large positive twist (blue) does not entrain and loses its periodicity. Both oscillators, besides their twist value, are otherwise identical (same free running period $\tau = 24$ h, amplitude $A = 1$ and amplitude relaxation rate $\lambda = 0.05$ h⁻¹).

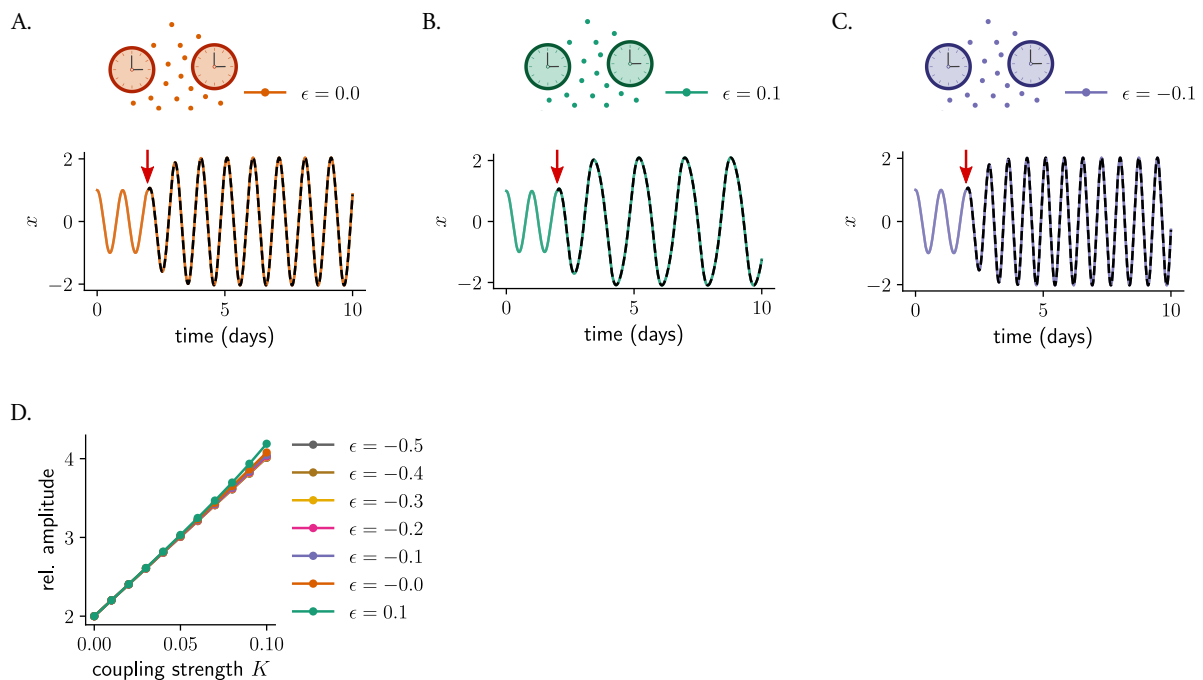


Figure S2. Coupling induces amplitude expansions and period changes that depend on the twist ϵ . Effect of mean-field coupling on two oscillators with (A) $\epsilon = 0$, (B) $\epsilon = 0.1$ or (C) $\epsilon = -0.1$ but otherwise identical (free running period $\tau = 24$ h, amplitude $A = 1$ and amplitude relaxation rate $\lambda = 0.05$ h⁻¹). The time series show how the period of the coupled network depends on the twist parameter ϵ . The moment in which mean-field coupling is turned on, at a coupling strength $K = 0.1$, is shown with a red arrow; the mean-field is shown with a dashed black line. (D) The amplitude of the coupled network increases with coupling strength K , but with no major differences for networks with different twist values.

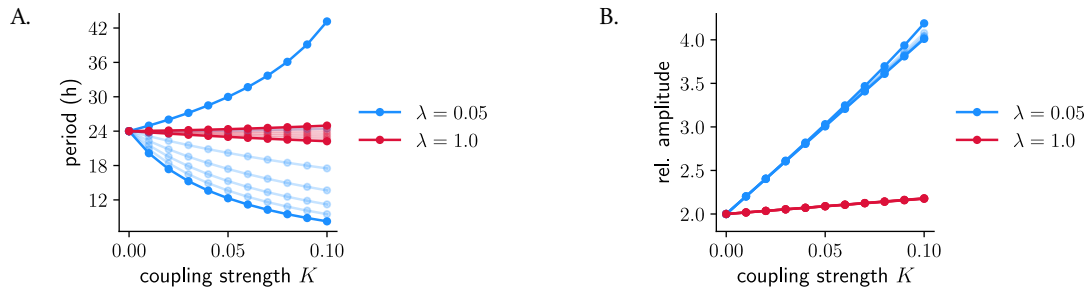


Figure S3. Role of amplitude relaxation rate on coupling-induced period and amplitude changes. More rigid clocks (higher λ values, red lines) are less sensitive to twist-induced effects and thus display less coupling-induced period changes (A) or amplitude expansions (B) than weaker clocks (in blue).

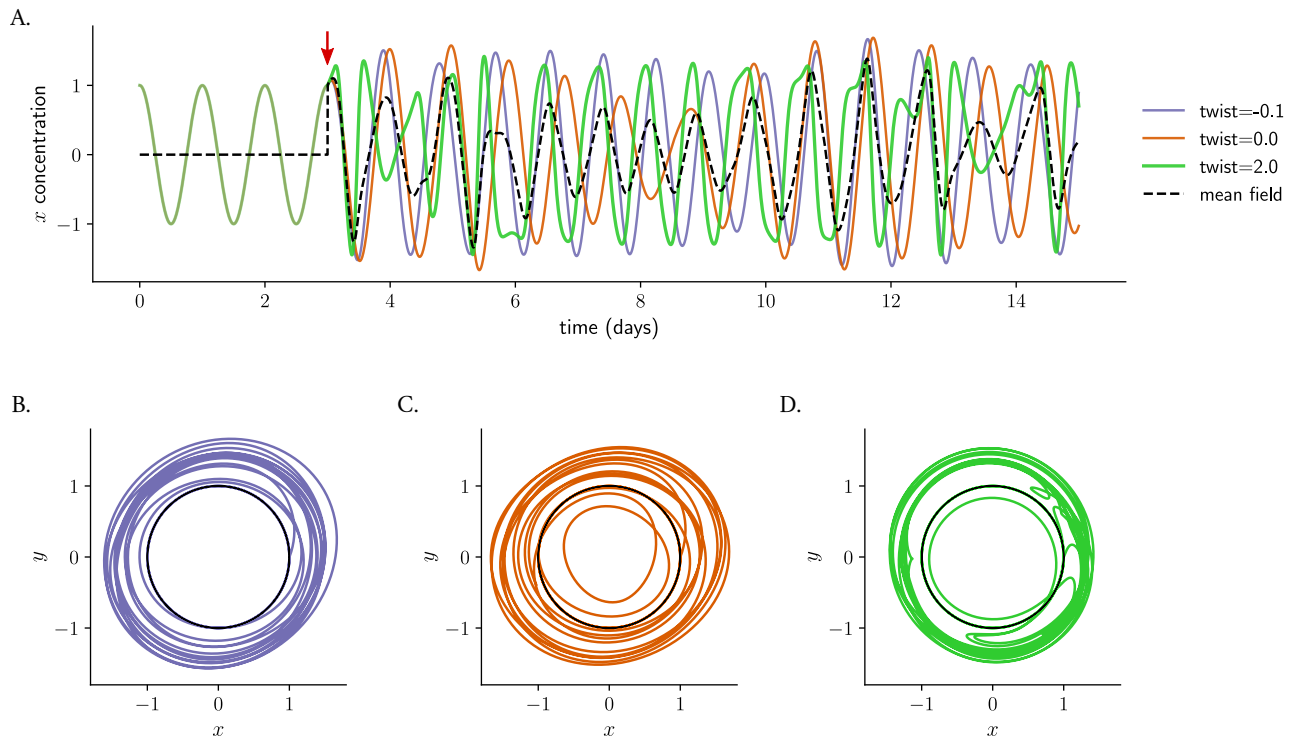


Figure S4. Twist-induced chaos upon mean-field coupling. (A) Oscillators with $\epsilon = -0.1, 0, 2$ oscillate in a self-sustained manner in the absence of coupling, but when mean-field coupling at $K = 0.1$ is turned on (red arrow), periodicity is lost. The oscillator with $\epsilon = 2$ (green line) cannot synchronize due to its large twist and thus affects the rest of the network (purple and orange lines become arrhythmic too). As a result, the mean-field (shown in black and with dashed line) also becomes arrhythmic. (B–D) Twist-induced chaotic trajectories of the oscillators from (A) shown in phase space. The limit cycles from the uncoupled clocks are shown in black.

References

- [1] Almeida, S., Chaves, M. & Delaunay, F. Transcription-based circadian mechanism controls the duration of molecular clock states in response to signaling inputs. *J Theor Biol* **484**, 110015 (2020).