

**Mathematical Cell Biology Graduate Summer Course**  
**University of British Columbia, May 1-31, 2012**  
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# Biochemical motifs (5)



[www.math.ubc.ca/~keshet/MCB2012/](http://www.math.ubc.ca/~keshet/MCB2012/)

# Basic GTPase signaling modules: other feedback and context



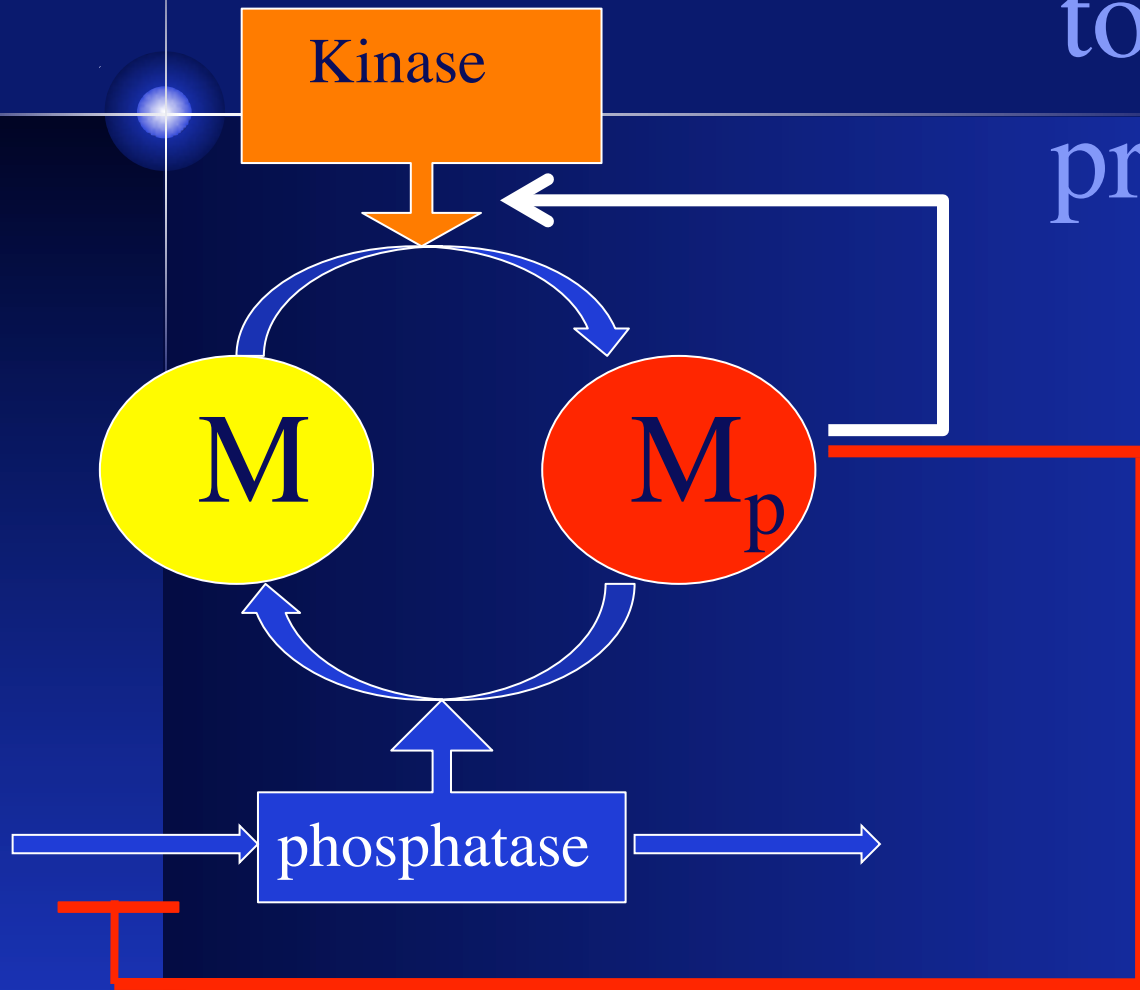
B.N. Kholodenko. Cell-signalling dynamics in time and space.  
Nature Reviews Molecular Cell Biology, 7(3):165–176, 2006

# Other feedback

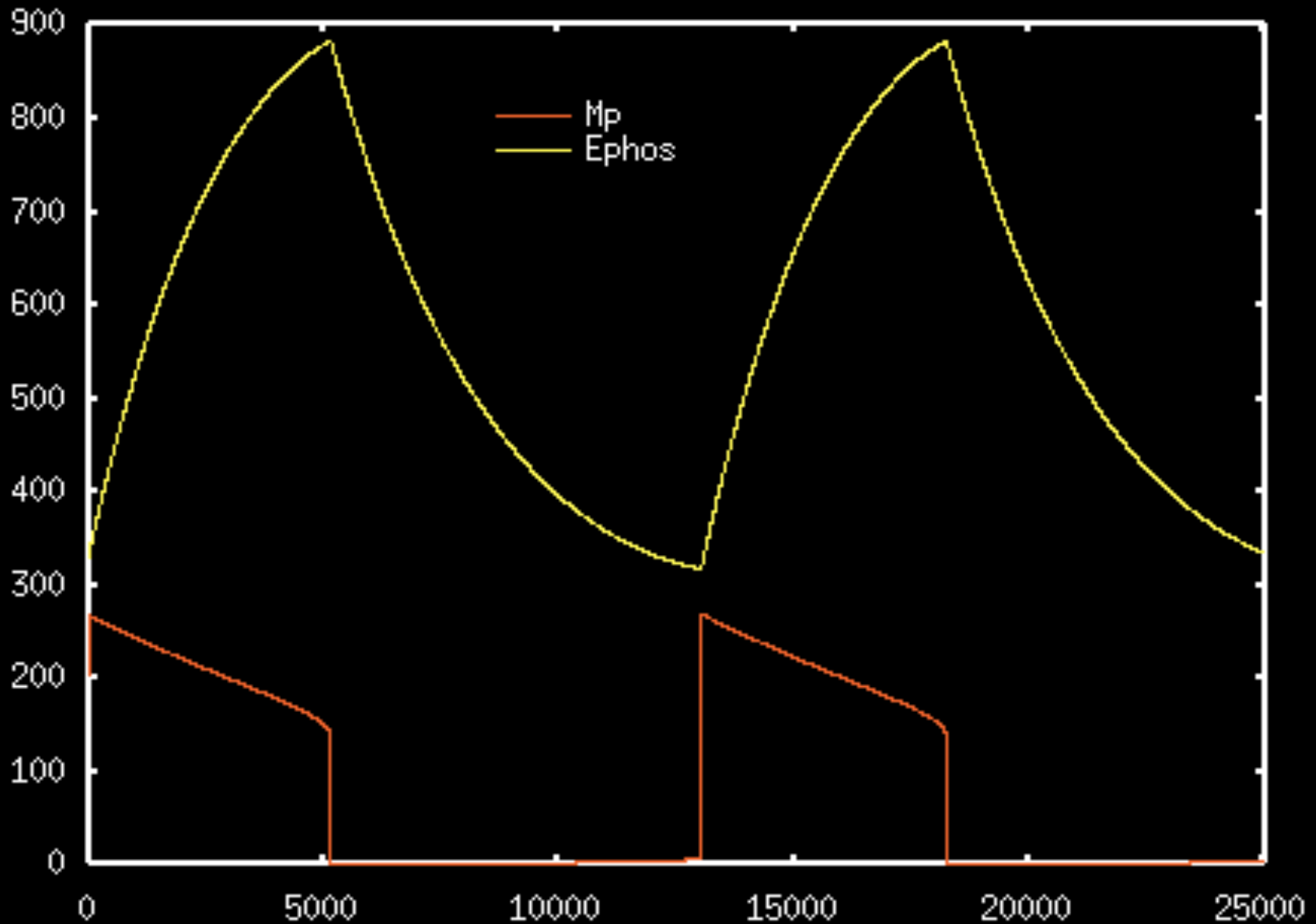
There are many other feedback effects possible.

Here is one more

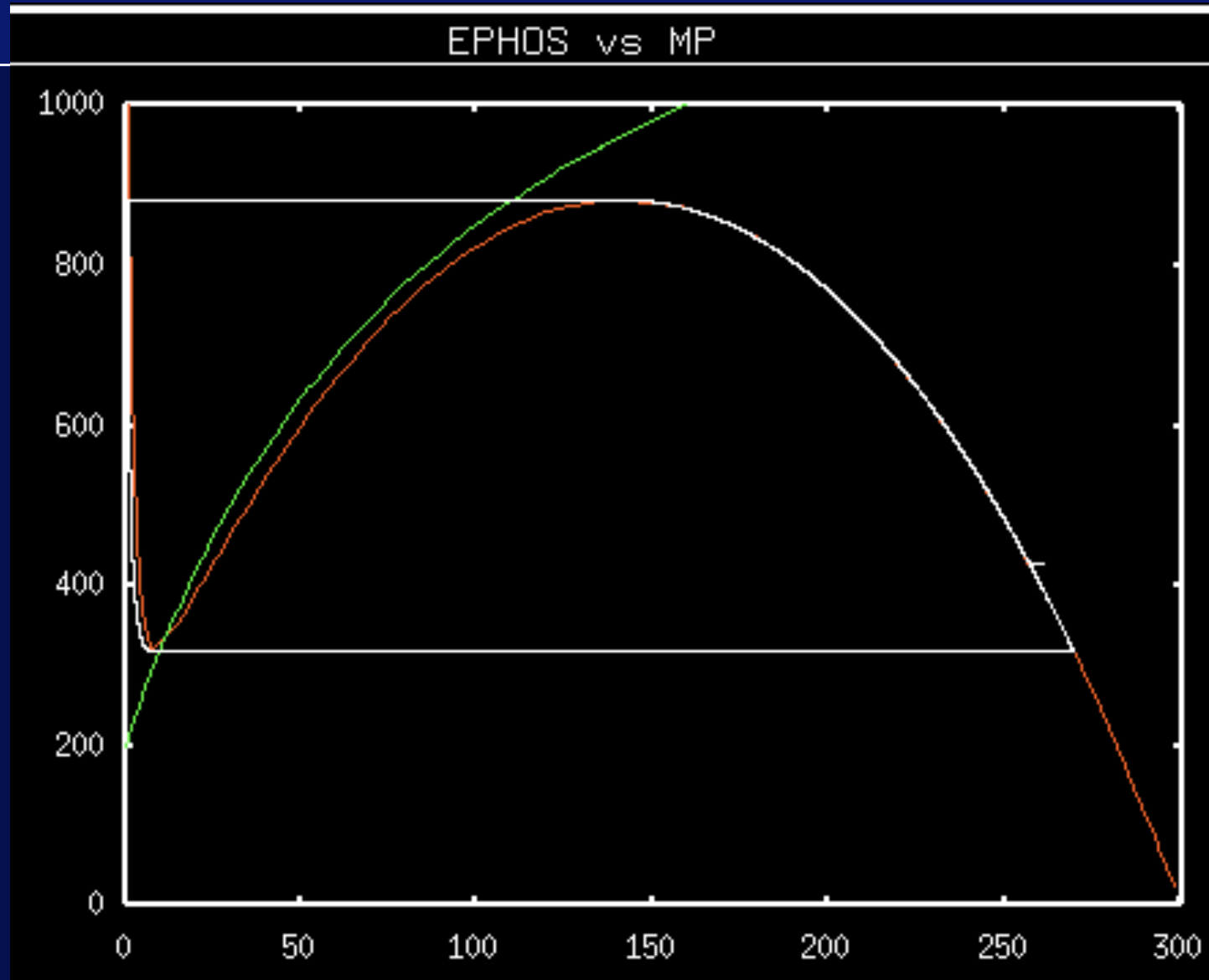
# Positive feedback to phosphatase production rate



# Time Behaviour of GTPase model



# Stable limit cycle



# XPP file

```
• # Kholod06Oscill_c.ode
• #
• # Kholodenko (2006) Nat Rev Mol Cell Bio 7, p 165
• # Box 2 Example (c)
• #
• # M is protein and Mp is its phosphorylated form
• # Ephos is the phosphatase

• Mp'=vkin(Mp,Mtot-Mp)-vphos(Mp,Mtot-Mp)
• Ephos'=v_phossyn(Mp)-v_phosdeg(Ephos)

• vkin(Mp,M)=((k_kincat*Ekin*M)/(Km1+M))*((1+A*(Mp/Ka))/(1+(Mp/Ka)))
• vphos(Mp,M)=k_phoscat*Ephos*Mp/(Km2+Mp)

• v_phossyn(Mp)=V_phos0*(1+Ap*(Mp/Kd))/(1+(Mp/Kd))
• v_phosdeg(Ephos)=k_phosdeg*Ephos

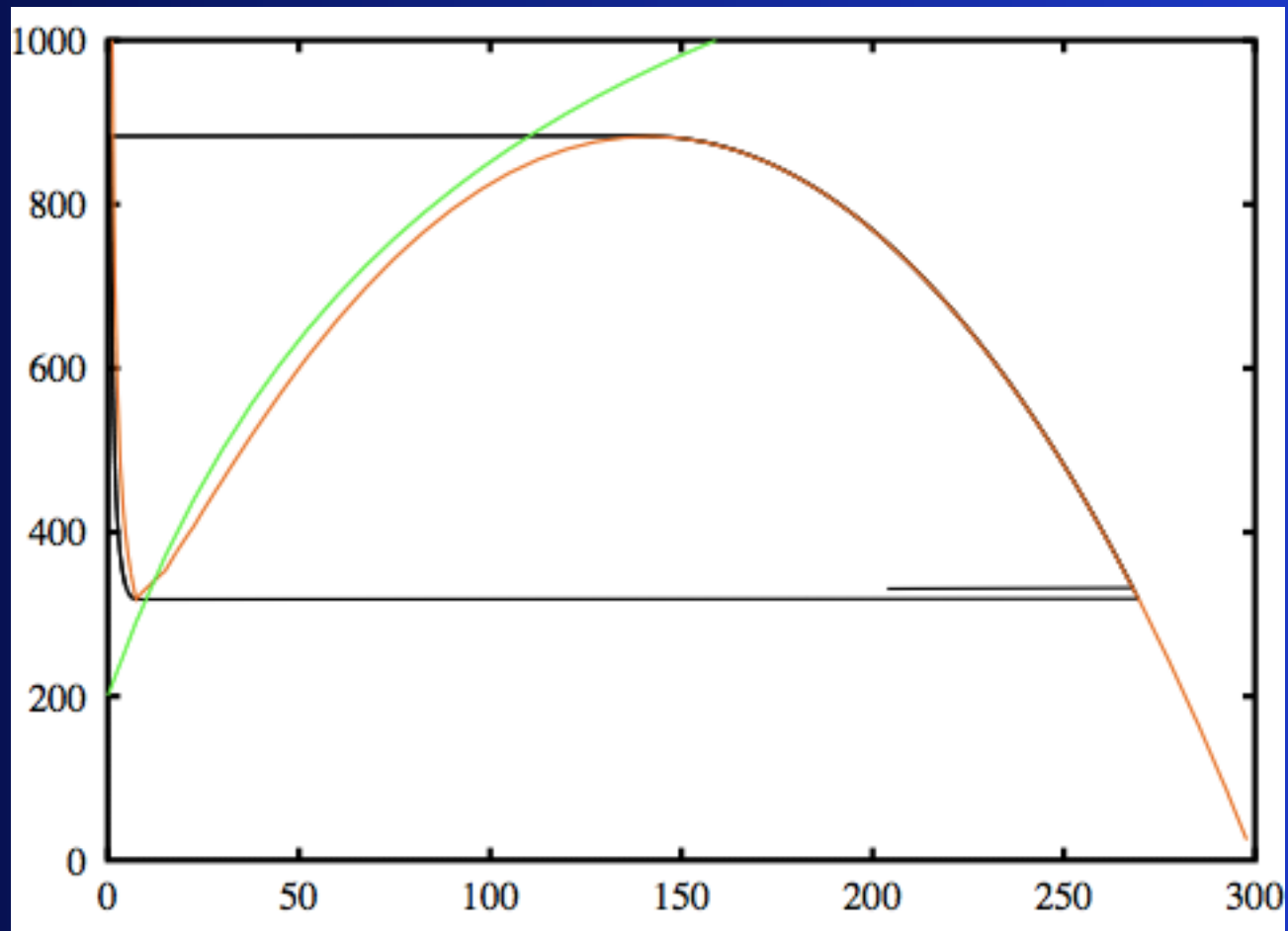
• # these params have units that include per sec already
• par k_kincat=1,A=100,Ka=500,Km1=500,Ekin=150
• par k_phoscat=1,Km2=10
• par Mtot=300

• #Convert Kholodenko's params to per sec
• # par V_phos0=200nm/hr, k_phosdeg=1/hr now in per sec:
• par V_phos0=0.0555,Kd=100,Ap=7.5,k_phosdeg=0.000277

• init Mp=300,Ephos=100

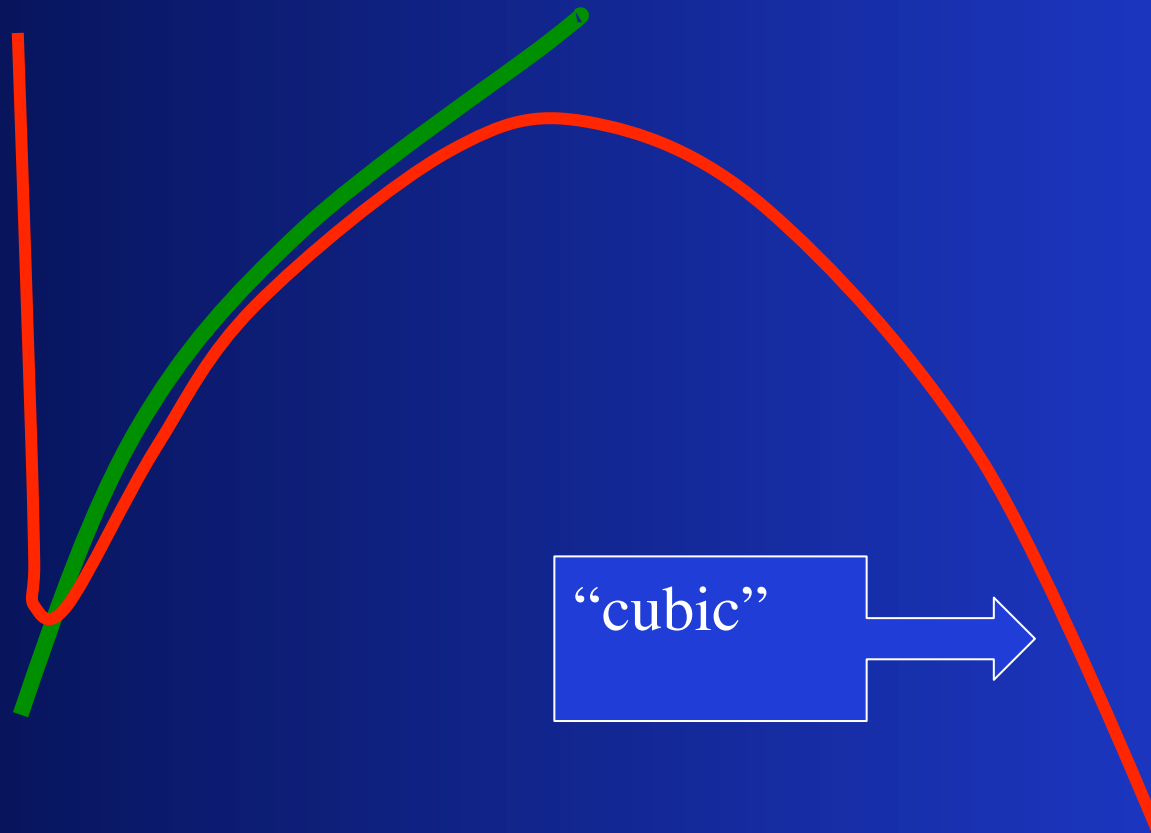
• @ dt=0.01,total=50000,nout=500,xp=Mp,yp=Ephos,xlo=0,xhi=300,ylo=0,yhi=1000,bounds=10000
• done
```

# Stable limit cycle

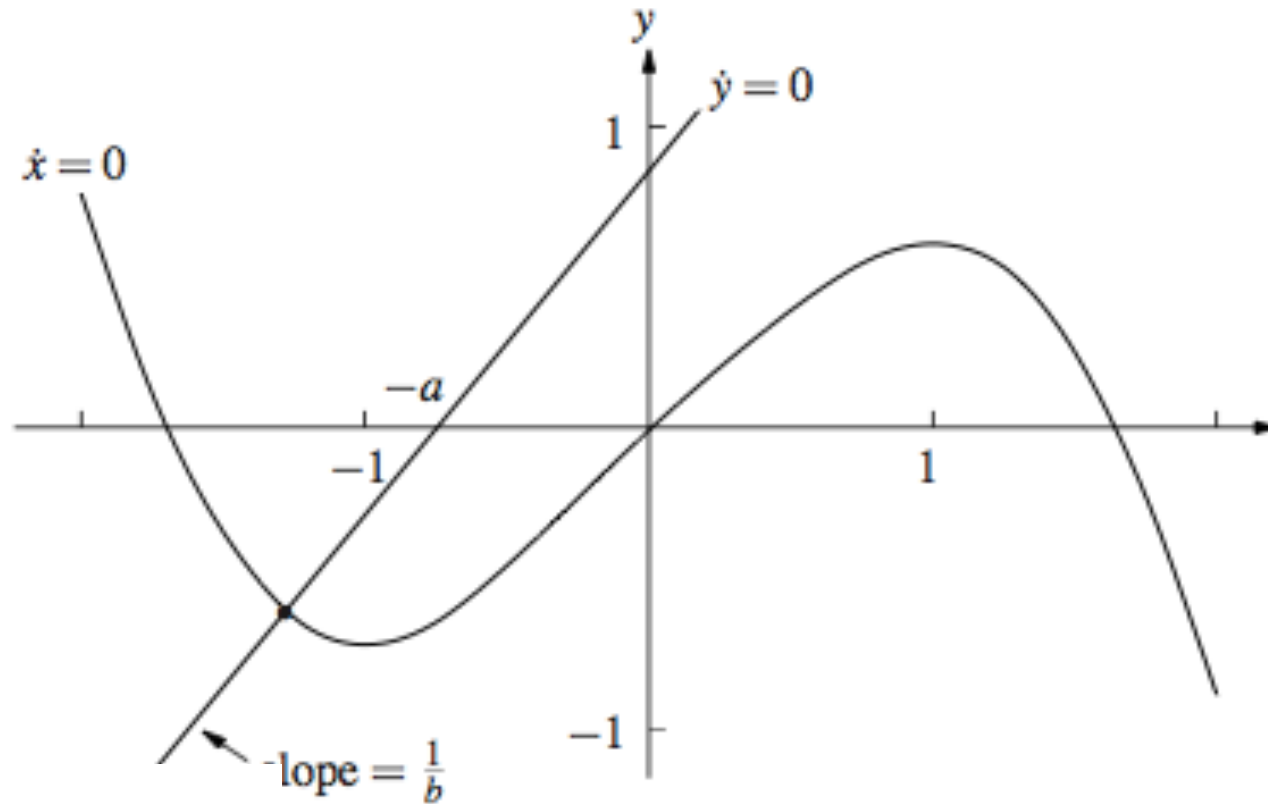




# Nullclines



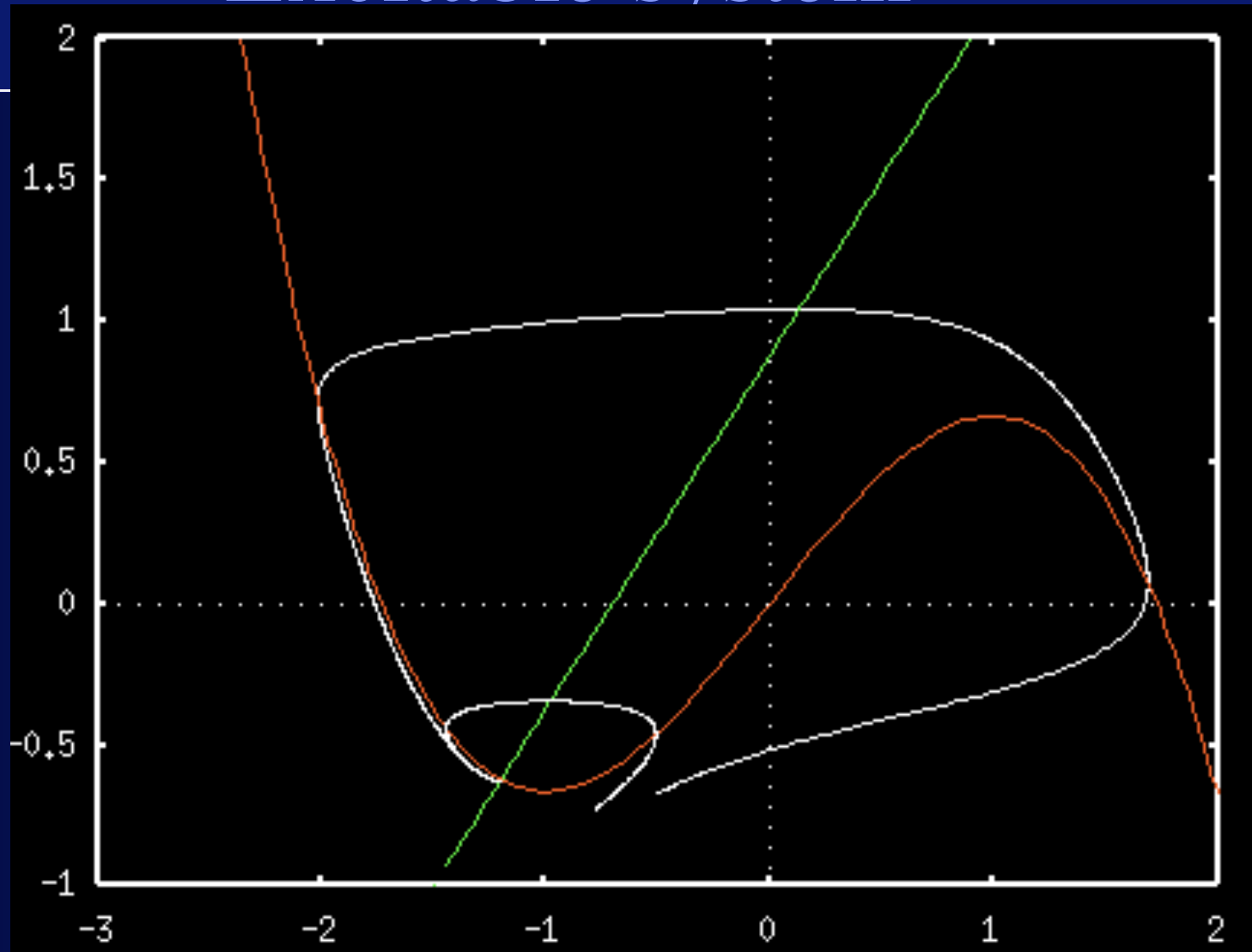
# Classic FitzHugh Model



$$\frac{dx}{dt} = c \left[ x - \frac{1}{3}x^3 - y + j \right],$$

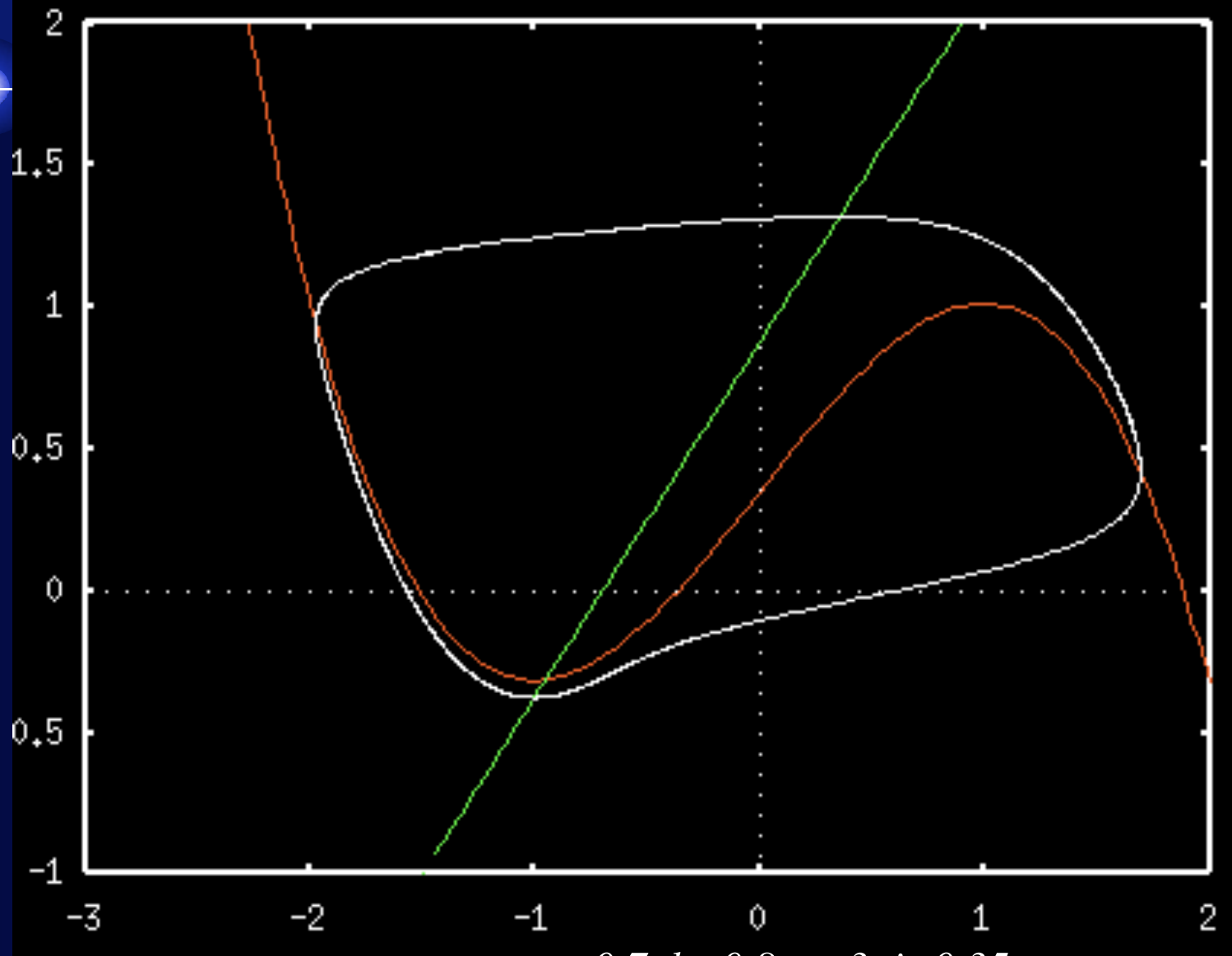
$$\frac{dy}{dt} = \frac{1}{c} [x + a - by].$$

# “Excitable system”



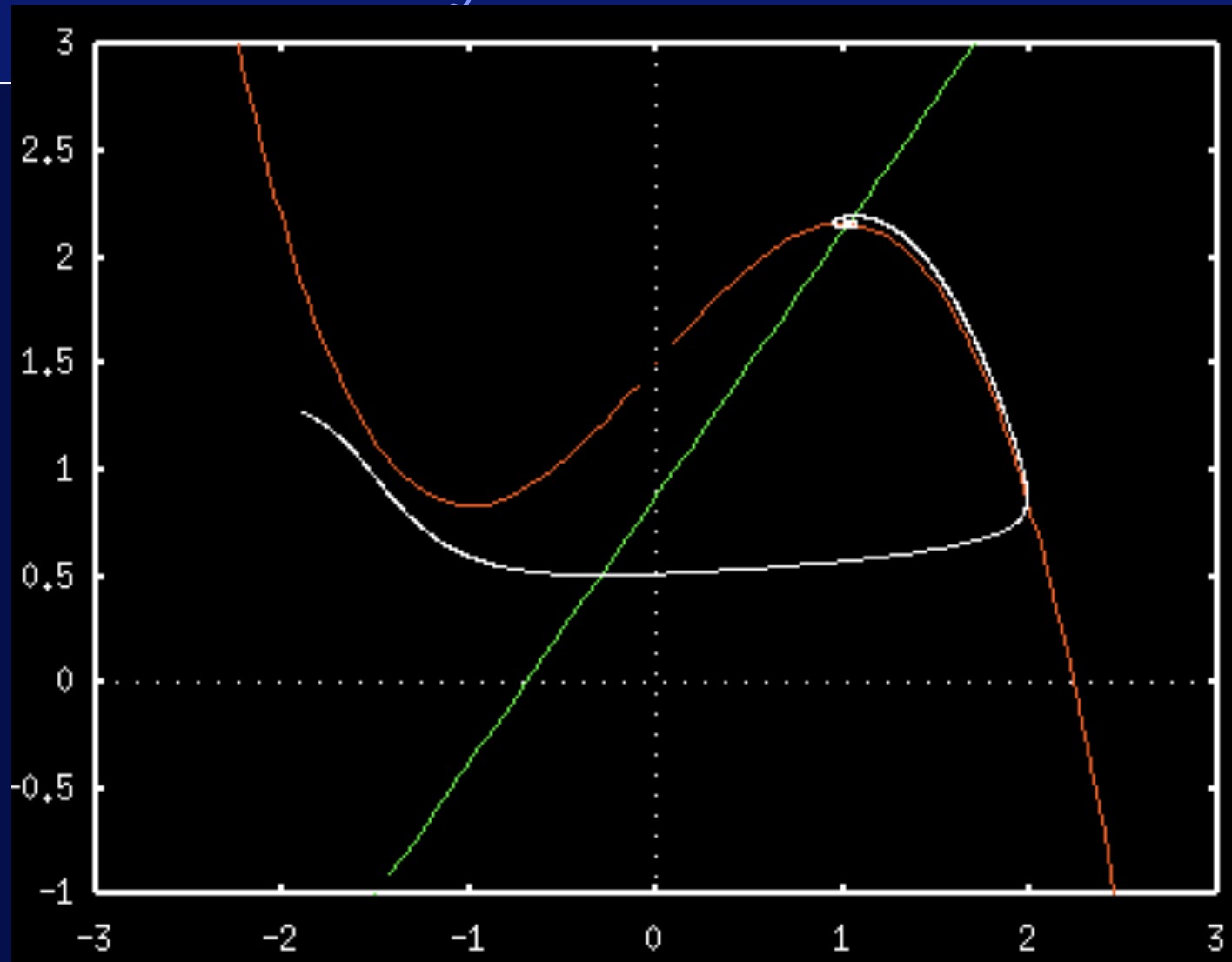
$$a=0.7, b=0.8, c=3, j=0$$

# FitzHugh Limit cycle



$$a=0.7, b=0.8, c=3, j=0.35$$

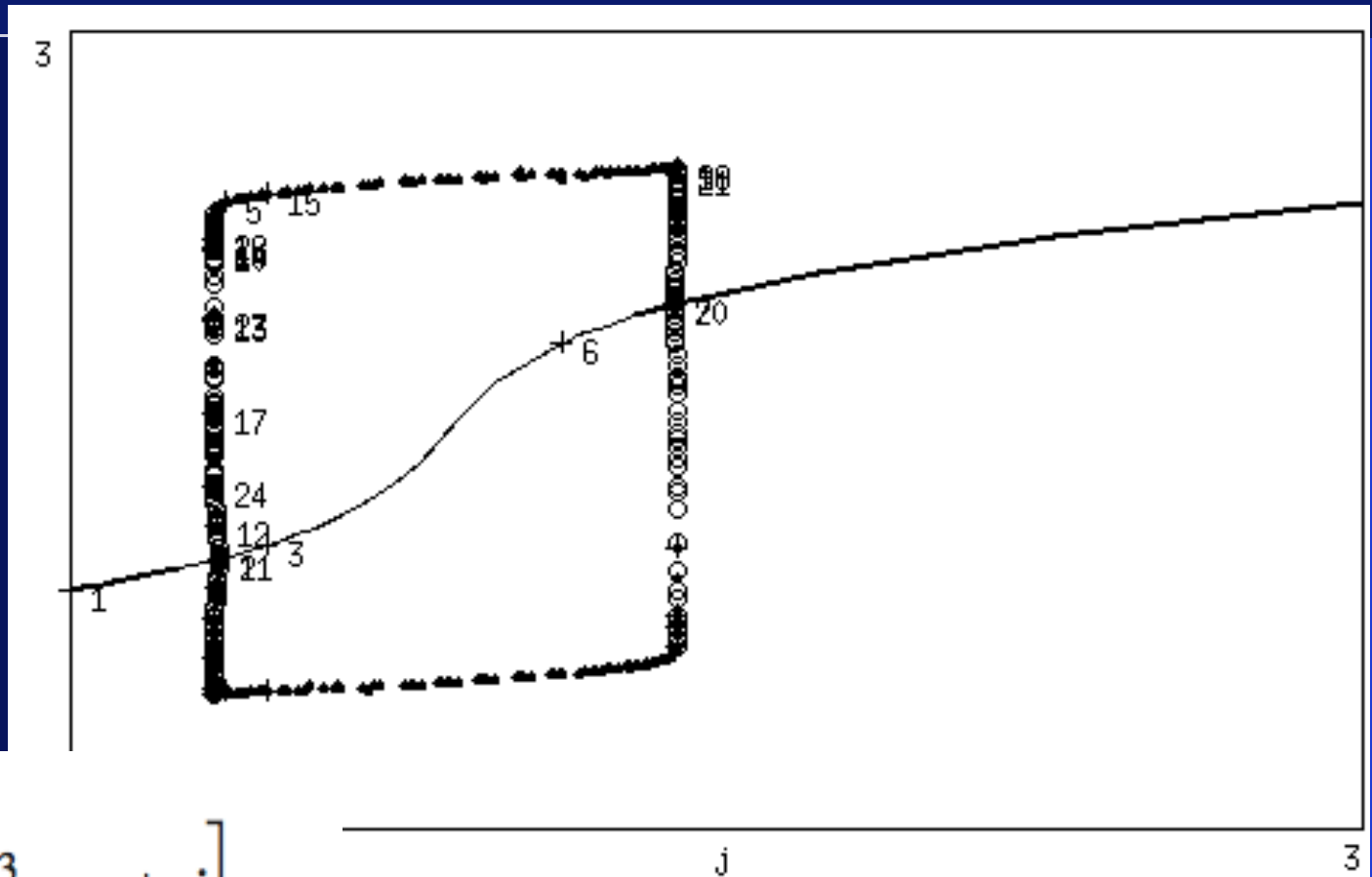
# Limit cycle vanishes



$$a=0.7, b=0.8, c=3, j=1.5$$

# Fitzhugh Bifurcation diagram

$x$



$$\frac{dx}{dt} = c \left[ x - \frac{1}{3}x^3 - y + j \right],$$
$$\frac{dy}{dt} = \frac{1}{c} [x + a - by].$$

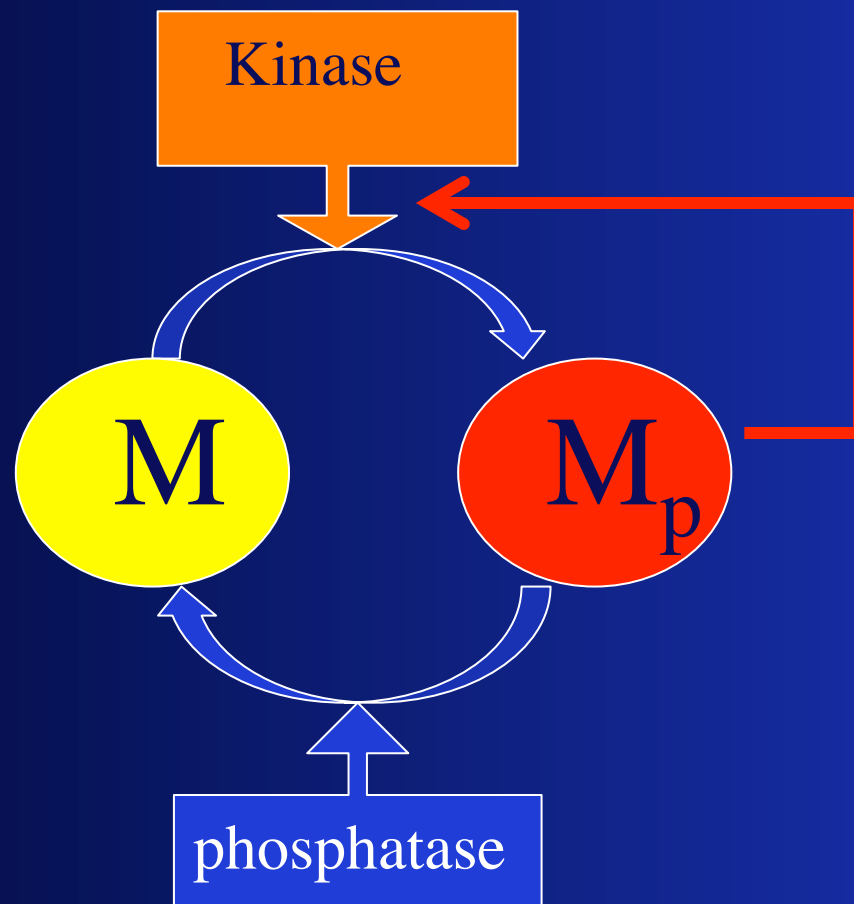
$j$

# Conclusions

- The basic GTPase cycle can become a bistable switch in 2 ways

# Ways of producing bistable switch

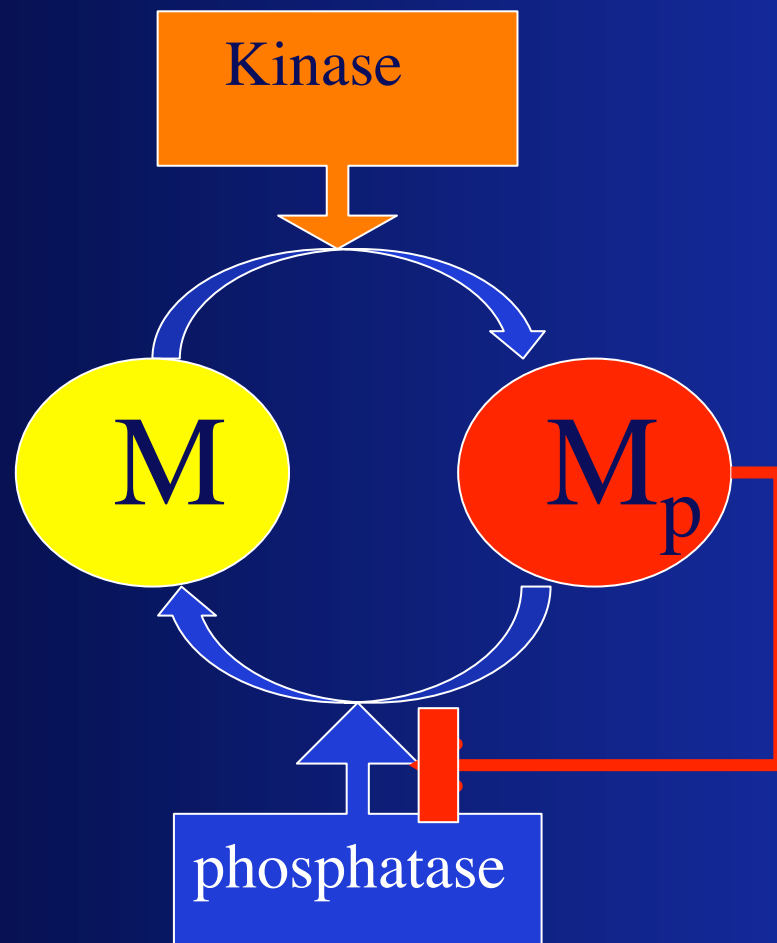
(1)





# Ways of producing bistable switch

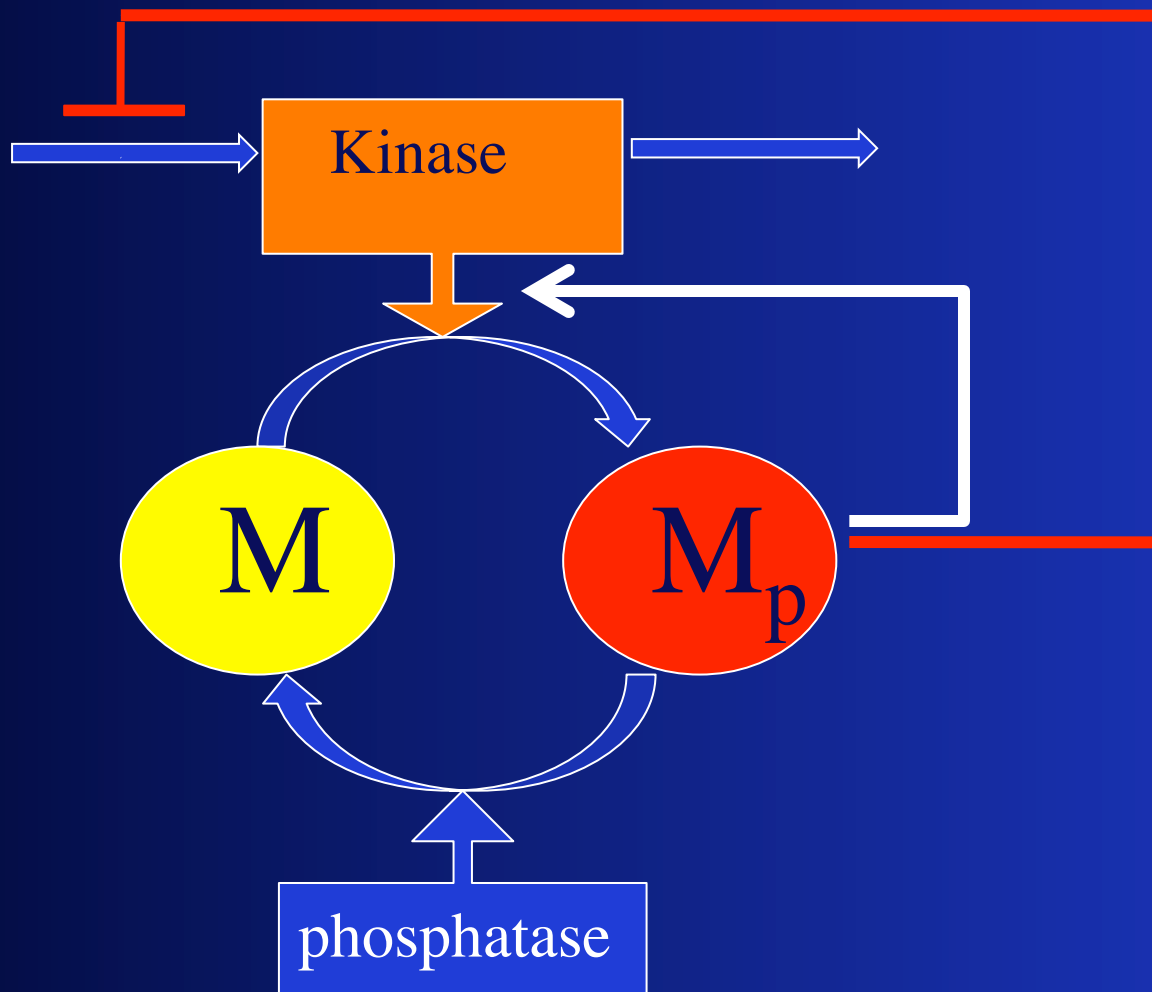
(2)



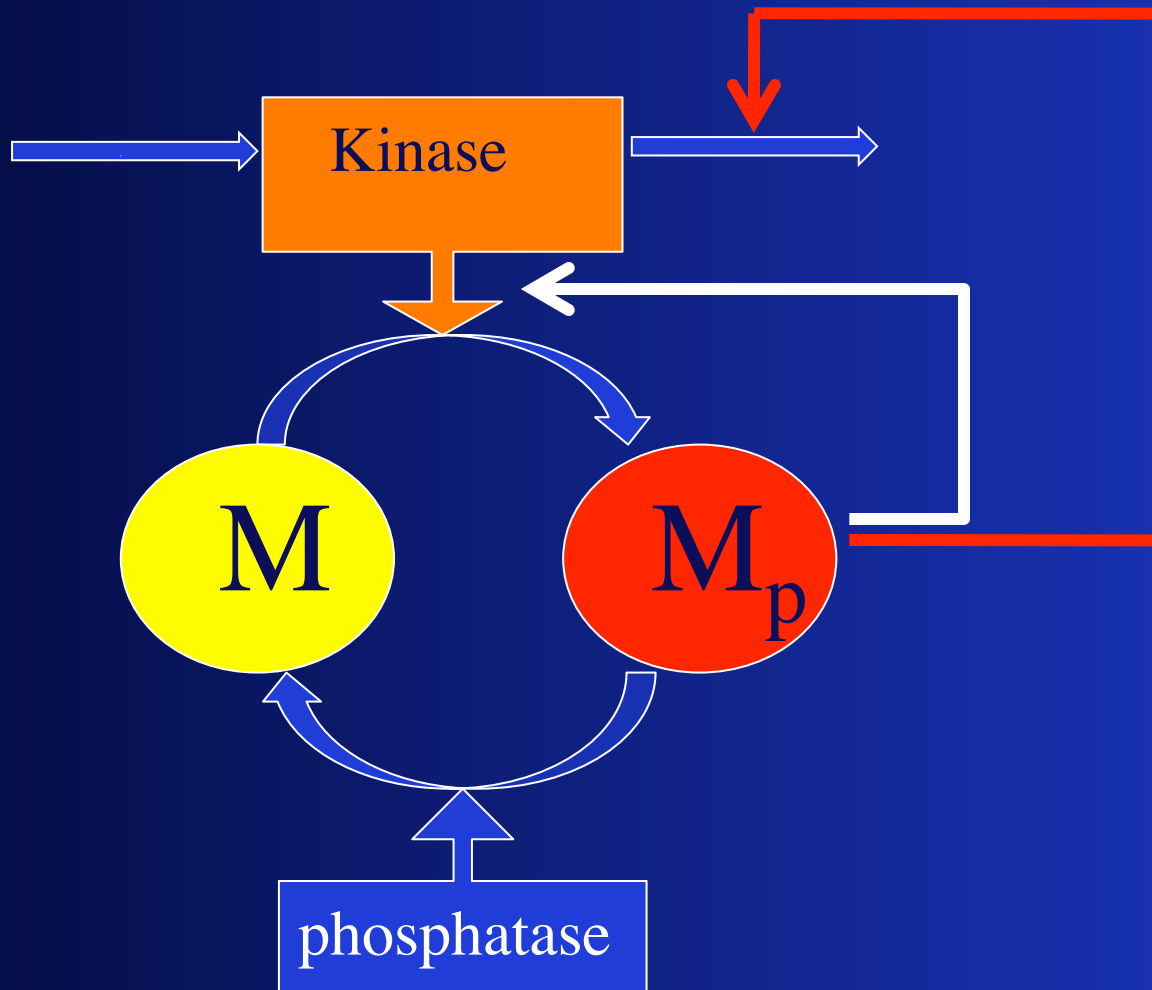
# An additional feedback can lead to relaxation oscillations

There are eight ways of achieving such behaviour

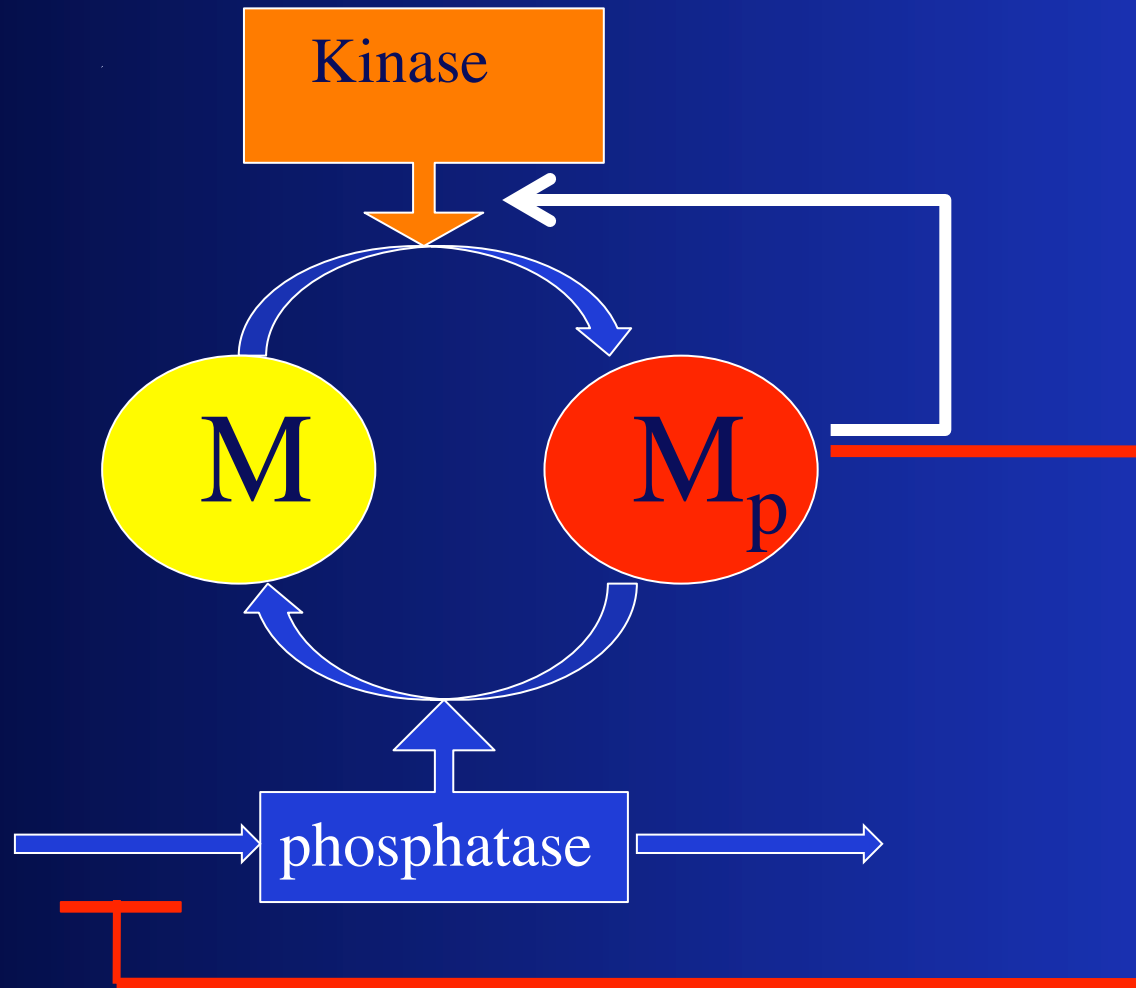
# Feedback to kinase (1)



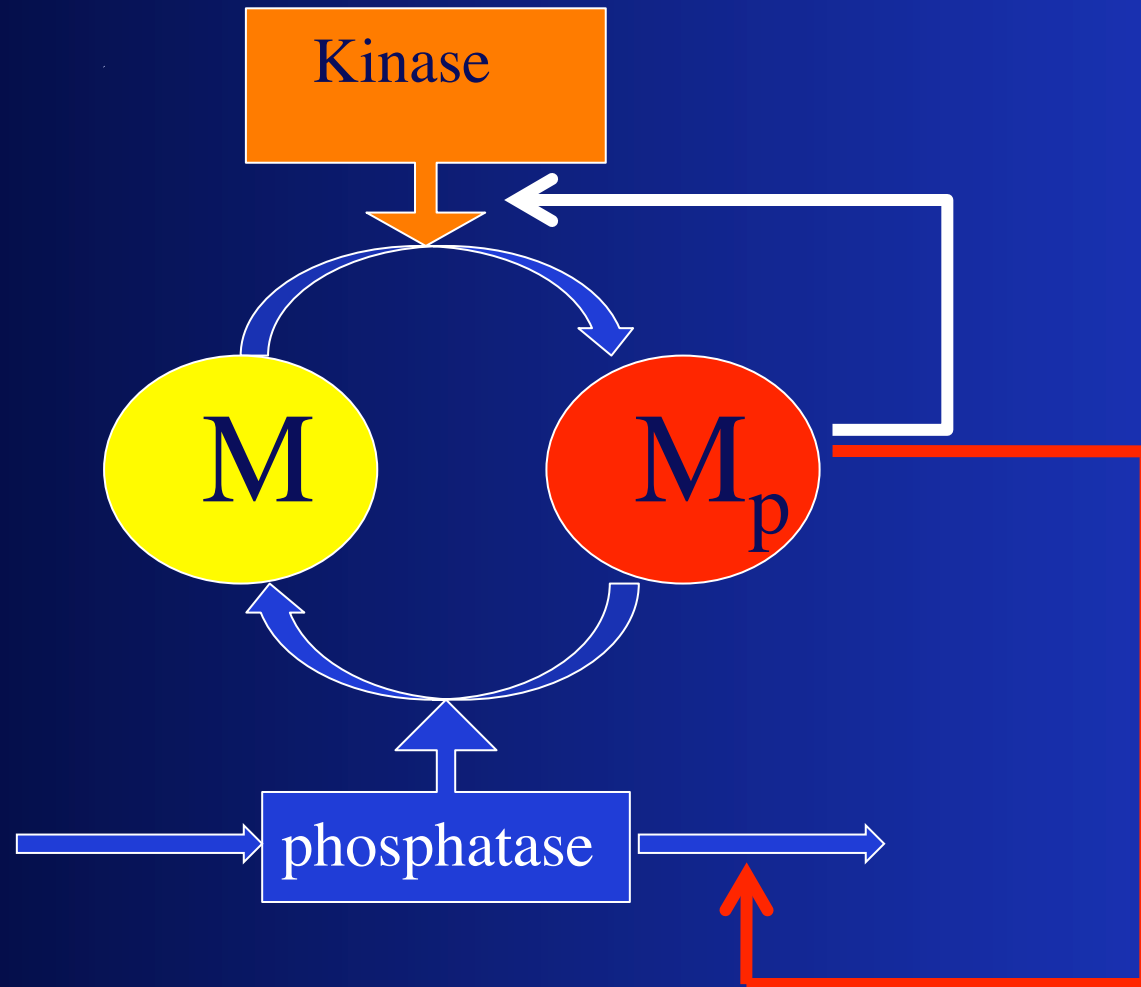
# Feedback to kinase (2)

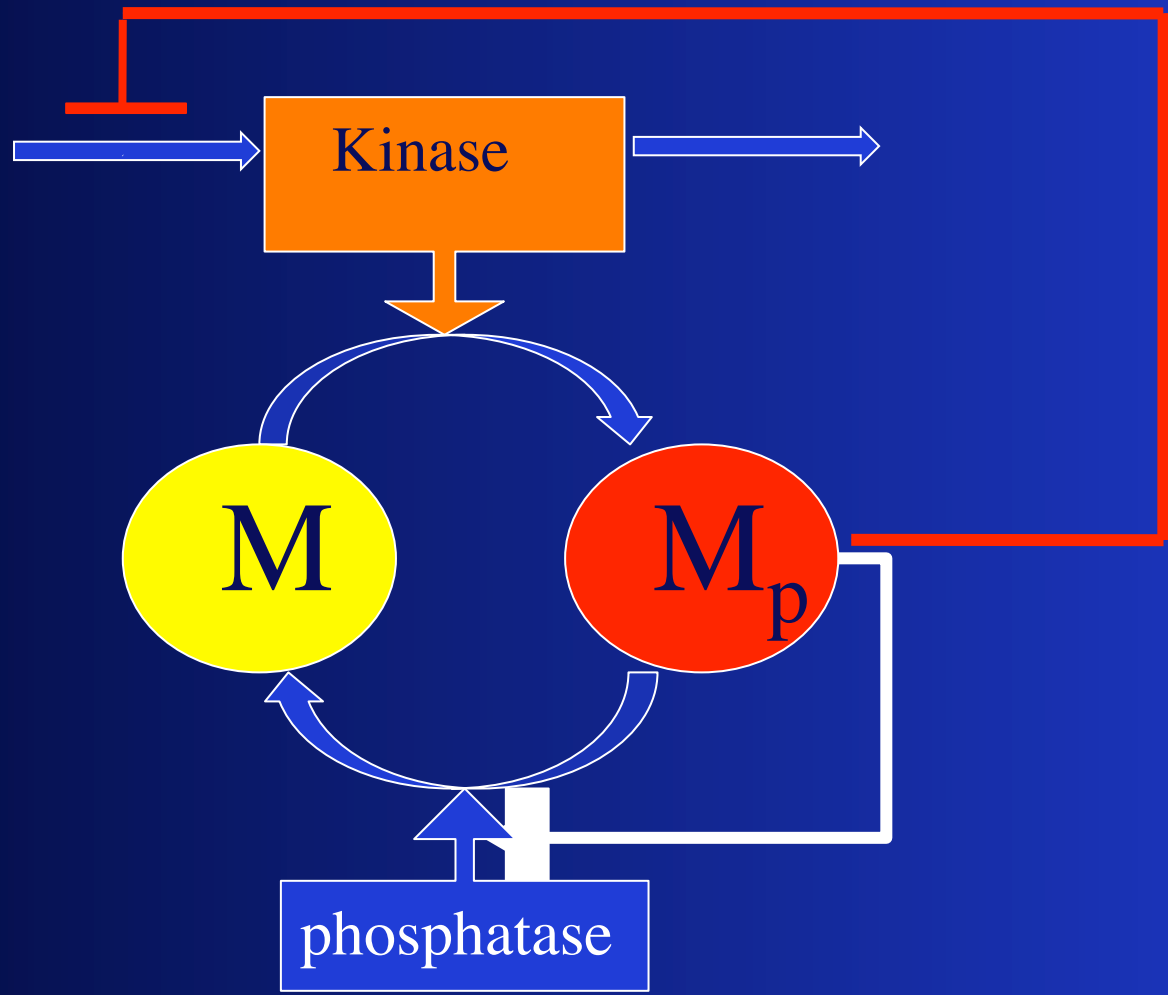


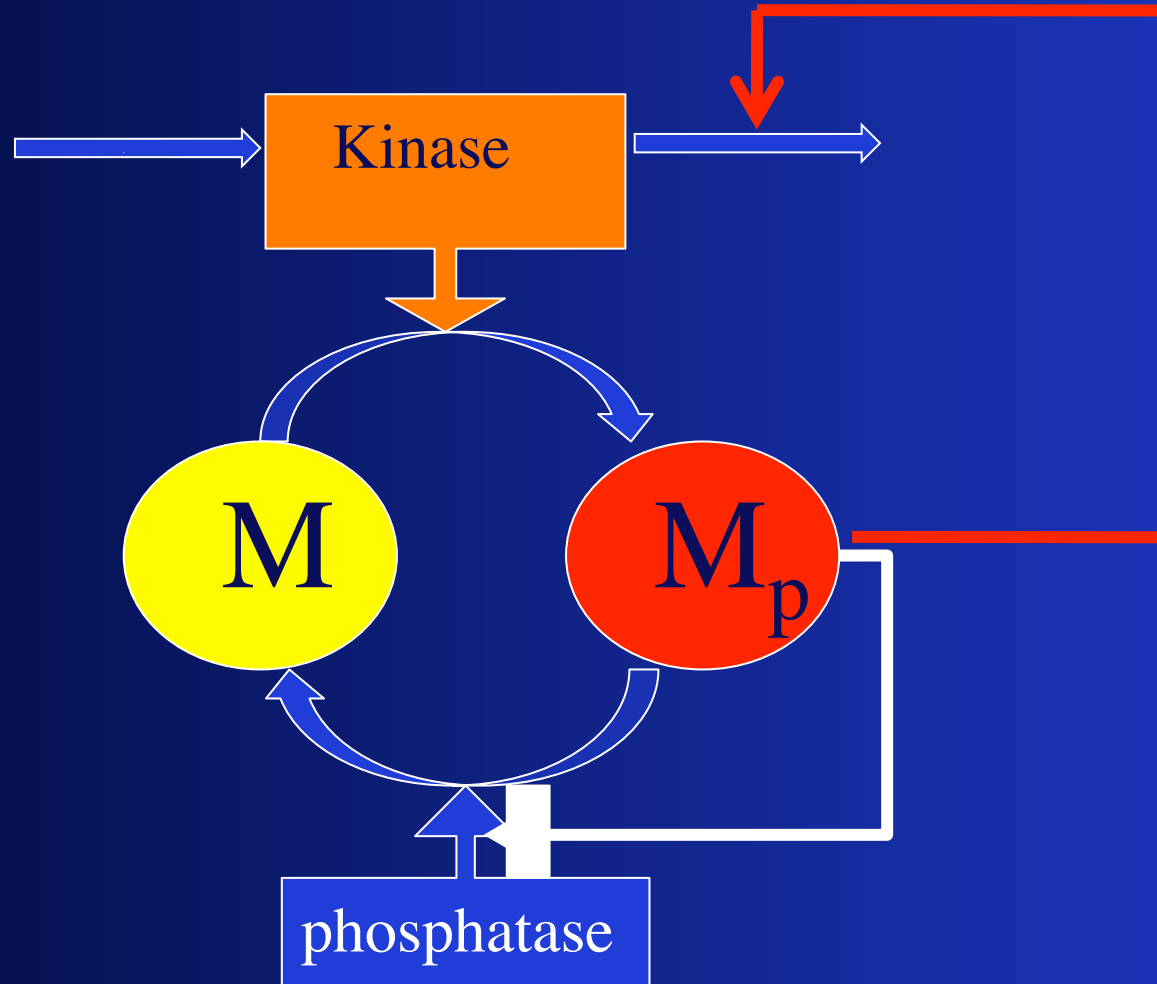
# Feedback to phosphatase (1)



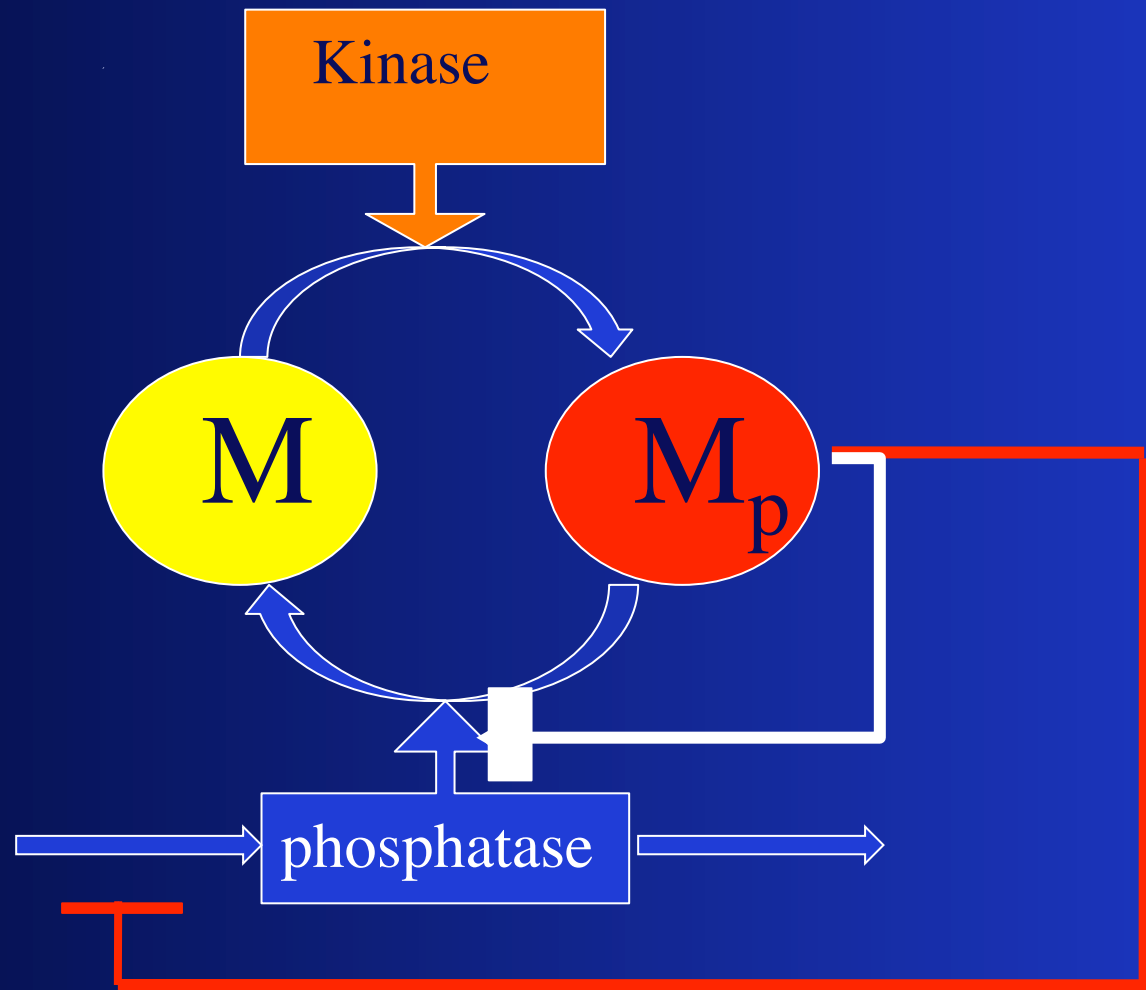
# Feedback to phosphatase (2)

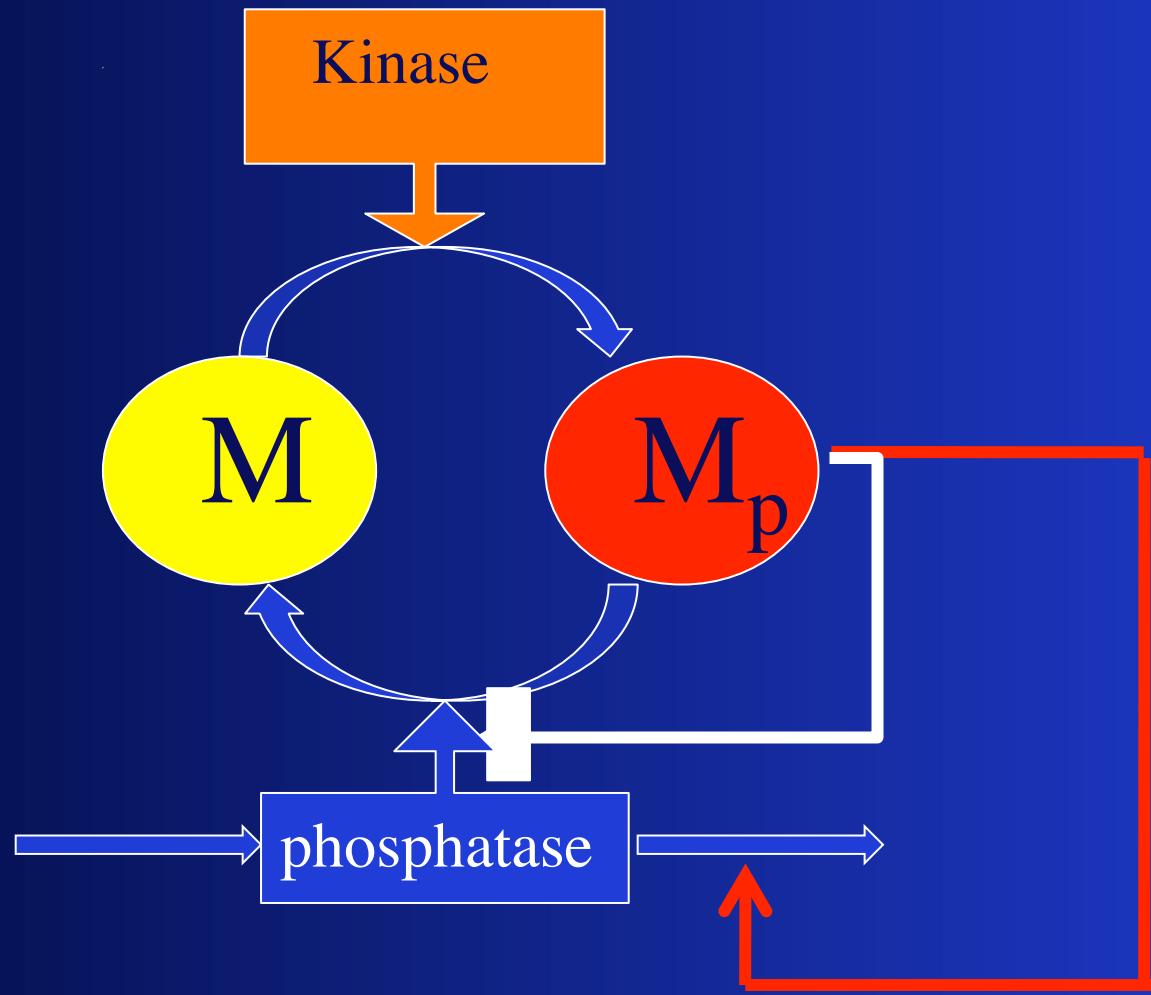












# Further comments

Distinct GTPases can also interact with one another in pairs, trios, etc, producing larger circuits of positive and negative feedback networks.

See:

M.A. Tsyganov, W. Kolch, and B.N. Kholodenko.  
The topology design principles that determine the spatiotemporal dynamics of G-protein cascades. *Mol. BioSyst.*, 8(3):730–743, 2012.