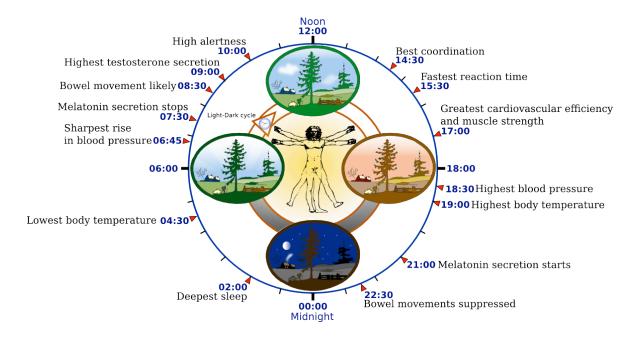
Simulation of the KaiABC Circadian Oscillation by Matlab

Xuan Zhu, BCMB June 3, 2009

Circadian Rhythm



- A roughly-24-hour cycle in the biochemical, physiological or behavior processes of living entities, including plants, animals, fungi and **cyanobacteria**
- Endogenously generated
- Can be entrained by **external cues** (called Zeitgebers), such as daylight
- Allow organisms to anticipate and prepare for precise and regular environmental changes

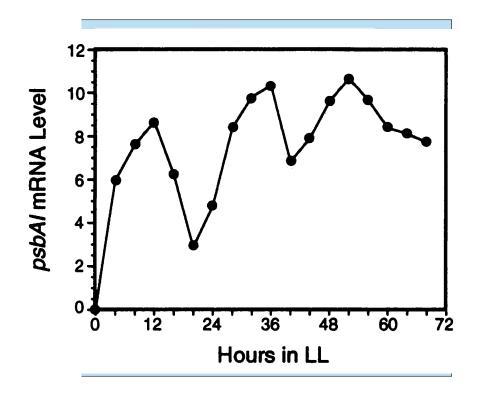
Three General Criteria

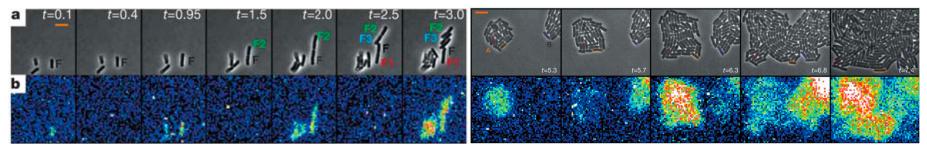
- 1. The rhythms persist in the absence of cues.
- 2. They persist equally precisely over a range of temperatures (i.e. temperature-compensation).
- 3. The rhythms can be adjusted to match the local time.

- Are prokaryotes capable of circadian rhythmicity?
- "Why have a timer for a cycle that is longer than your life time?"

Bacterial Circadian Rhythms

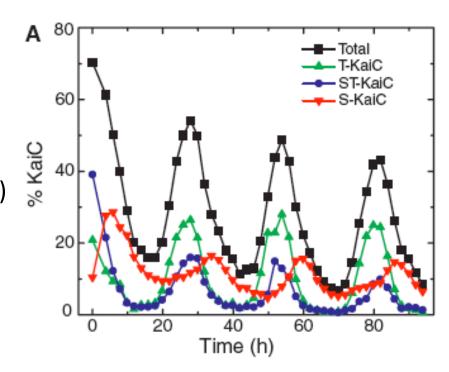
- Cyanobacteria display daily rhythms of nitrogen fixation in both LD cycle and in constant light (1985-1986)
- Satisfy the three criteria
- Keep track of two timing processes
- Adaptive significance





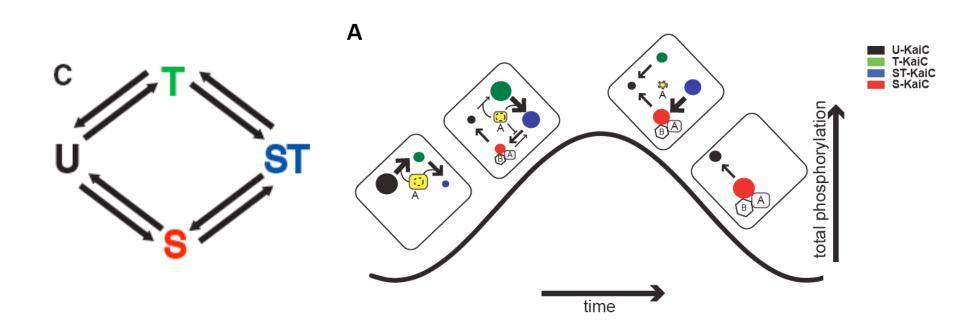
Molecular Mechanism of the Cyanobacterial Clockwork - KaiABC

- Traditional view: transcriptional feedback oscillators
- Reconstitution in vitro using only KaiA, KaiB, and KaiC
- KaiC: a hexameric enzyme that can autophosphorylate (KaiA-dependent) and autodephosphorylate at both \$431 and T432
- KaiA: its dimer enhances the autophophorylation of KaiC
- KaiB: antagonizes the activity of KaiA



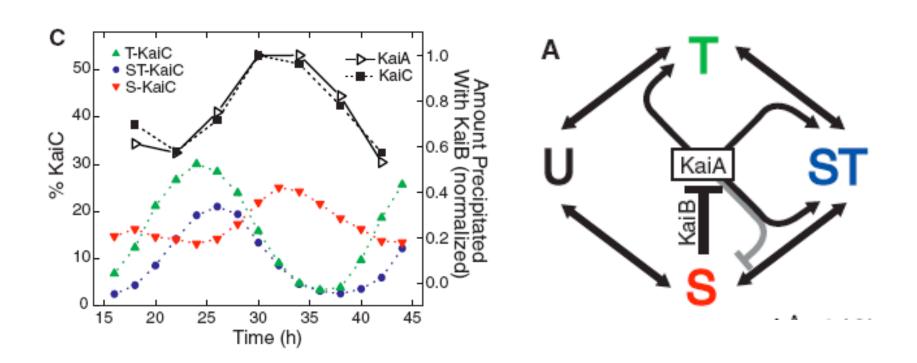
New model: The phosphoform distribution (or a combination tightly linked to the phosphorylation state) determines the phase of the oscillator.

Four-state model with first-order kinetics of interconversion of KaiC phosphoforms

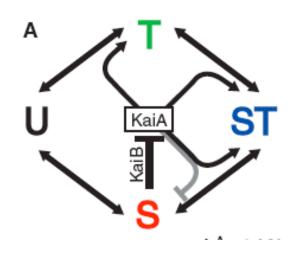


New model: The phosphoform distribution (or a combination tightly linked to the phosphorylation state) determines the phase of the oscillator.

- 1. KaiA activity alters the first-order rate constants for interconversion of KaiC phosphoforms
- 2. KaiB suppresses KaiA activity in an S-KaiC-dependent manner



Description of the Model



Assumptions:

 The concentrations of the three phosphorylated species are the only slow dynamical variables;

$$\frac{dT}{dt} = k_{UT}(S) U + k_{DT}(S) D - k_{TU}(S) T - k_{TD}(S) T \tag{1}$$

$$\frac{dT}{dt} = k_{UT}(S) U + k_{DT}(S) D - k_{TU}(S) T - k_{TD}(S) T$$

$$\frac{dD}{dt} = k_{TD}(S) T + k_{SD}(S) S - k_{DT}(S) D - k_{DS}(S) D$$

$$\frac{dS}{dt} = k_{US}(S) U + k_{DS}(S) D - k_{SU}(S) S - k_{SD}(S) S$$
(1)

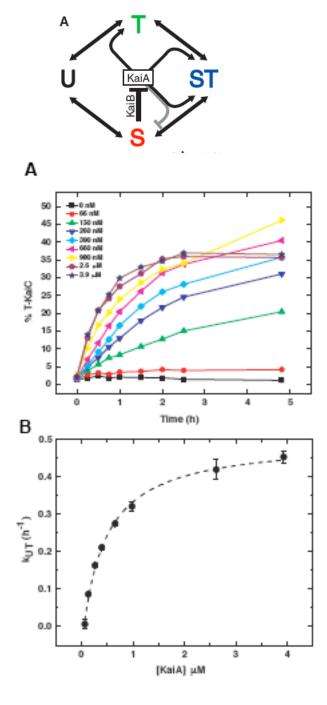
$$\frac{dS}{dt} = k_{US}(S) U + k_{DS}(S) D - k_{SU}(S) S - k_{SD}(S) S$$
 (3)

$$k_{XY}(S) = k_{XY}^0 + \frac{k_{XY}^A A(S)}{K_{1/2} + A(S)}$$

$$A = \max\{0, [KaiA] - 2mS\}$$

Assumptions:

- 1. The concentrations of the three phosphorylated species are the only slow dynamical variables;
- 2. The interconversions are first-order reactions with rates that dependent hyporbolically on the concentration of active KaiA;



Summary of the model

$$\frac{dT}{dt} = k_{UT}(S) U + k_{DT}(S) D - k_{TU}(S) T - k_{TD}(S) T$$

$$\frac{dD}{dt} = k_{TD}(S) T + k_{SD}(S) S - k_{DT}(S) D - k_{DS}(S) D$$

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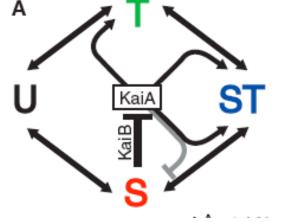






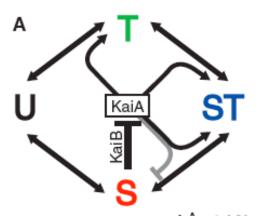






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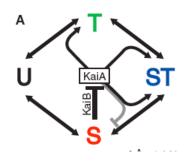
Assumptions:

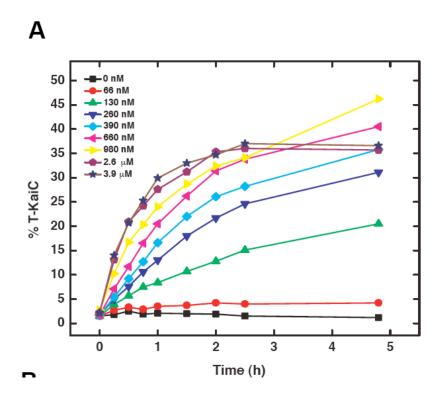
- 1. The concentrations of the three phosphorylated species are the only slow dynamical variables;
- 2. The interconversions are first-order reactions with rates that dependent hyporbolically on the concentration of active KaiA;
- Each S-KaiC monomer (together with KaiB) inactivates one KaiA dimer (m=1).

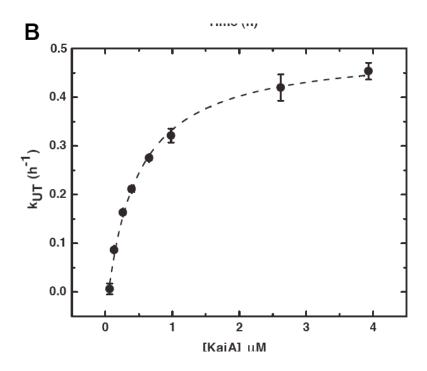
Determine $K_{1/2}$: $K_{1/2}=0.43\pm0.05\mu M$

$$k_{XY}(S) = k_{XY}^{0} + \frac{k_{XY}^{A}A(S)}{K_{1/2} + A(S)}$$

 $A = \max\{0, [\text{KaiA}] - 2mS\}$

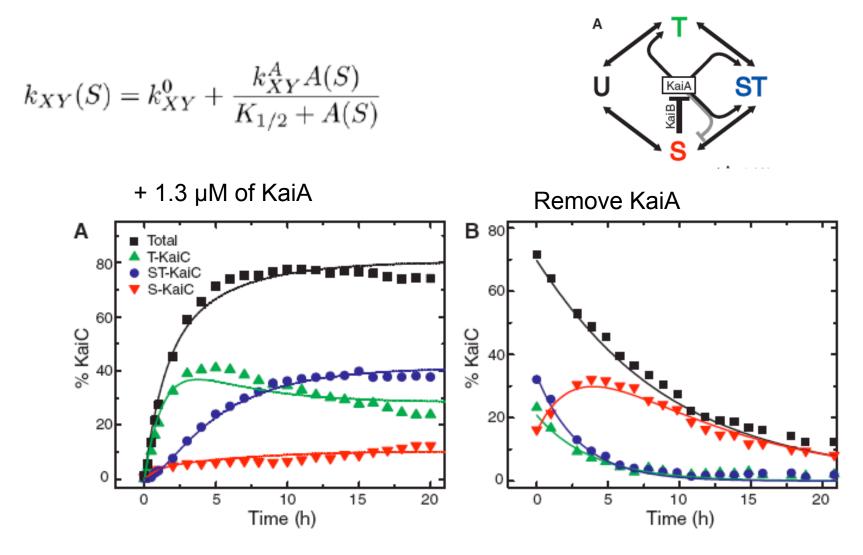






Rust, M.J., et. al. Science 2007.

Determine k⁰ and k^A:

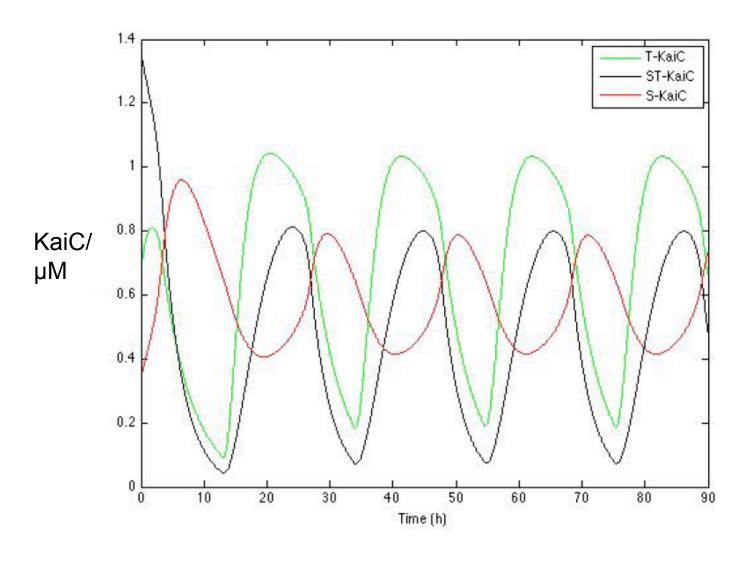


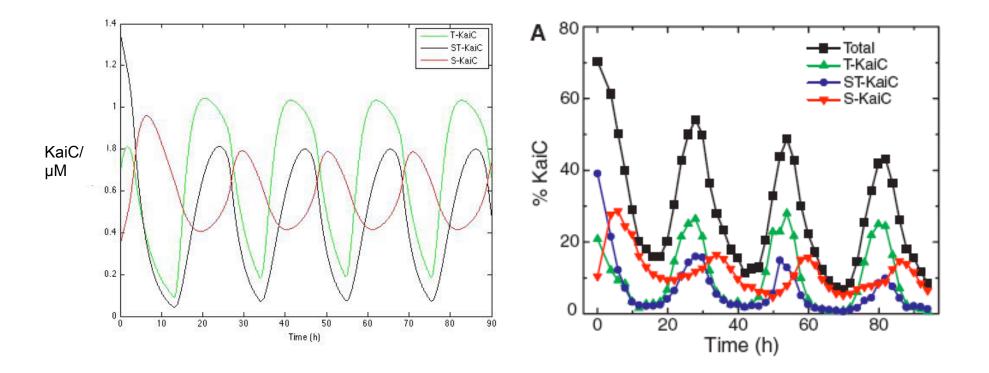
Rust, M.J., et. al. Science 2007.

Table S2

Category	Process	Parameter name	Value	Experiment
Basal rates (- Kai A)	U→T	k_{or}^{o}	0 h ₁₊	Figure 2B*
	T → ST	k_{ID}^{ϕ}	0 h ₁₊	Figure 2B*
	S → ST	k _{sp}	0 h ⁻¹ *	Figure 2B*
	U→S	k _{to}	0 h ⁻¹ *	Figure 2B*
	T→U	k_{ro}^{ϕ}	0.21 h ⁻¹	Figure 2B
8	ST → T	k_{Dr}^{0}	0 h ⁻¹	Figure 2B
Ã	ST → S	k ⁰ _{če}	0.31 h ⁻¹	Figure 2B
	S→U	k_{sv}^{o}	0.11 h ⁻¹	Figure 2B
	U→T	k_{or}^{A}	0.479077 h ⁻¹	Figure 2A
i A	T → ST	k_{TD}^A	0.212923 h ⁻¹	Figure 2A
of K	S → ST	k_{sp}^{A}	0.505692 h ⁻¹	Figure 2A
.8	U→S	k4	0.0532308 h ⁻¹	Figure 2A
Maximal Effect of Kai A	T→U	k_{TU}^A	0.0798462 h ⁻¹	Figure 2A
B.	ST → T	k_{DT}^{A}	0.1730000 h ⁻¹	Figure 2A
Max	ST → S	k-A ze	-0.319385 h ⁻¹	Figure 2A
_	S→U	k_{SU}^A	-0.133077 h ⁻¹	Figure 2A
Other	Concentration of KaiA causing half- maximal effect on KaiC	K _{1/2}	0.43 µМ	Figure S4
	Stoichiometry of inactivation of KaiA dimers by S-KaiC	т	1	assumed
	Concentration of KaiA	[KaiA]	1.3 μΜ	Bradford assay (see Materials and Methods)
	Concentration of KaiC	[KaiC]	3.4 μ M	Bradford assay (see Materials and Methods)
Simulation parameters for Figure 4A	Initial Conditions	T ₀	0.68 μМ	Figure 1A
		D_{b}	1.36 µM	Figure 1A
		S ₀	0.34 μΜ	Figure 1A

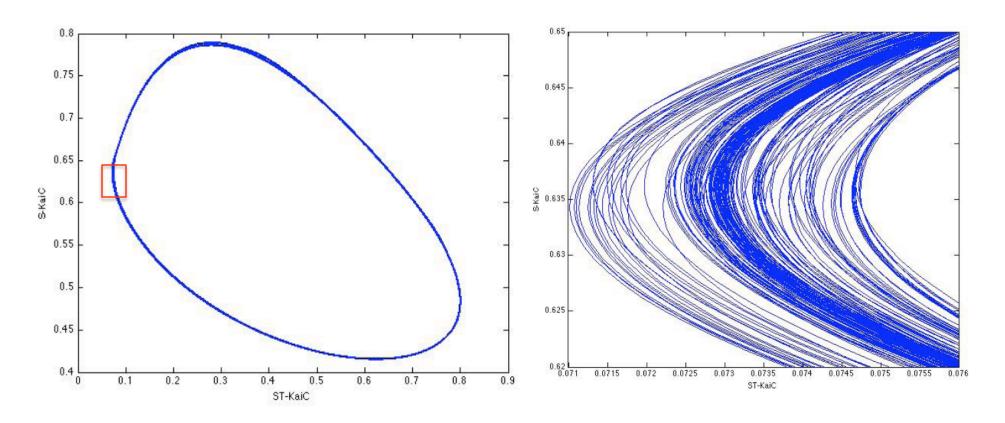
Simulation results of oscillation with matlab:



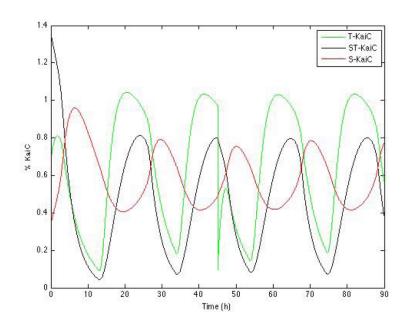


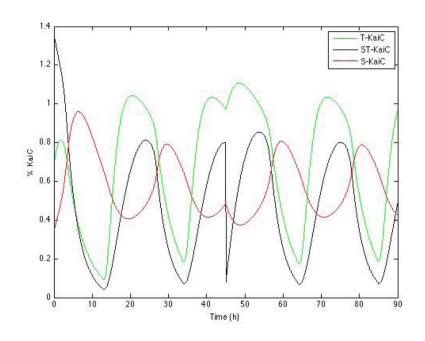
Rust, M.J., et. al. Science 2007.

2-D phase portrait:



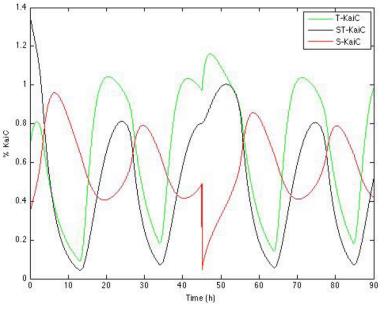
- Not exactly periodical, but "chaotic" or periodical with a very long period
- Long-term average period is 20.6865h



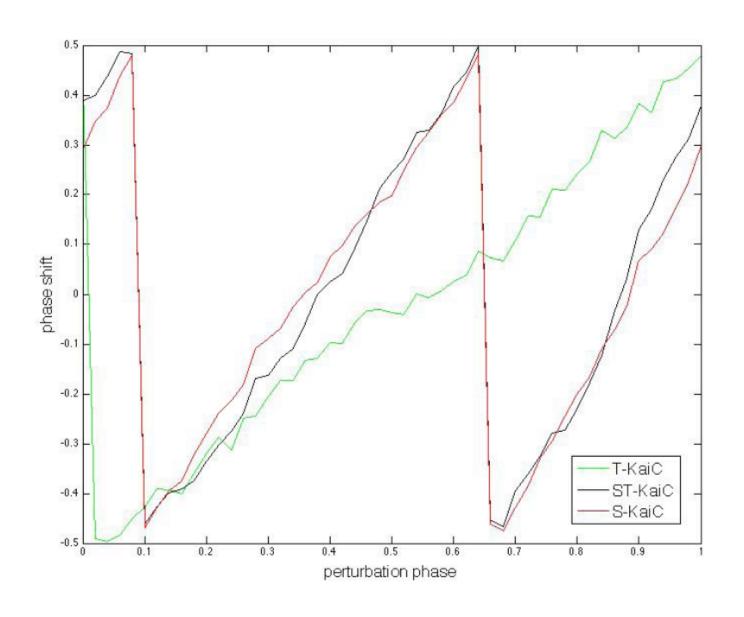


- **Perturbation simulations:** 1/10 of the value of T-KaiC, ST-KaiC, or S-KaiC at 45h
- Phase shift: $2\pi \times$ the fraction of a period that the original unperturbed oscillatory curves would have to be shifted in time to match the perturbed oscillation once it recovers its full amplitude oscillation

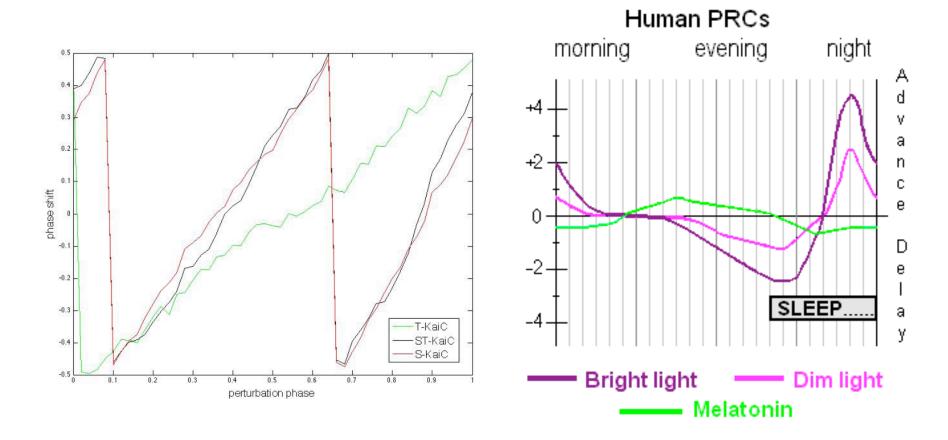
	T-KaiC	ST-KaiC	S-KaiC
phase- shift	1.00	2.20	2.26



Phase response curve (normalize 2π to 1 here)



Phase response curves



Summary

- Successful simulation of the KaiABC minimal model by Matlab
- Discussion about periodicity and perturbations in the KaiABC oscillator
- Phase response curve (PRC) generation for the model and comparison to natural PRC
- Some discussion