	j	k	1	m	n
Index	PER	CRY	Kinase	Location	BMALs-CLK
0	No PER bound	No CRY	No Kinases	Cytoplasm	No BMALs-
		bound	bound		CLK bound
1	PER1	CRY1	CKI	Nucleus	BMALs ^P -CLK ^P
2	PER1 ^P by CKI	CRY2	GSK3		
3	PER2		CKI&GSK3		
4	PER2 ^p by CKI		Inh:CKI		
5	PER2 ^P by		Inh.CVI 0.CCV2		
	GSK3		Inh:CKI&GSK3		
6	PER2 ^P by both				
	GSK3 and CKI				

Table S7. The variables of protein complexes used in the mathematical model. We represent each complex as x[j][k][l][m][n], where j, k, l, and n refer to the proteins that are present in the complex and m refers the location of the complex. Newly added variables that describe protein complex bound to CK1 inhibitor (PF-670462) are highlighted in bold. x[j][k][4 or 5][0][n] and x[j][k][4 or 5][1][n] correspond Inh:CK1_c and Inh:CK1_n in Figure S2, respectively. Details for the interpretation of variables can be found in Supplementary Table S2 in Kim and Forger, MSB (2012).