

Cellular networks

Combinatorial complexity of gene regulation

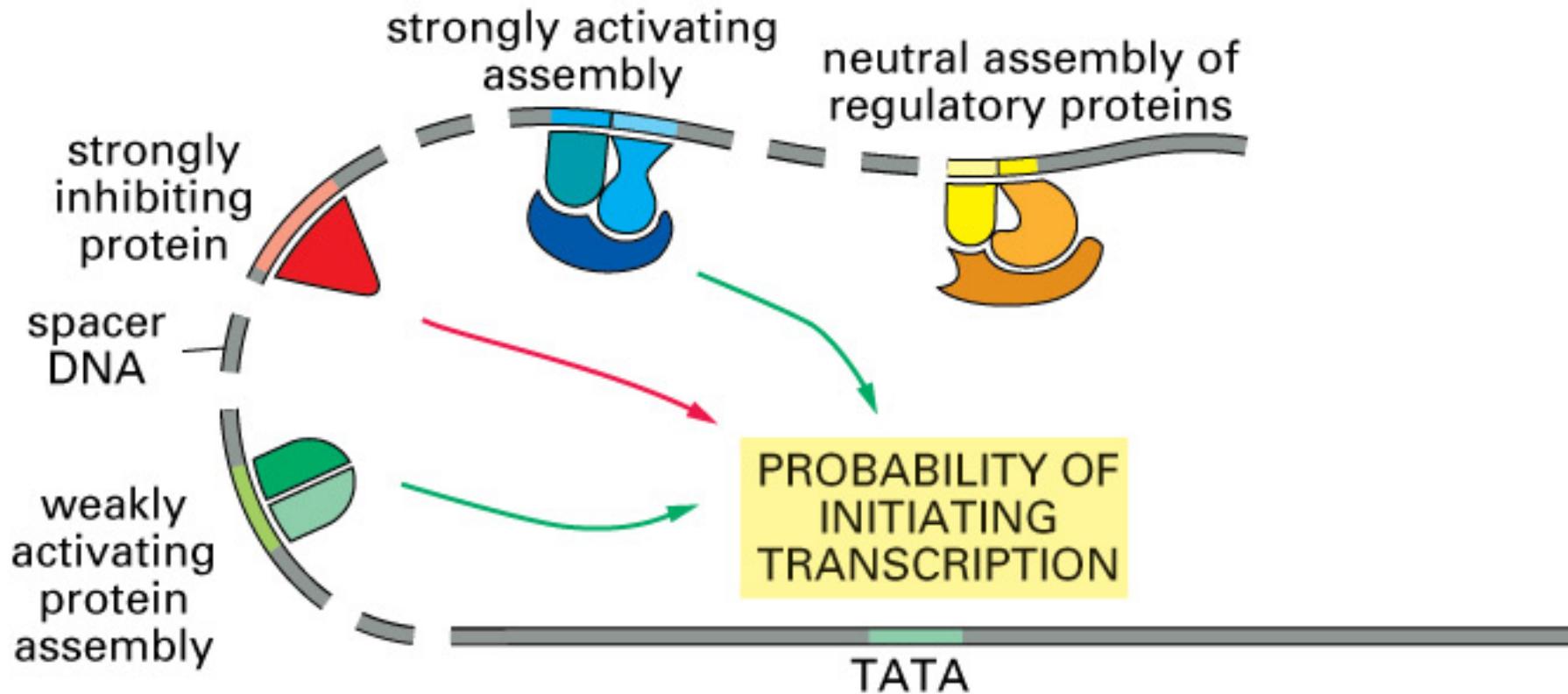
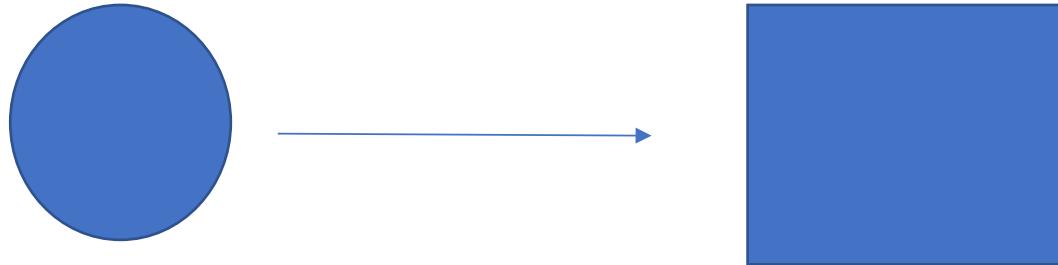


Figure 7–57. Molecular Biology of the Cell, 4th Edition.

Transcription regulatory networks (TRN)



Components of TRNs (also known as Gene Regulatory Networks (GRNs)):

Transcription regulatory network patterns

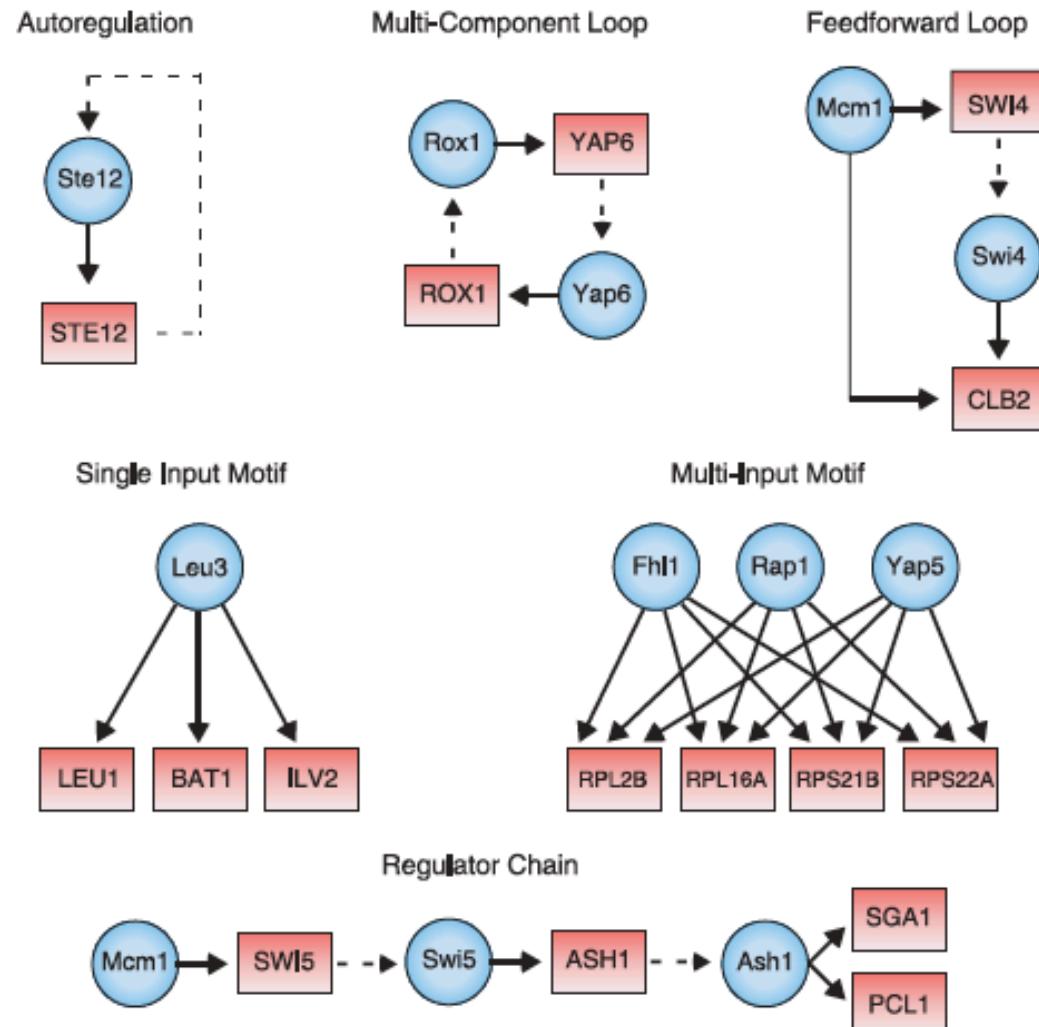
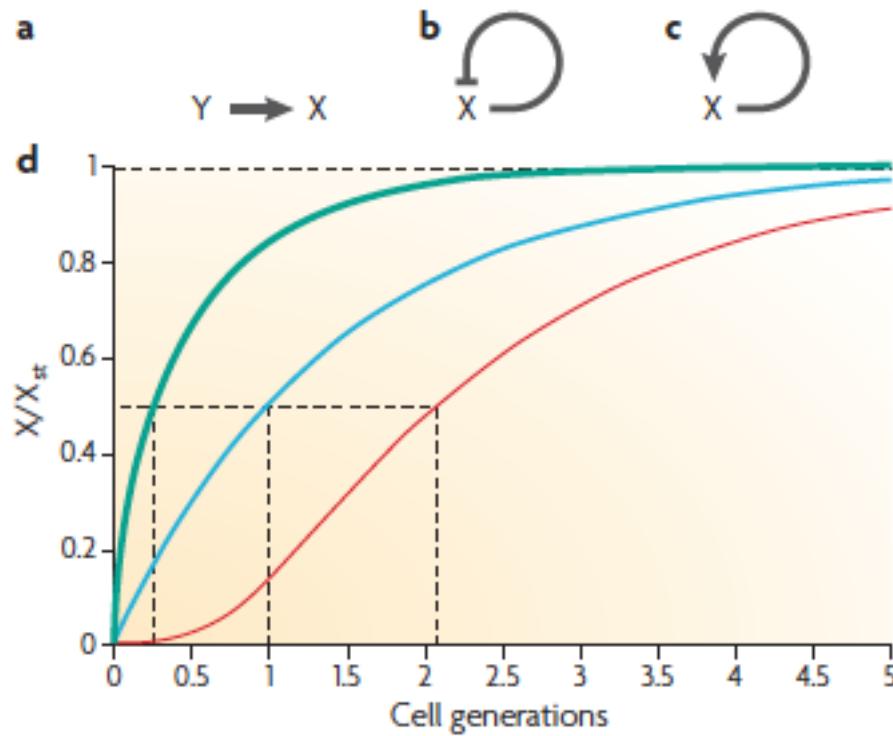


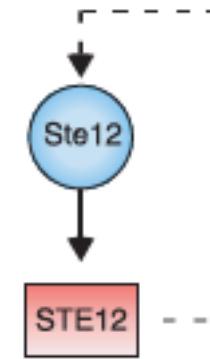
Fig. 3. Examples of network motifs in the yeast regulatory network. Regulators are represented by

Lee et al., Science 1999.

Connection: molecular mechanism \rightarrow motif topology \rightarrow gene expression dynamics



Autoregulation

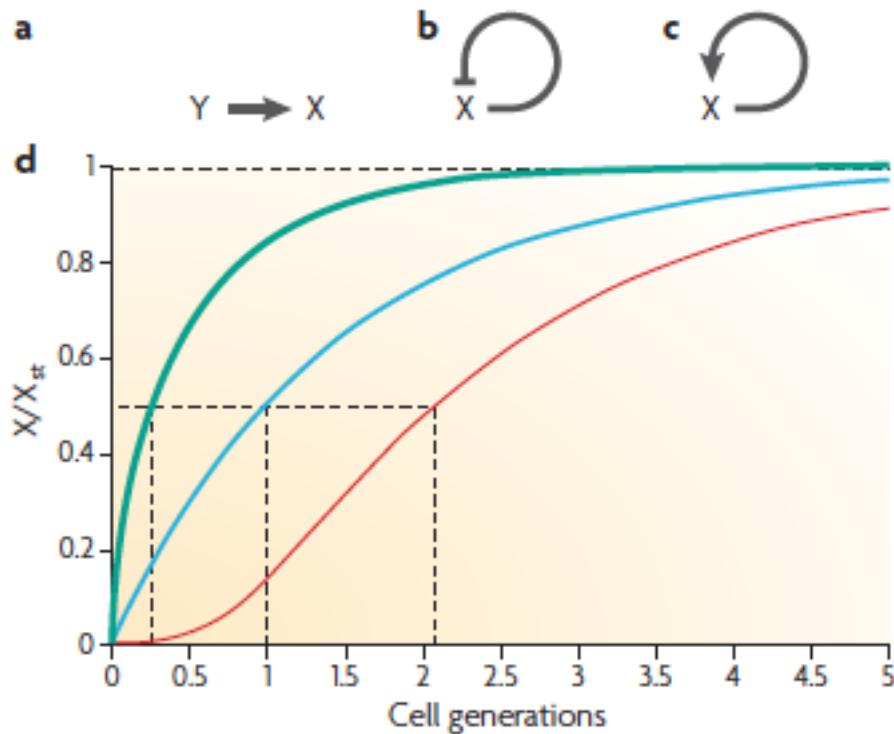


Positive autoregulation slows gene expression response time.

Negative autoregulation speeds gene expression response time.

Shen-Orr et al., 2002 Nature Genetics
Lee et al., Science 1999.

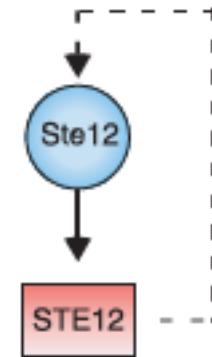
Autoregulation



Positive autoregulation _____

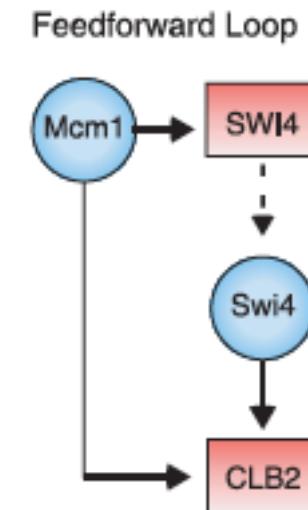
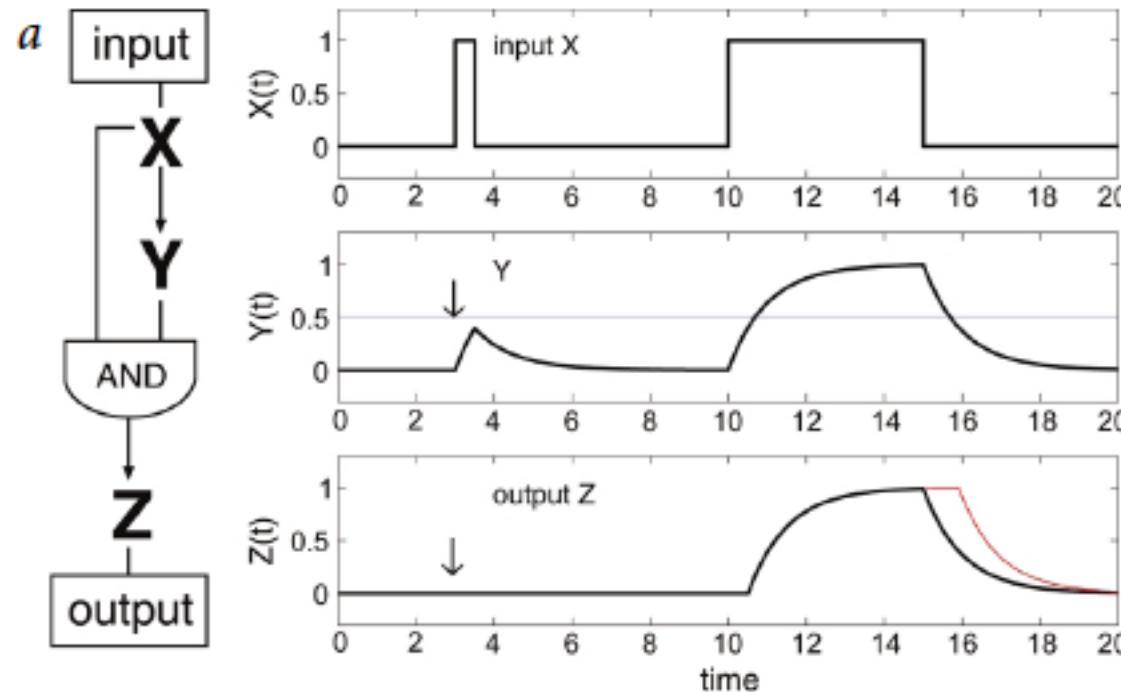
Negative autoregulation _____

Autoregulation



Shen-Orr et al., 2002 Nature Genetics
Lee et al., Science 1999.

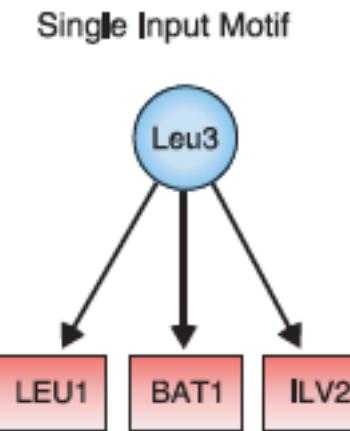
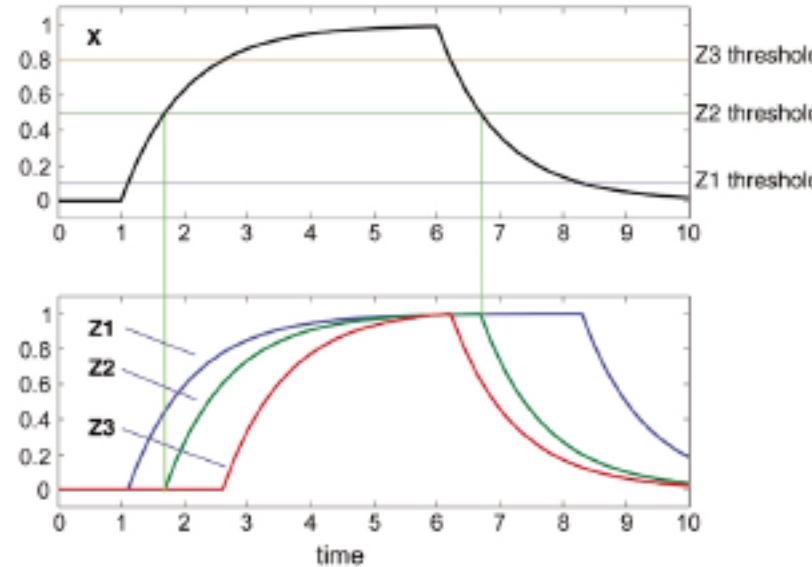
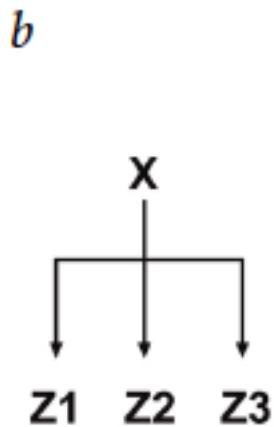
Feed forward loops.



Most common is coherent FFL. Dynamic behavior is _____.

Shen-Orr et al., 2002 Nature Genetics
Lee et al., Science 1999.

Single input motifs co-regulate genes that have similar functions.



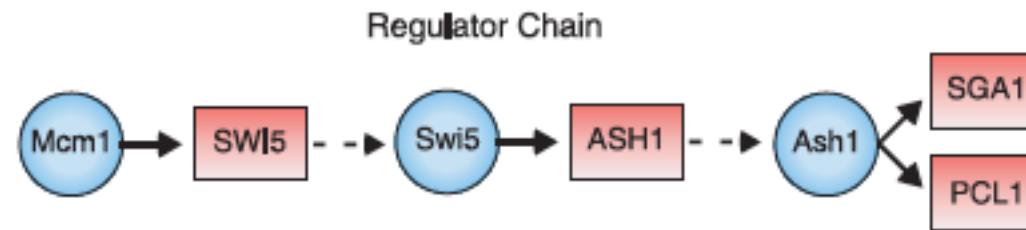
Shen-Orr et al., 2002 Nature Genetics
Lee et al., Science 1999.

Allows for _____ of expression with a single regulator.

Good for _____ processes or need for specific _____ in complexes.

If Z1, Z2, Z3 all have same _____, allows for same timing of expression

Connection: molecular mechanism → motif topology → gene expression dynamics and function



Inter-TF gene regulatory network of yeast

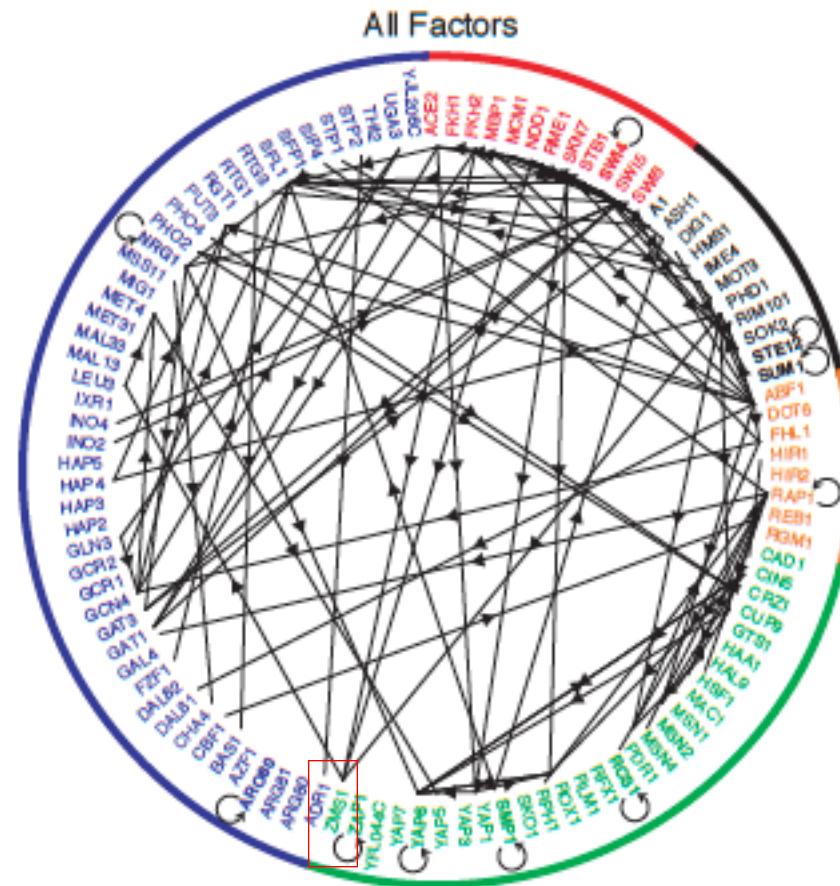
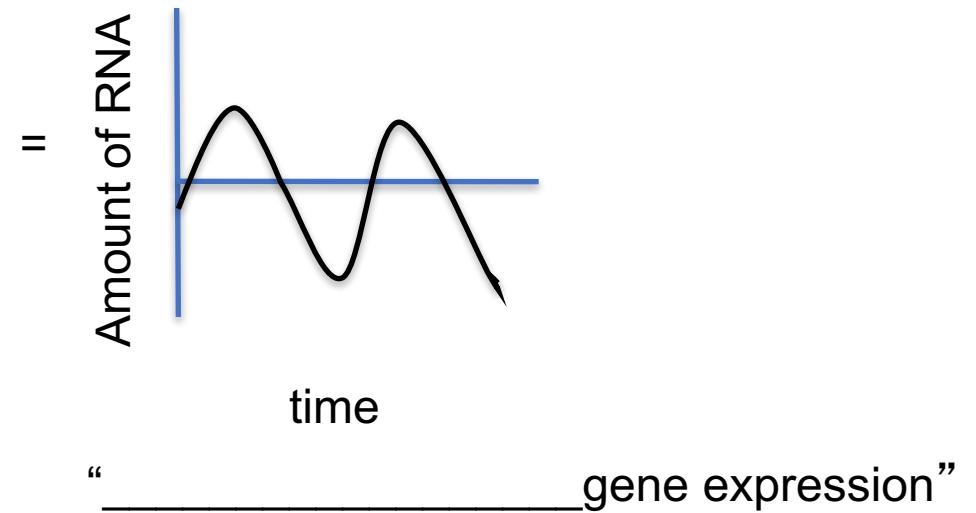
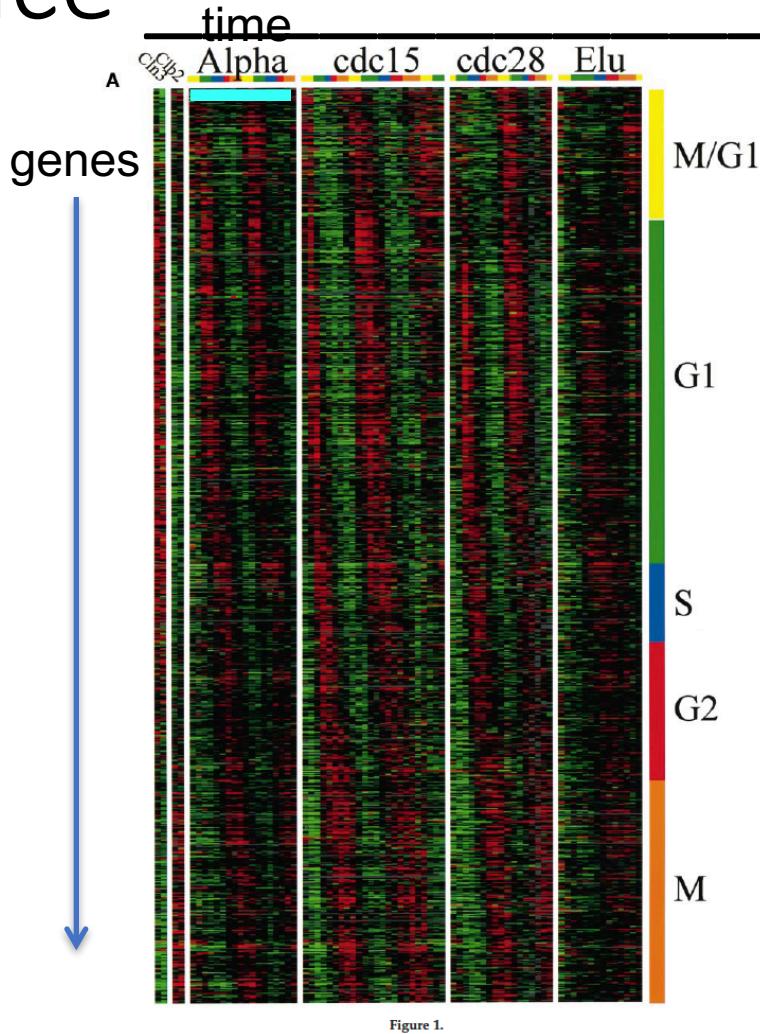


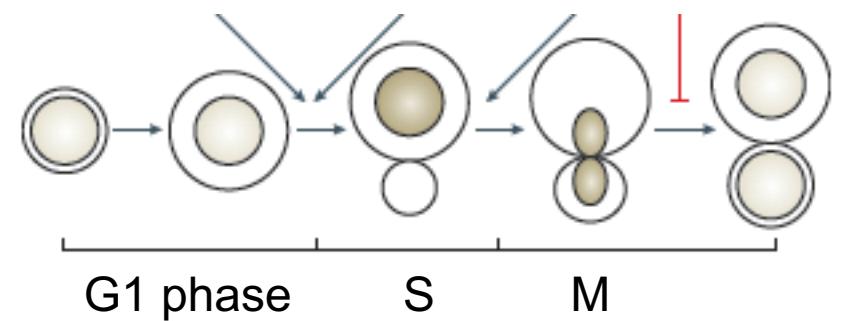
Figure 5, Lee et al., 2002

Transcription regulatory networks can produce oscillations in gene expression



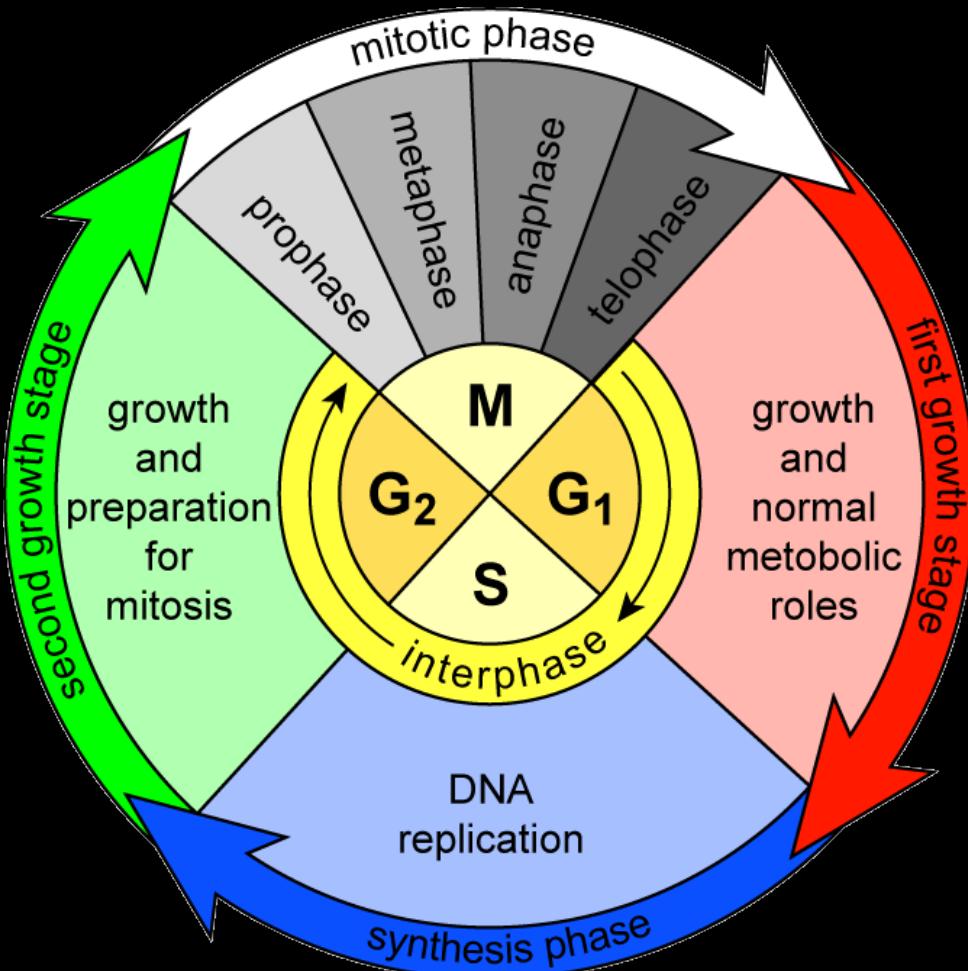
Spellman et al., Mol Biol Cell, 1998

Transcription networks produce gene expression oscillations to regulate the _____



from Bloom and Cross 2007

The cell cycle

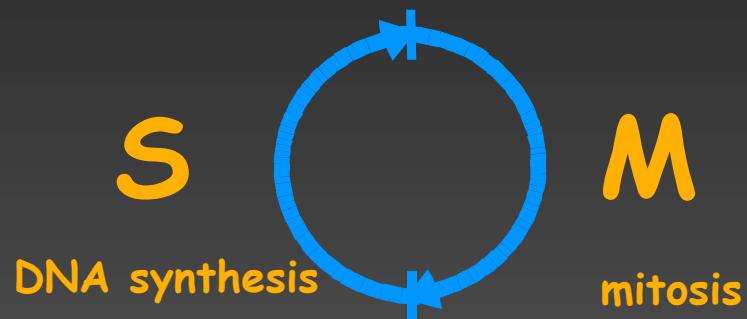


Mitosis

- Replicated _____, partitioning to two daughters.
- Mitosis in action: <https://www.youtube.com/watch?v=2J65DoinDKU>
- Think / pair / share: what would happen if cell division happened before mitosis was finished?
- How is progress through the cell cycle regulated?

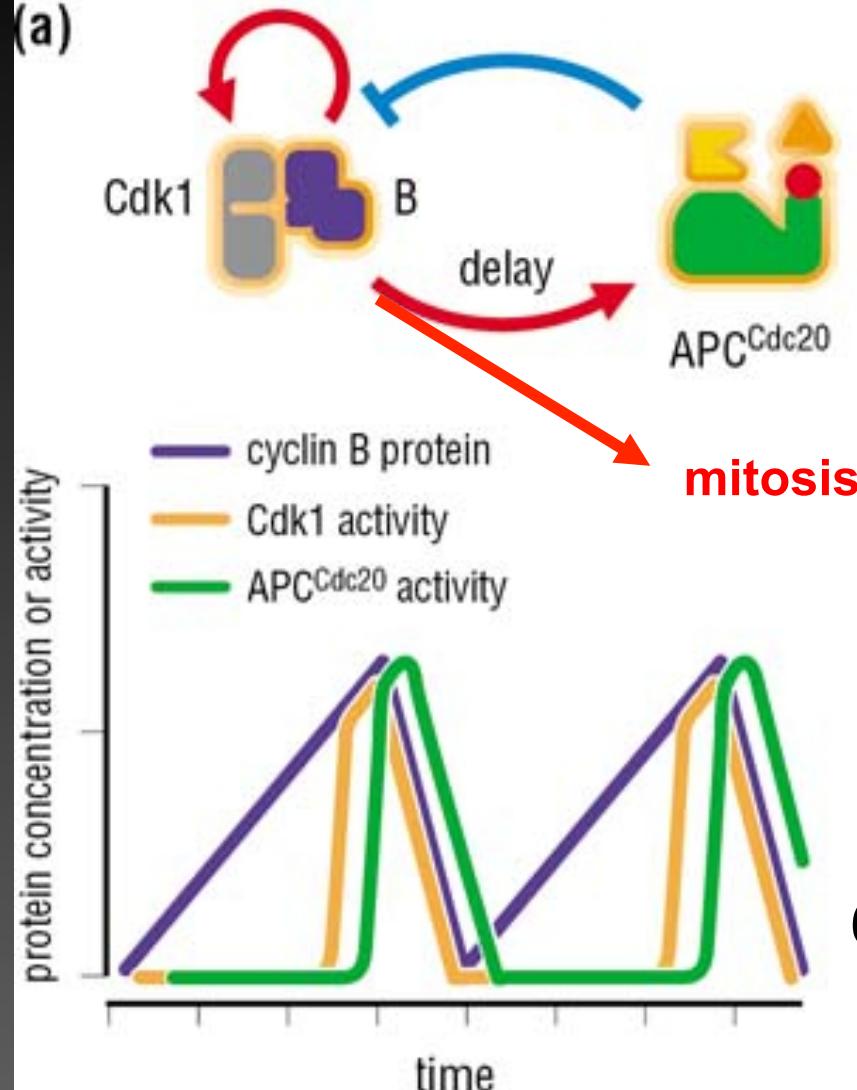
A simple cell-cycle oscillator model:

embryonic systems

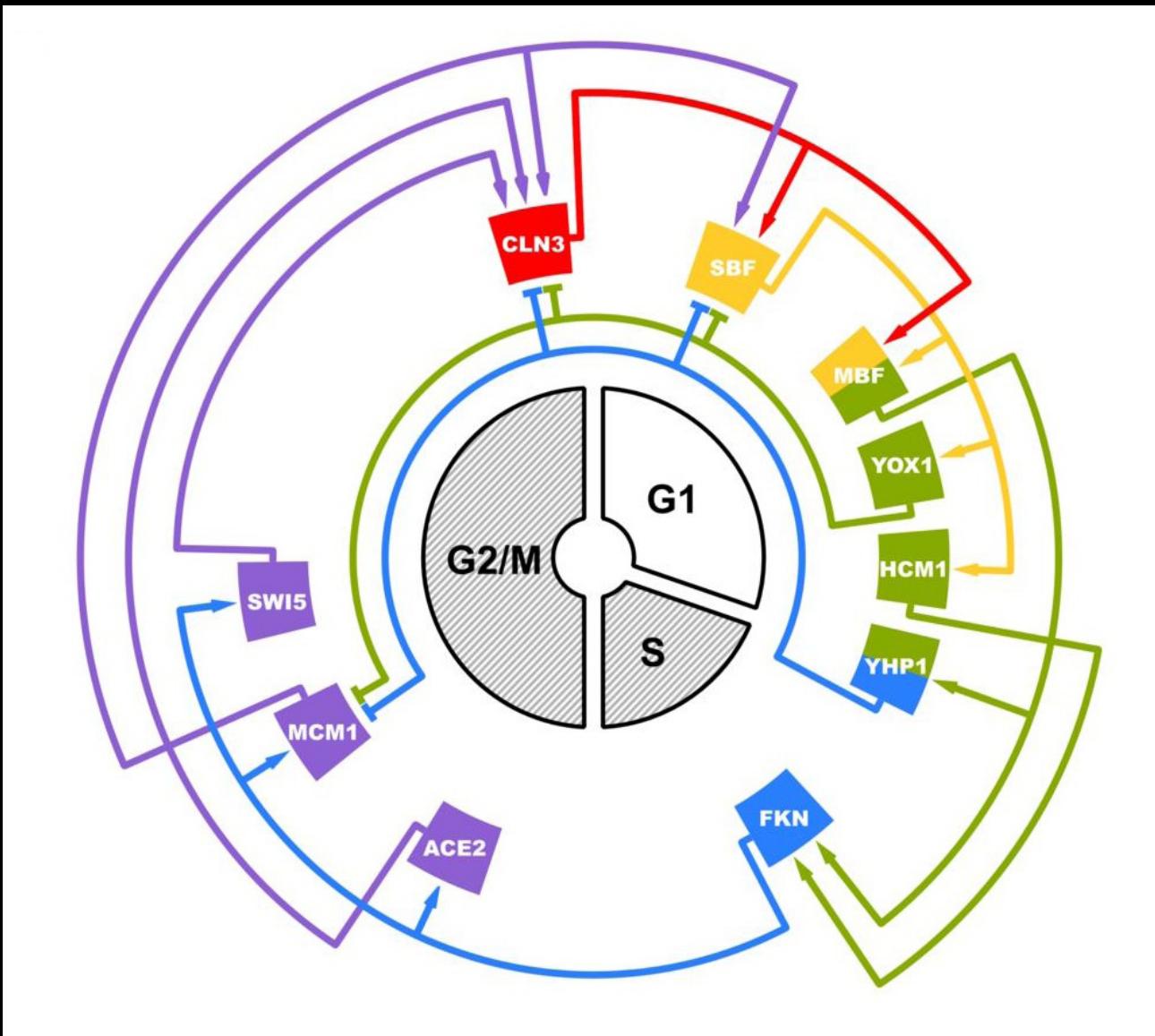


From [The Cell Cycle: Principles of Control](#)
by David O Morgan

(a)

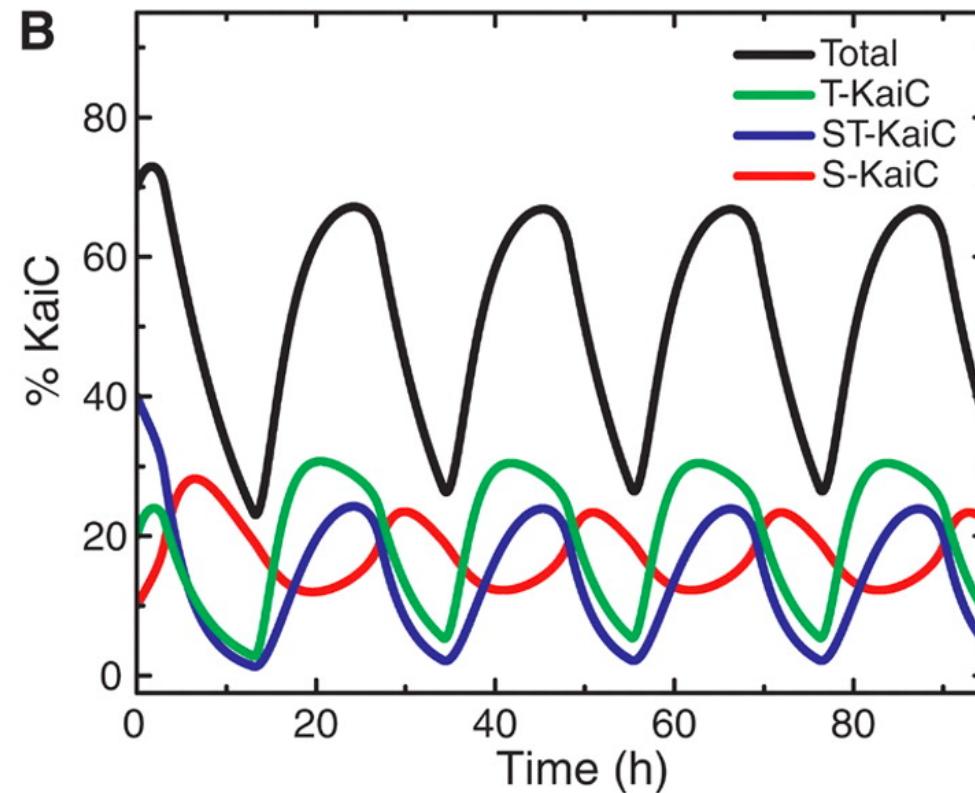
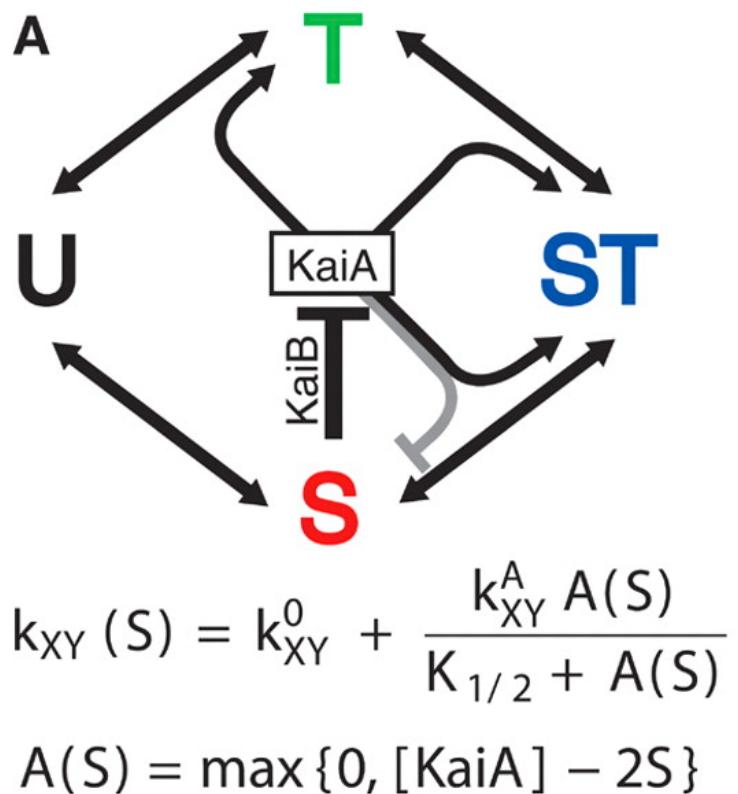


A synchronously updating Boolean model



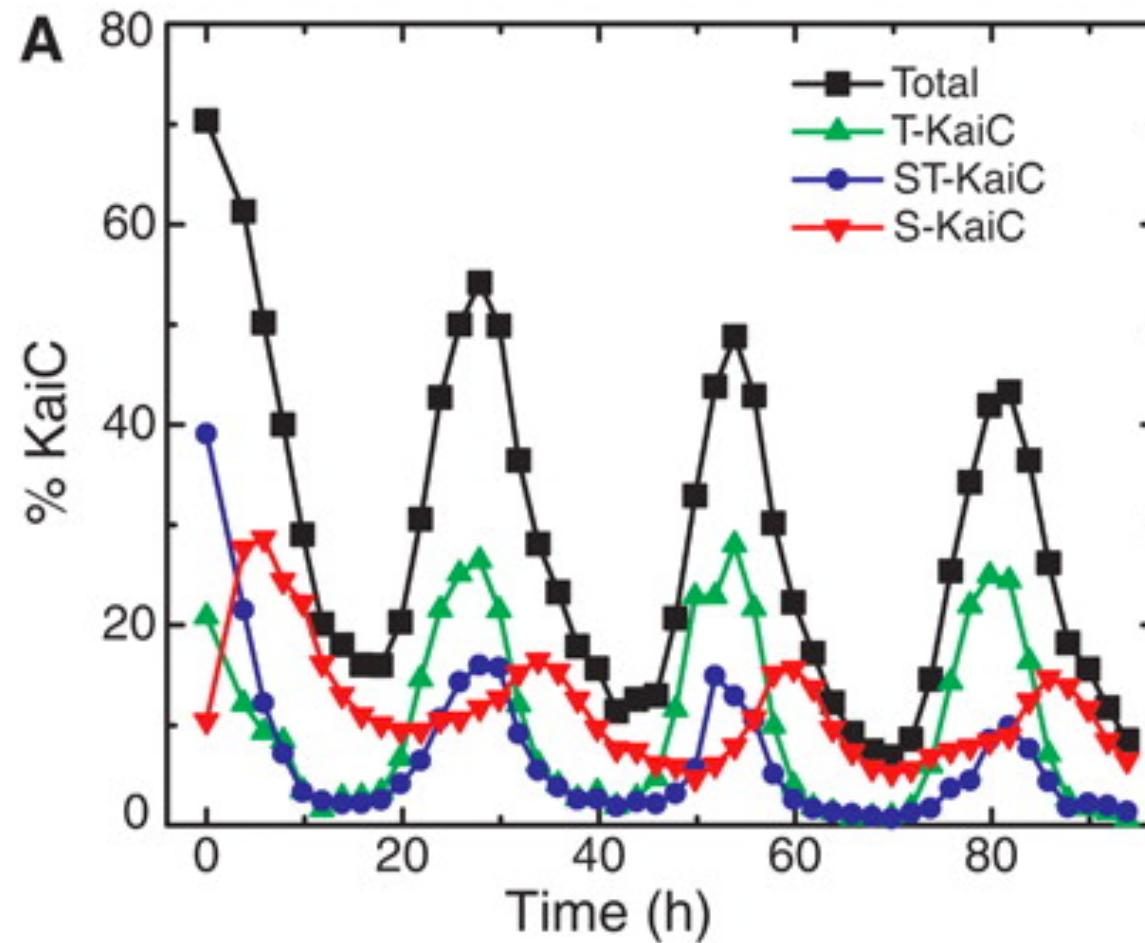
6.7

Transcription networks produce oscillations in _____ to regulate _____



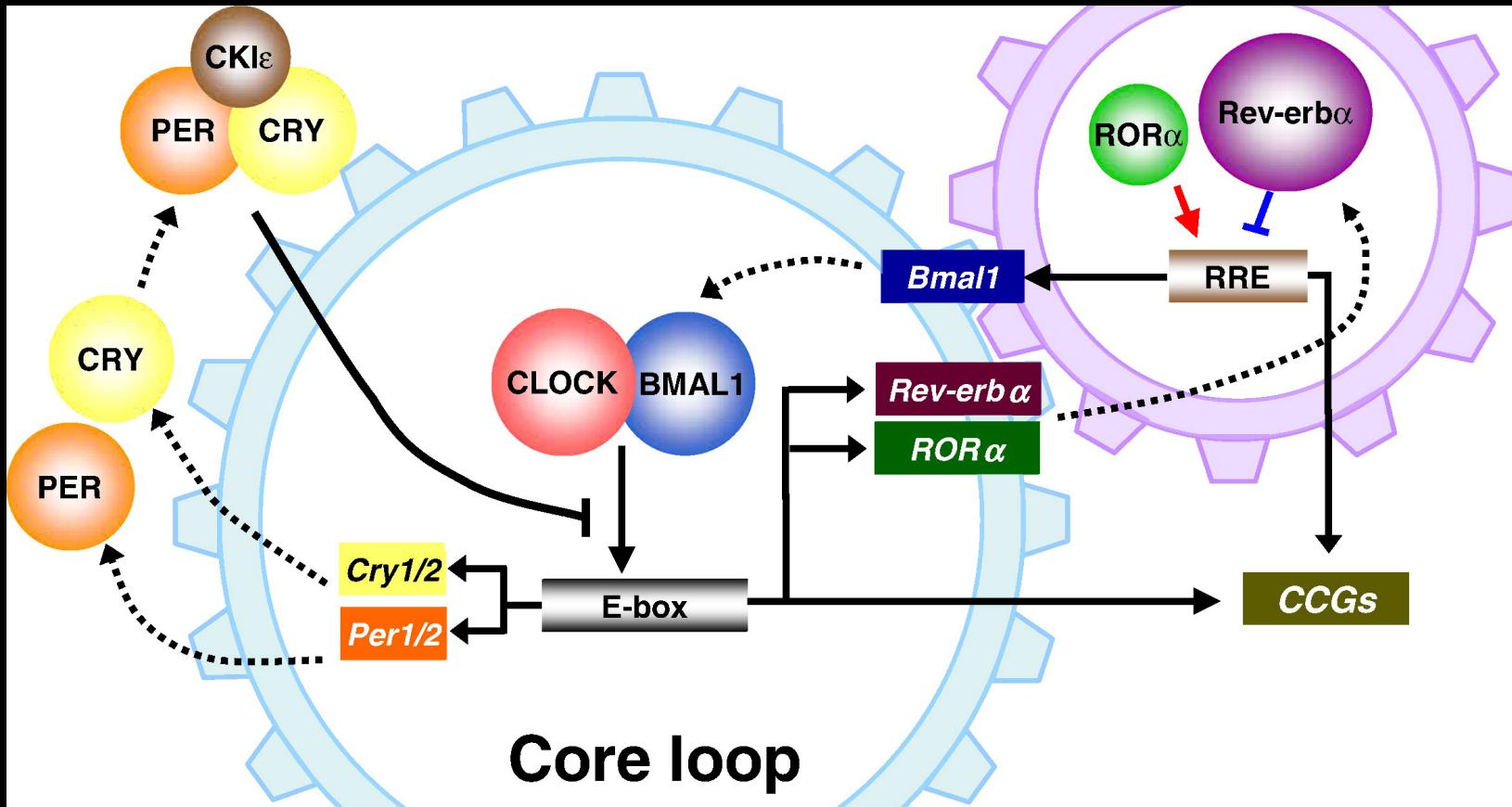
Rust et al.,
Science 2 November 2007:
Vol. 318 no. 5851 pp. 809–812
DOI: 10.1126/science.1148596

Oscillation requires phosphorylation control



Rust et al.,
Science 2 November 2007:
Vol. 318 no. 5851 pp. 809–812
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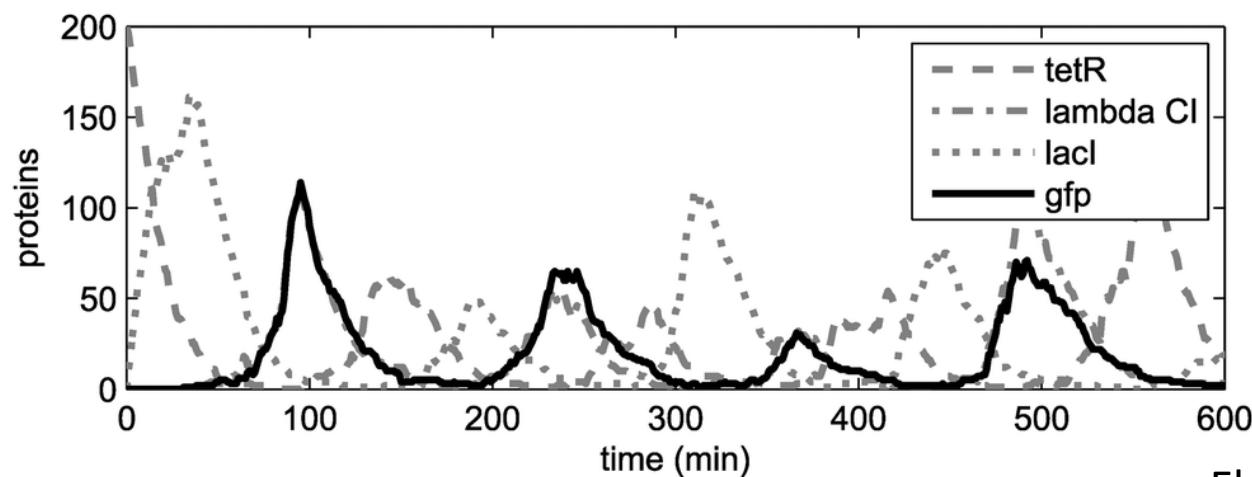
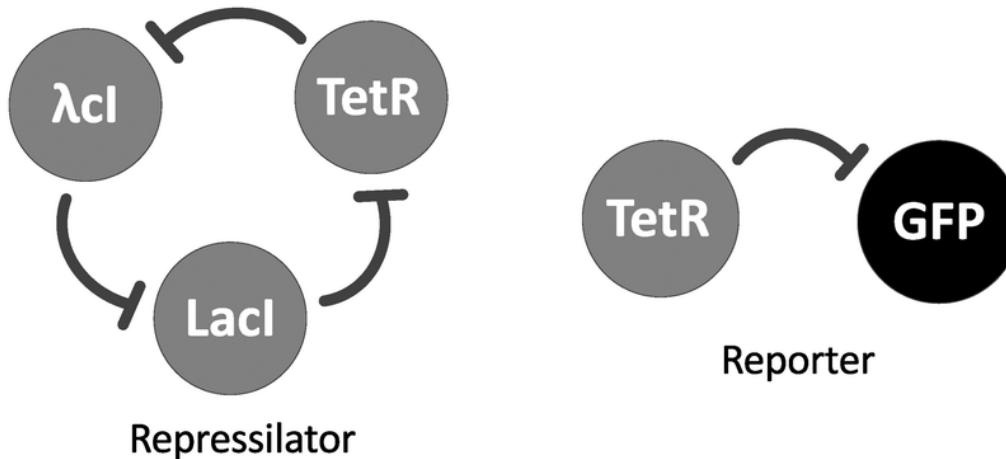
Circadian network



6.8

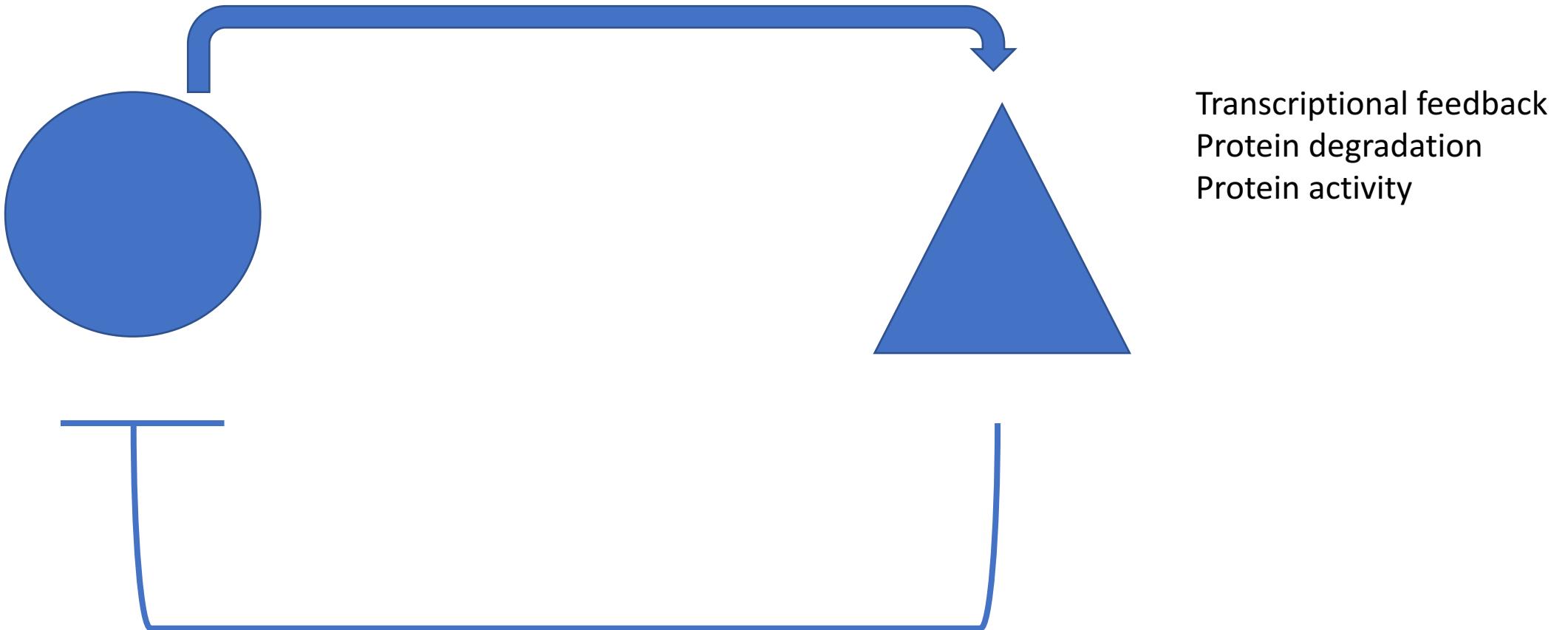
http://www.sciencedirect.com/cache/MiamiImageURL/1-s2.0-S0925443911000329-gr2_lrg.jpg/0?wchp=dGLbVIB-zSkWA&pii=S0925443911000329

A synthetic TRN: the Repressilator



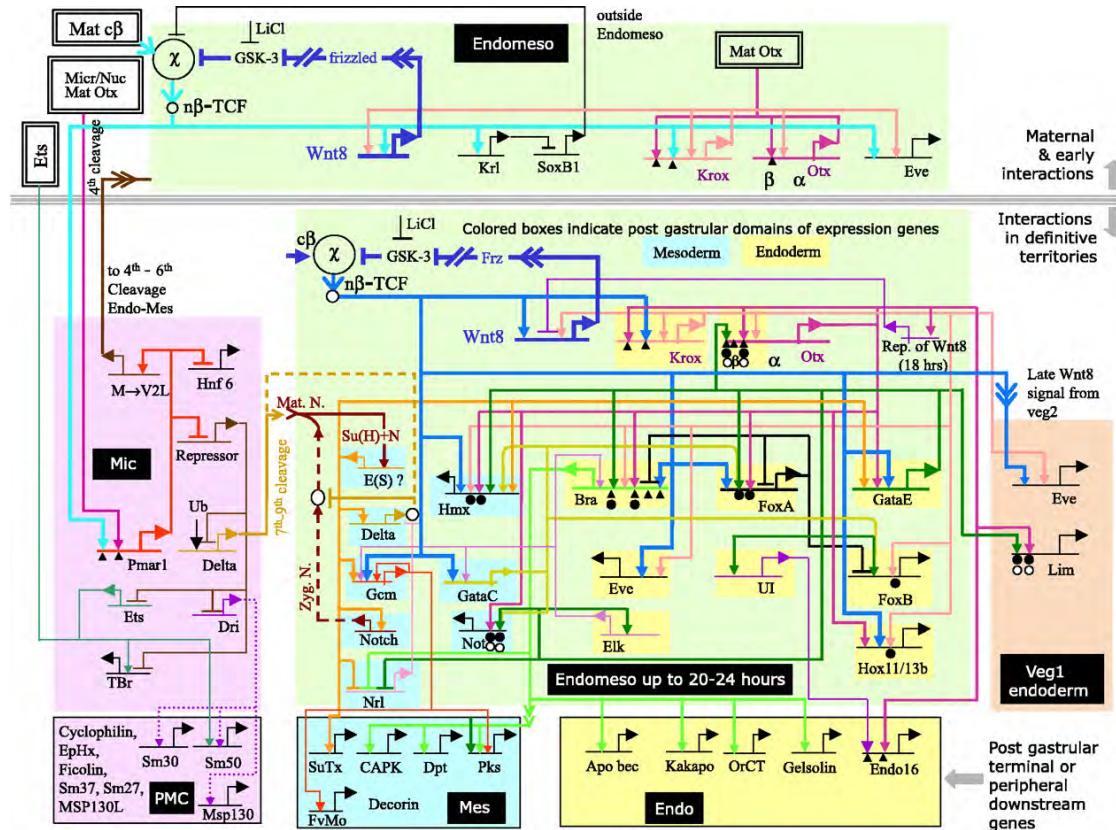
Elowitz et al., 2000. Nature. 403(6767):335-8.
Oliveira et al., 2015. Mol Biosyst. 11:1939-1945

Oscillations require negative feedback with delay



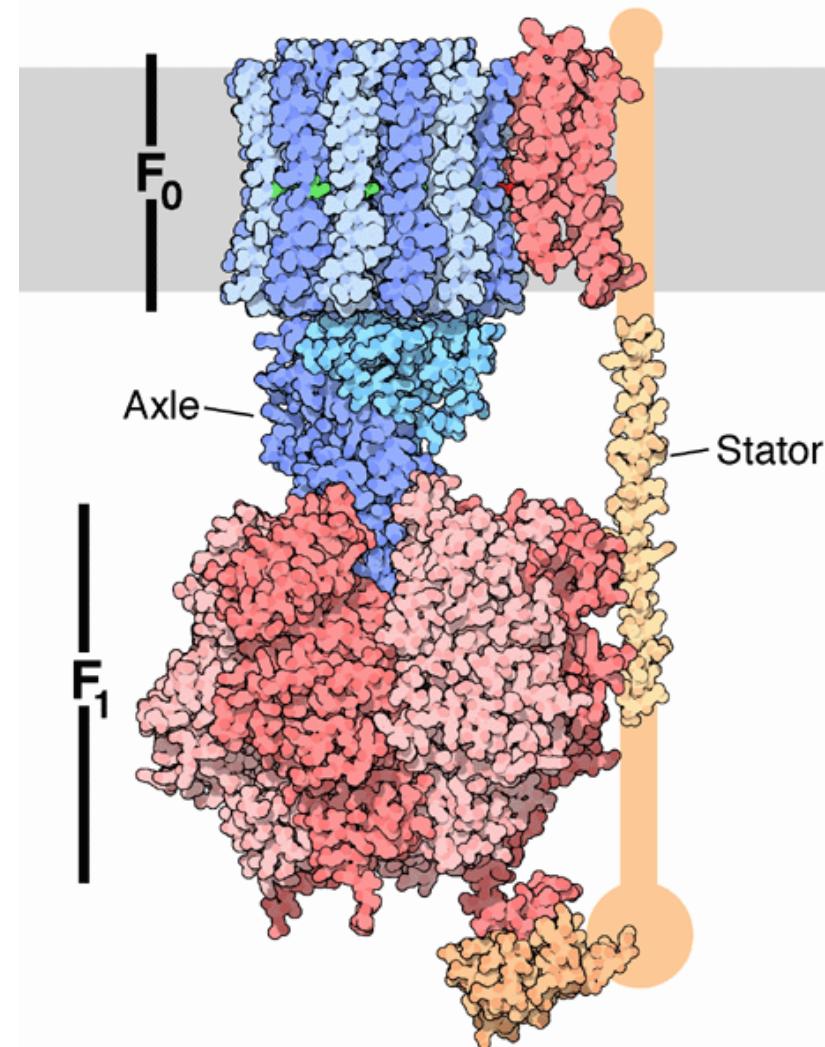
Development: Ordered hierarchy

Sea urchin embryogenesis



Protein-protein interactions

- Describes how all the proteins in the cell are connected together to make functional complexes.



Protein-protein interaction networks

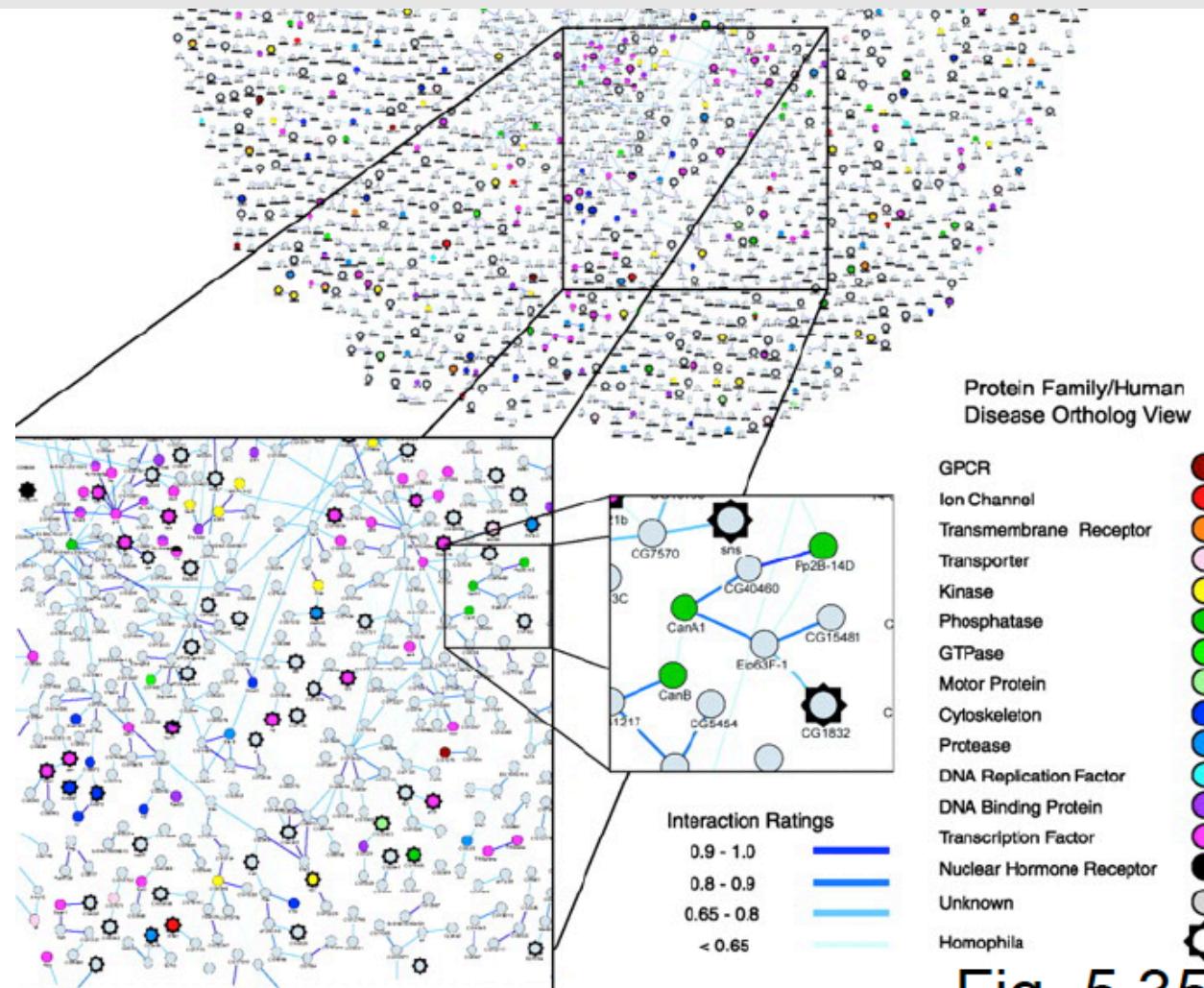
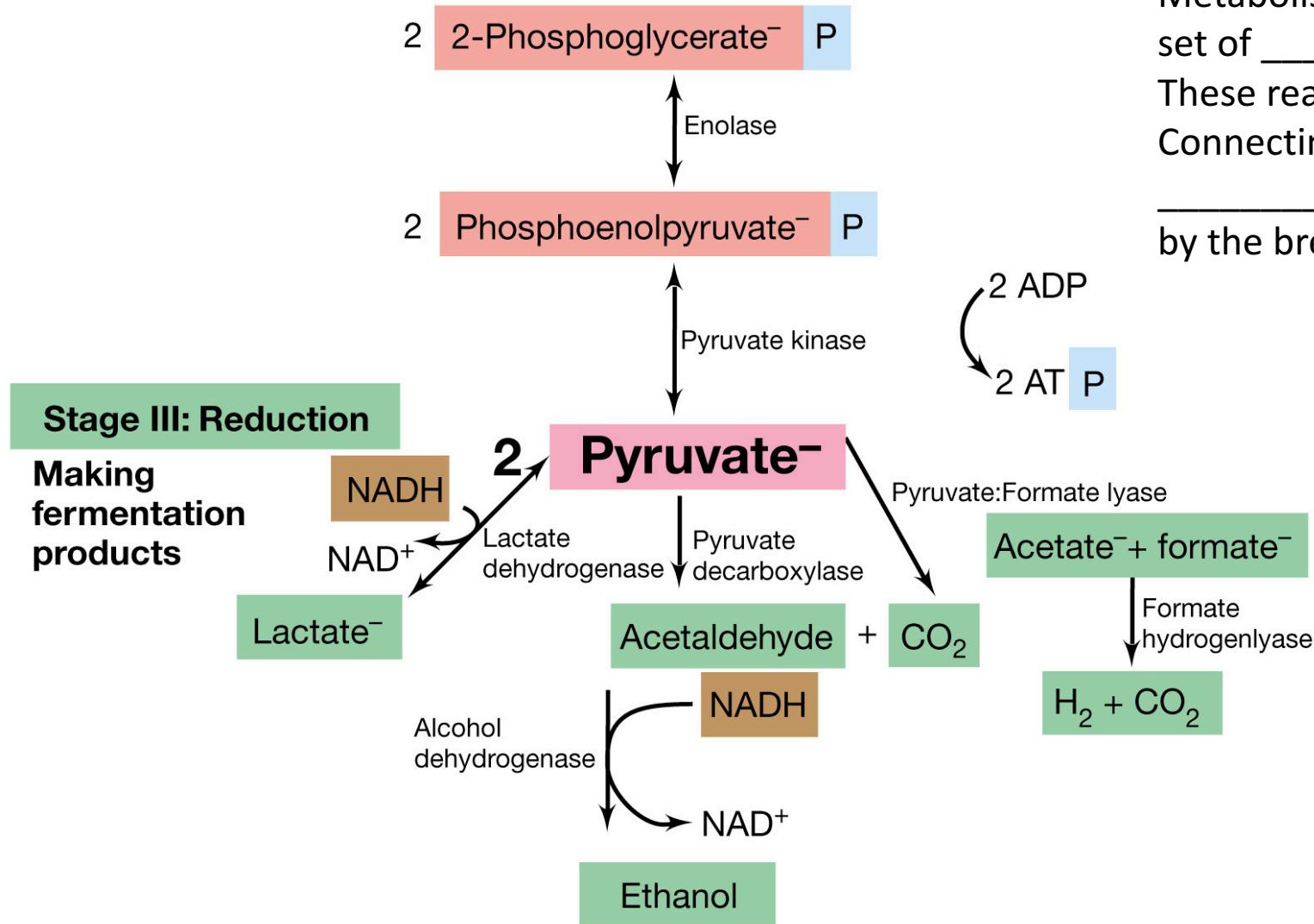


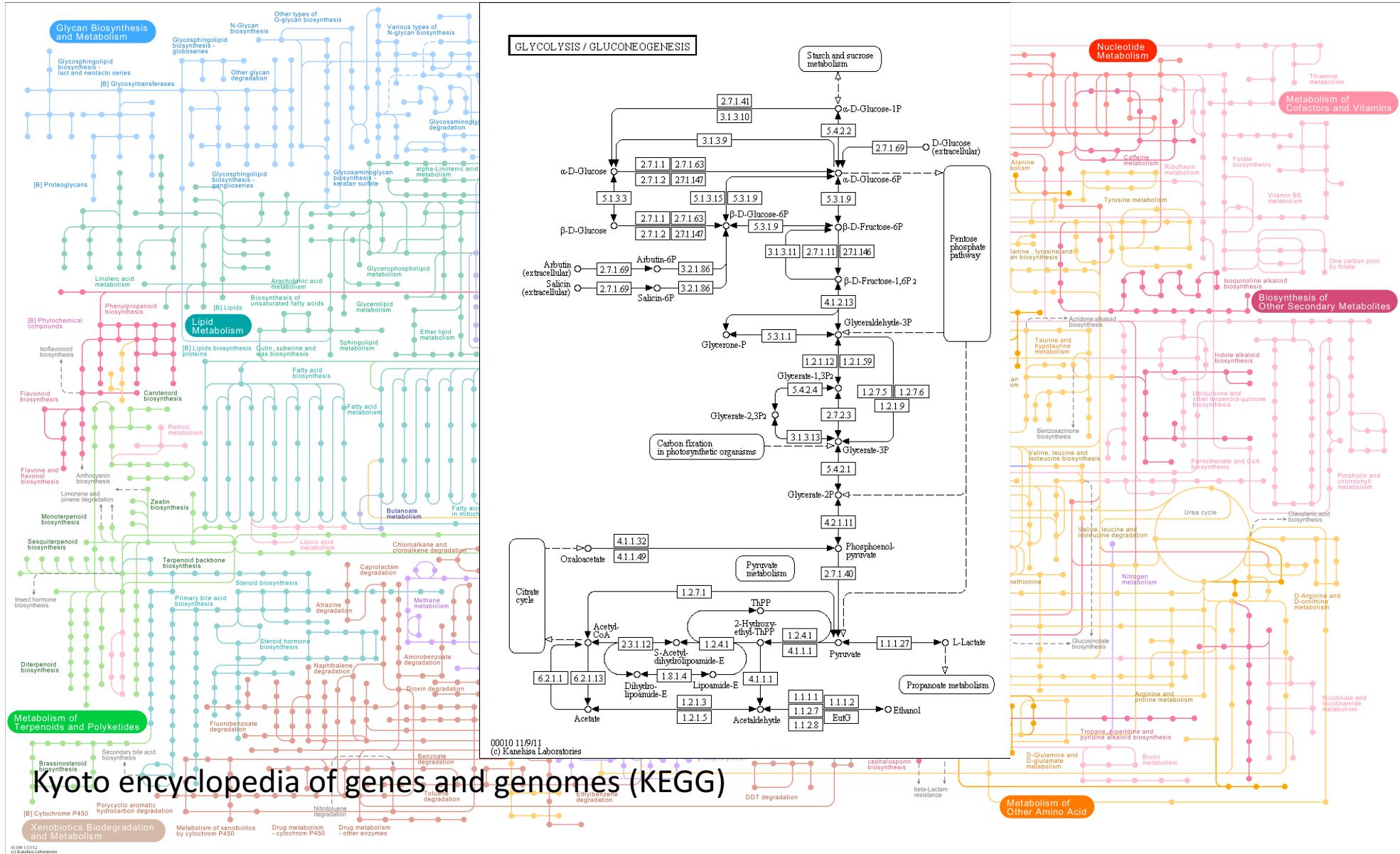
Fig. 5.35

Metabolism



Metabolism is defined as the set of _____ in the cell. These reactions can _____ by connecting small molecules together or _____ by the breakdown of small molecules.

Metabolic network –



Metabolic networks

- Describe chemical reactions for the production (anabolism) and breakdown (catabolism) of building blocks.
 - Amino acids
 - Sugars
 - Lipids
- Enzymes and energy required for synthesis and the processes by which the building blocks combine to make larger molecules
- Energy is released by catabolic reactions
- Metabolite profiling combined with gene expression analysis identifies how the TRN and metabolic networks are intercoordinated.