



GenPCCA: Markov State Models for Non-Equilibrium Steady States

M. Weber*, K. Fackeldey



Zuse Institute Berlin (ZIB)
Computational Molecular Design
<http://www.zib.de/weber>

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joint work with: S. Röblitz, C. Schütte, L. Reinmiedl, RG A. Gleixner,
N. Djurdjevac-Conrad

stochastic process \rightarrow finite Markov chain with transition matrix P

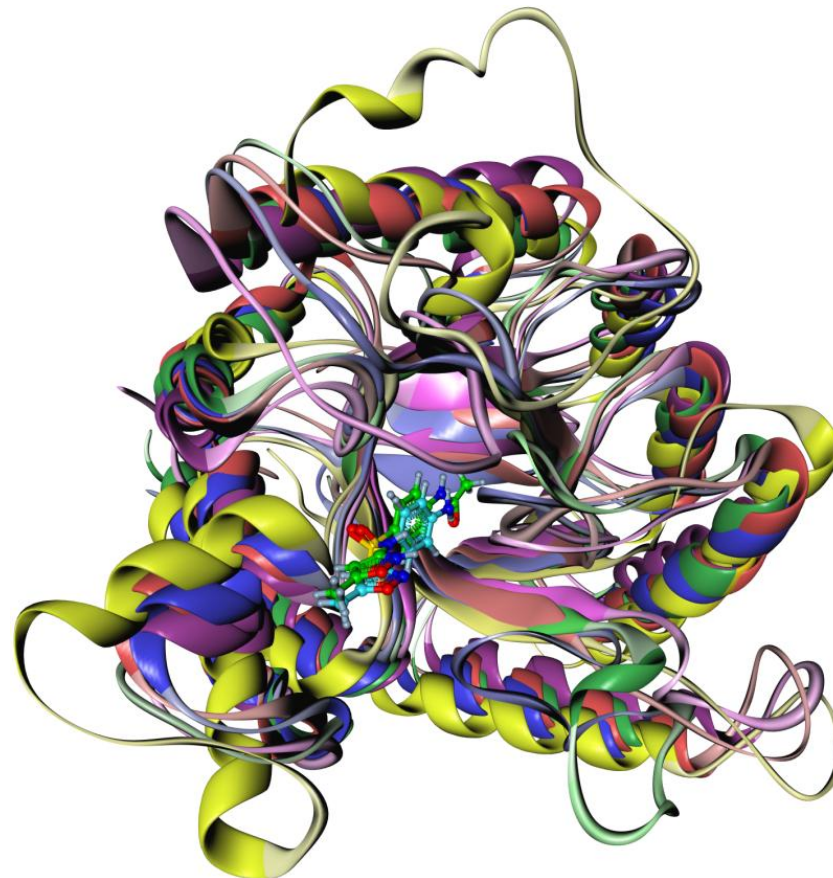
1) Cyclic behaviour in molecular processes

2) Robust Perron Cluster Analysis (PCCA+)

3) From Eigendecompositions to Schur Decompositions

4) Generalized PCCA for Dominant Cycles
(function-based clustering)

5) Example



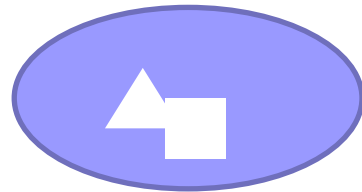
optimizing dihydropterat-synthetase

dihydropterat-synthetase (catalytic reaction)

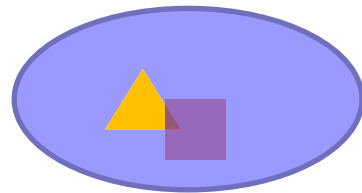
DHPP



pABA

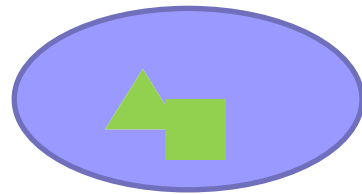


dihydropterat-synthetase
(catalytic reaction)



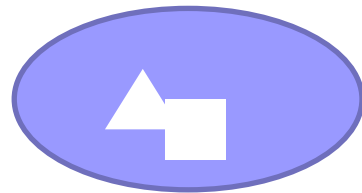
complex

dihydropterat-synthetase
(catalytic reaction)

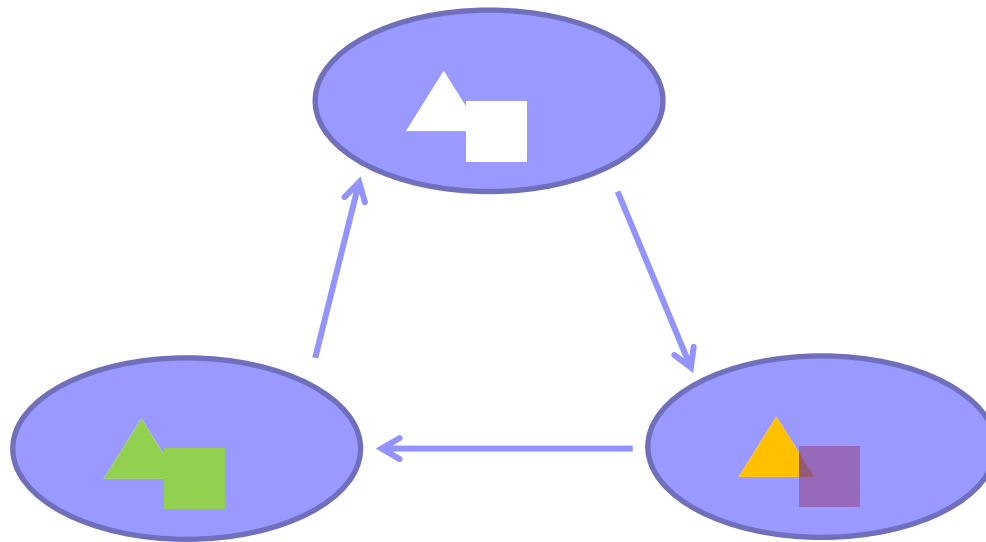


complex

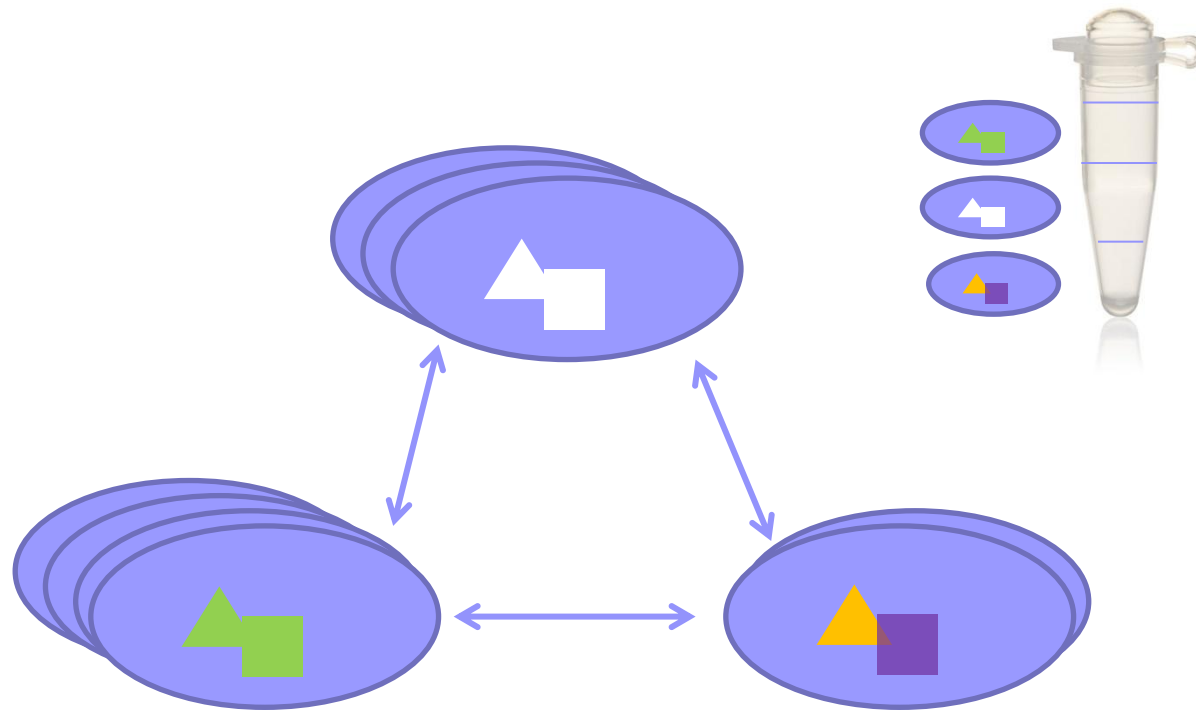
dihydropterat-synthetase
(catalytic reaction)



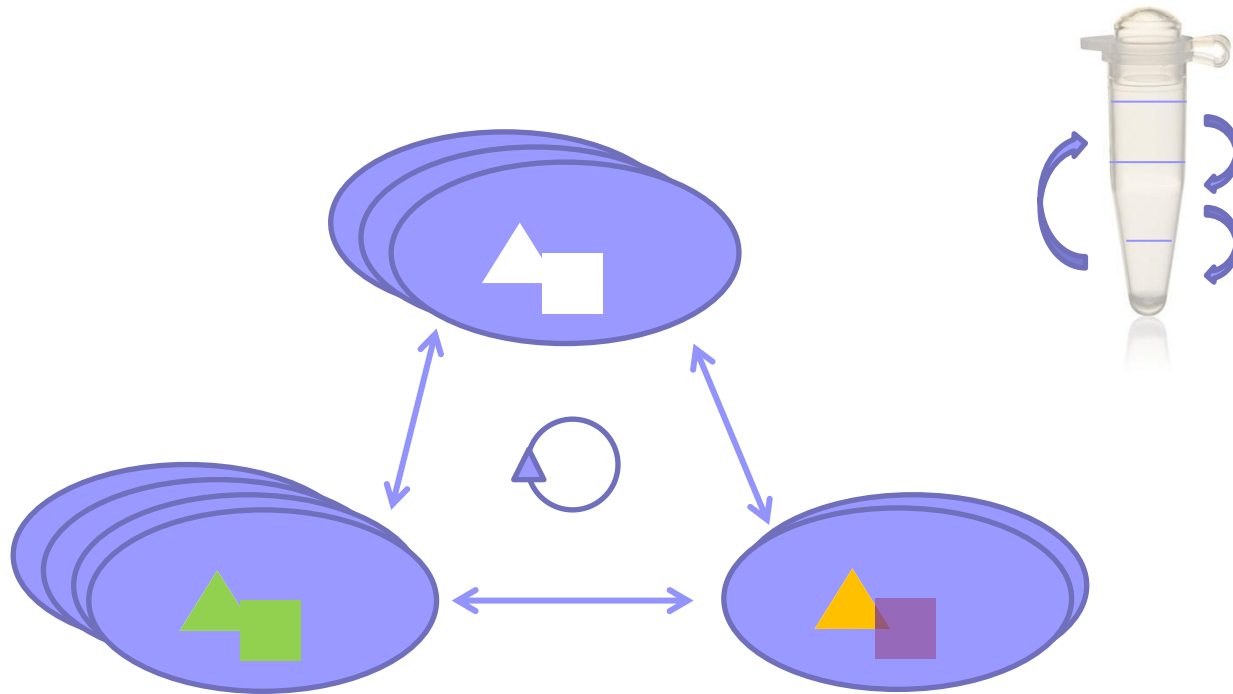
product
(-> acid folique)



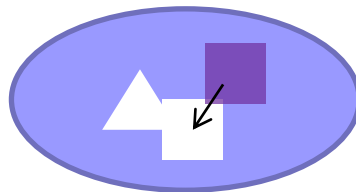
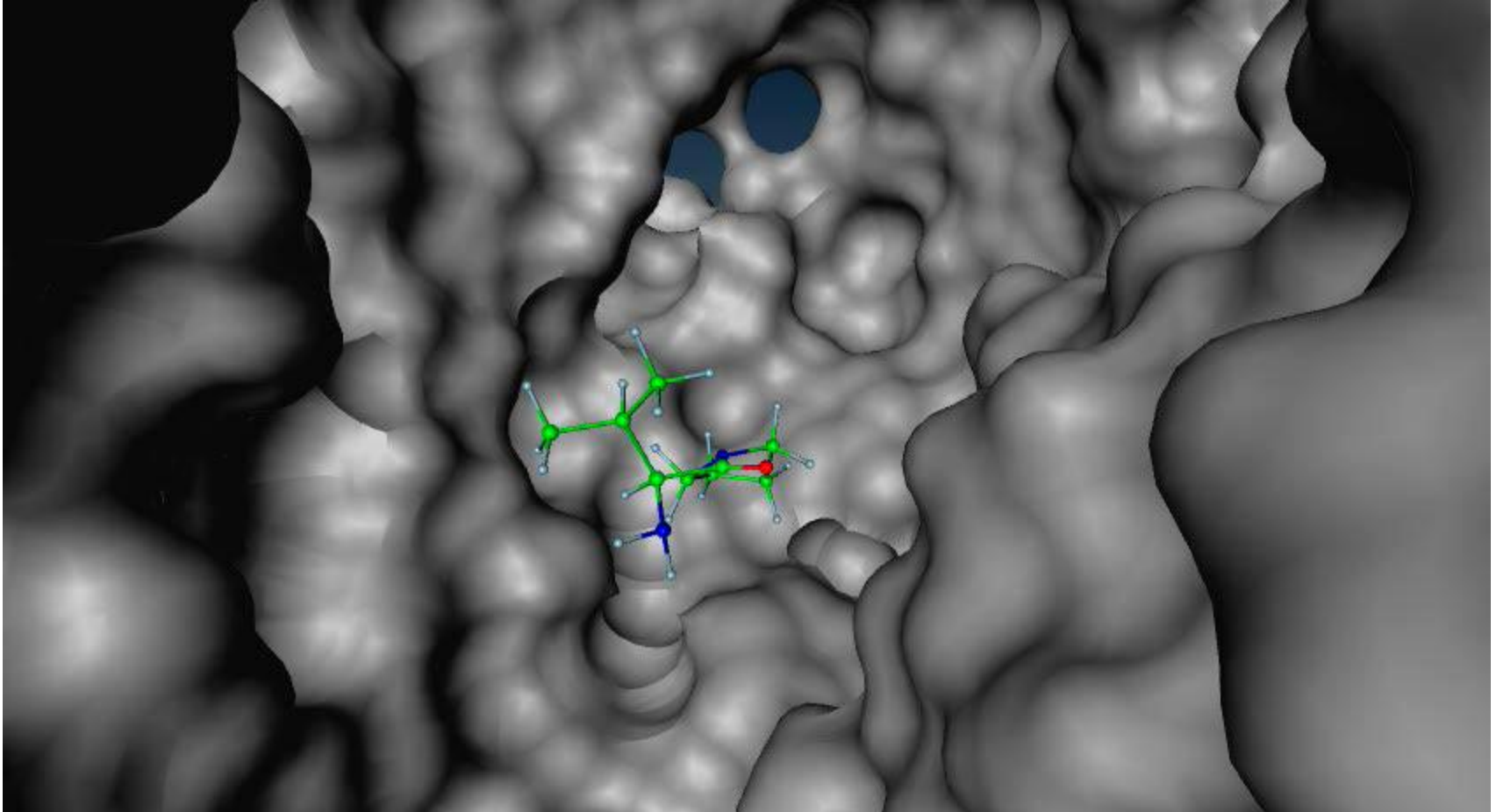
catalytic cycle



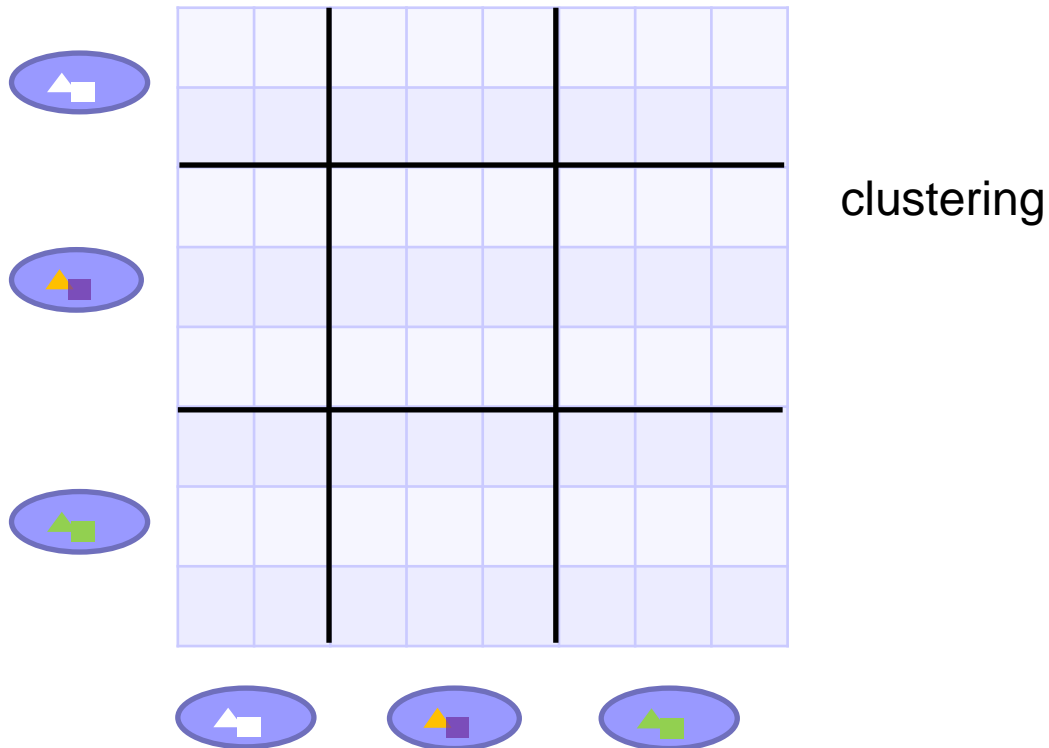
stationary distribution



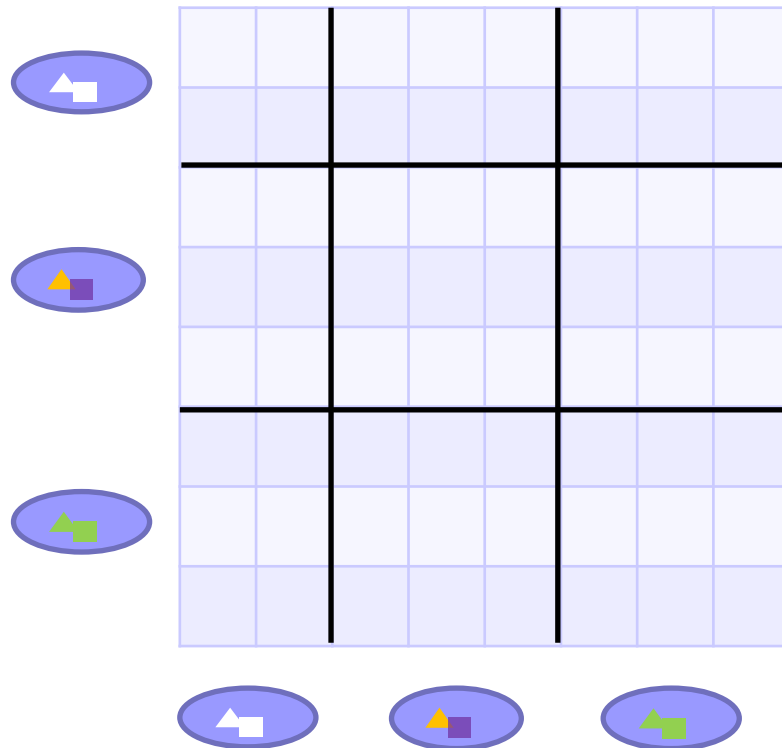
NESS
(non-equilibrium steady state)



detailed molecular description
of the system (transitions P)

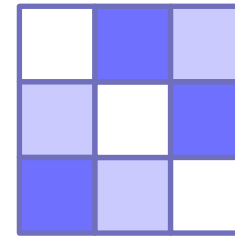


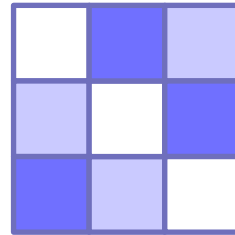
detailed molecular description
of the system (transitions P)



clustering/
projection
→

efficiency of
catalytic cycle

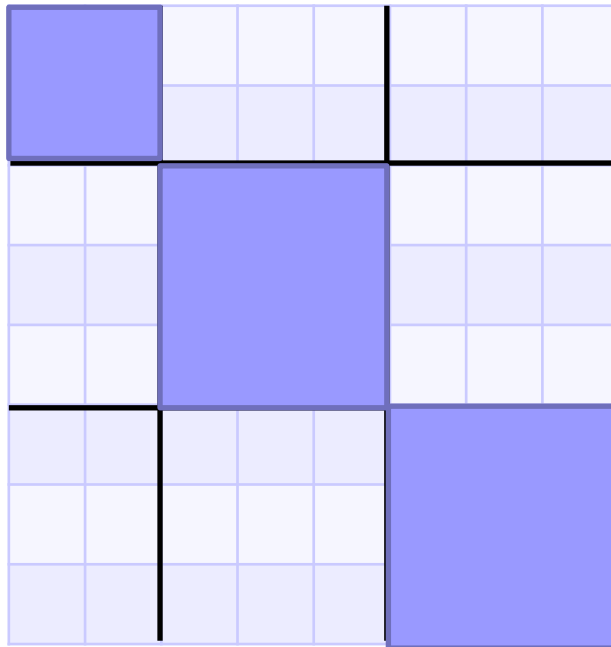




efficiency = non-reversibility of P_C

- 1) Cyclic behaviour in molecular processes
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- 5) Example

P



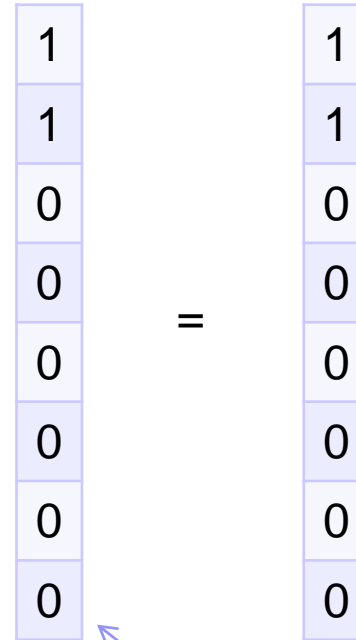
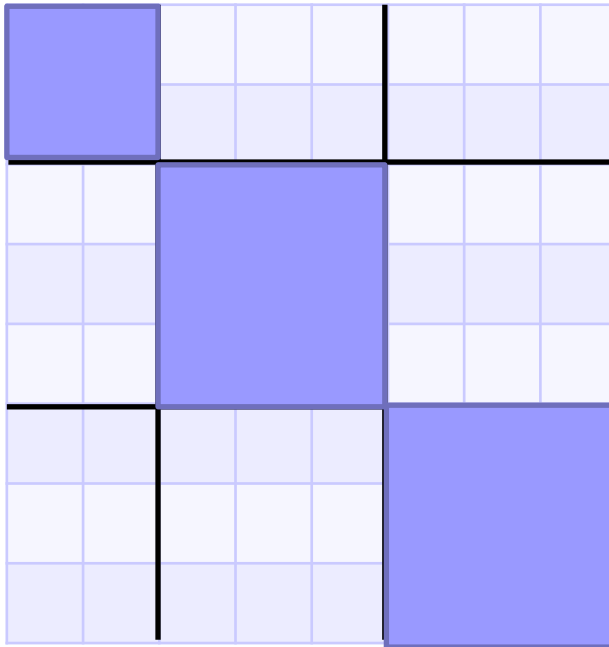
1
1
0
0
0
0
0
0

=

1
1
0
0
0
0
0
0

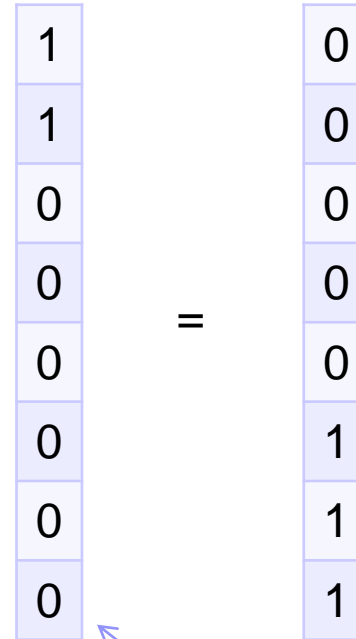
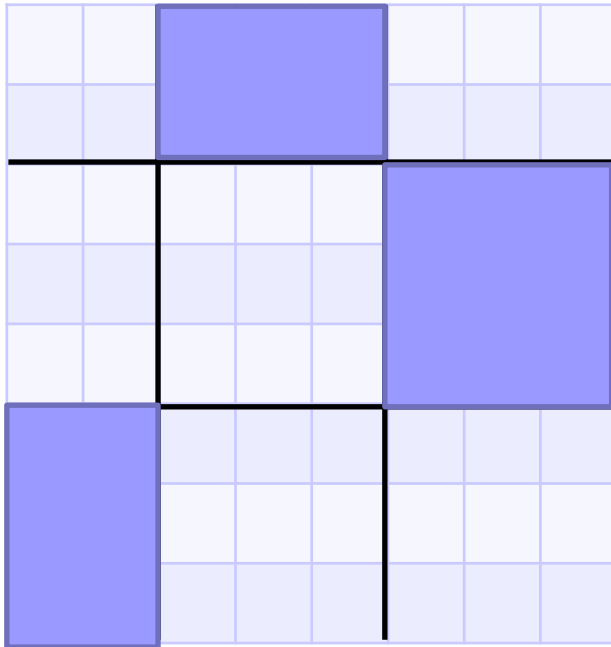
Weber, Galliat, 2002
Deuffhard, Weber, 2005
Weber, 2006
Röblitz, Weber, 2013

P

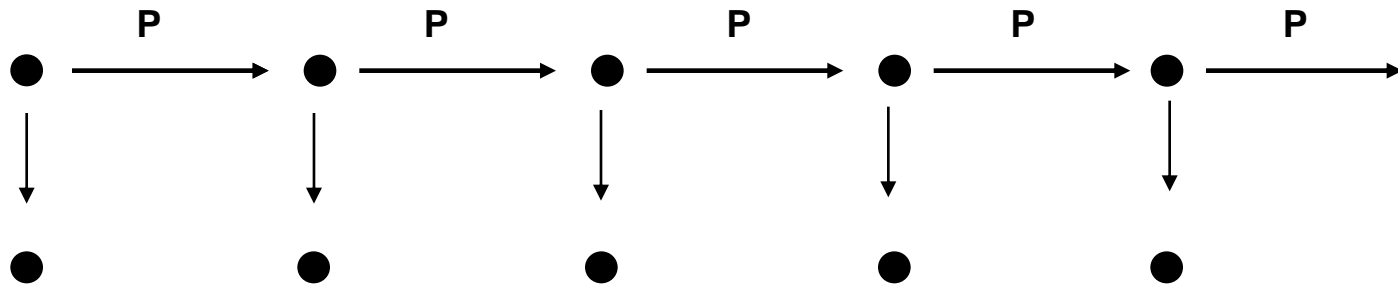


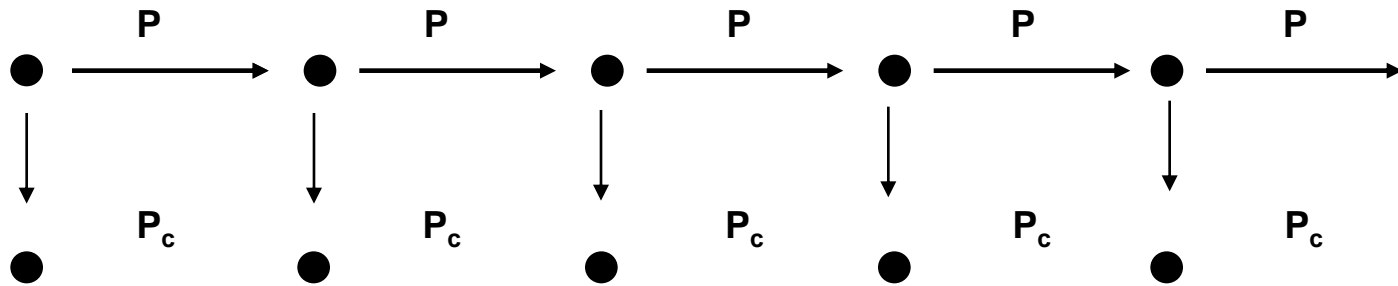
Probability for a process starting here for ending up in that set in 1 step

P

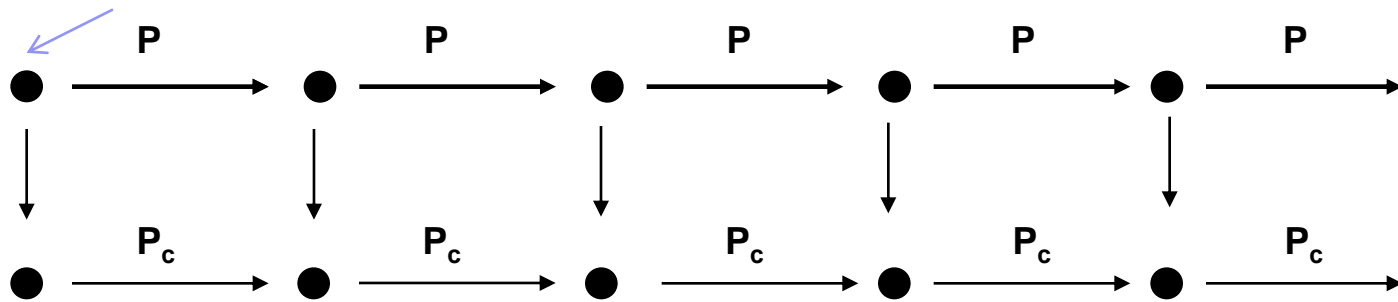


Probability for a process starting here for ending up in that set in 1 step





membership vectors




linear coefficients of the membership vectors

- P_c is a Galerkin discretization of P
- Membership vectors form an invariant subspace X of P
- starting point of P -chain inside the invariant subspace

Weber, 2011

Röblitz, Weber, 2013


Fackeldey, Weber, 2016


$$PX = X\Lambda$$

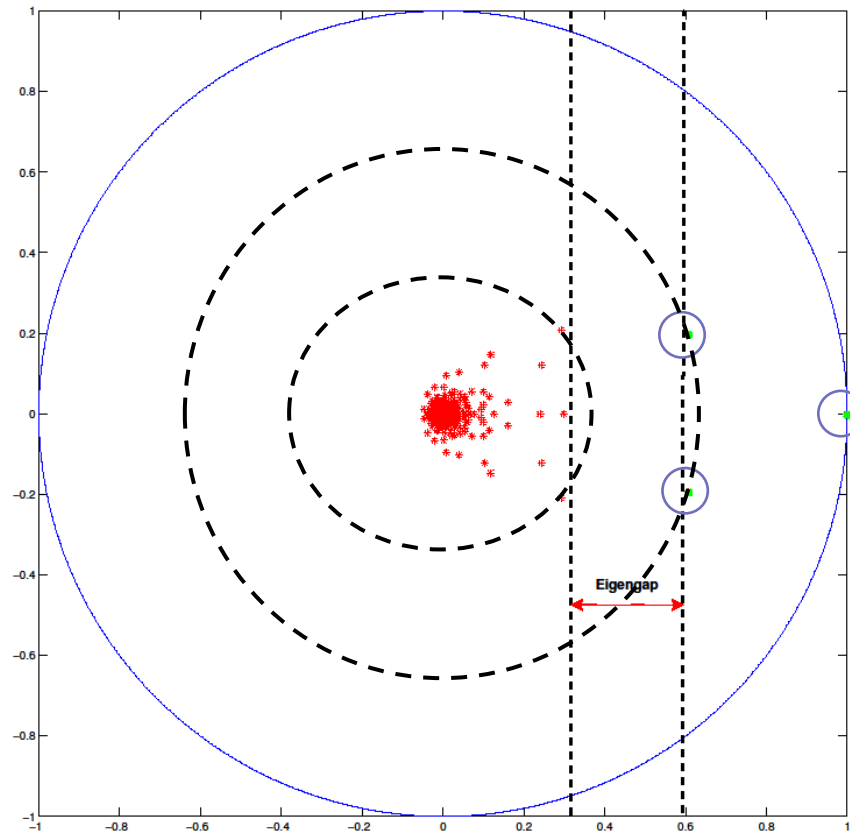
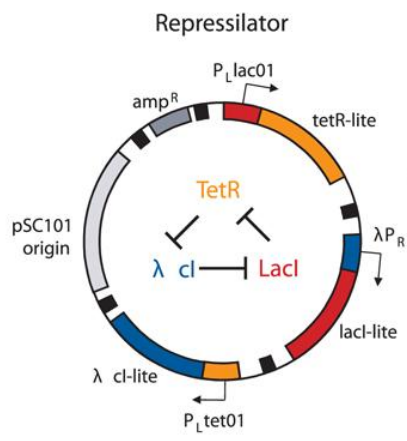
$$\chi = XA$$


X orthogonal matrix with regard to the stationary distribution (separability = orthogonality of A)

- 1) Cyclic behaviour in molecular processes
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$$PX = X\Lambda$$

X orthogonal matrix with regard to the stationary distribution




$$PX = X\Lambda$$

diagonal matrix??

X orthogonal matrix with regard to the stationary distribution

Schur
decomposition

$$PX = X\Lambda$$

+	+	+	+	+	+	+	+
	+	+	+	+	+	+	+
		+	+	+	+	+	+
			+	+	+	+	+
				+	+	+	+
					+	+	+
						+	+
							+

Schur
decomposition

$$PX = X\Lambda$$

+	+	+	+	+	+	+	+
	+	+	+	+	+	+	+
		+	+	+	+	+	+
			+	+	+	+	+
				+	+	+	+
					+	+	+
						+	+
							+

real Schur
decomposition

$$PX = X\Lambda$$

+	+	+	+	+	+	+	+
	+	+	+	+	+	+	+
		+	+	+	+	+	+
		+	+	+	+	+	+
				+	+	+	+
					+	+	+
						+	+
							+

real Schur
decomposition

$$PX = X\Lambda$$

+	+	+	+	+	+	+	+
	+	+	+	+	+	+	+
		+	+	+	+	+	+
		+	+	+	+	+	+
				+	+	+	+
					+	+	+
						+	+
							+

„Schur = Eigen“ for reversible matrices

Schur is well-conditioned

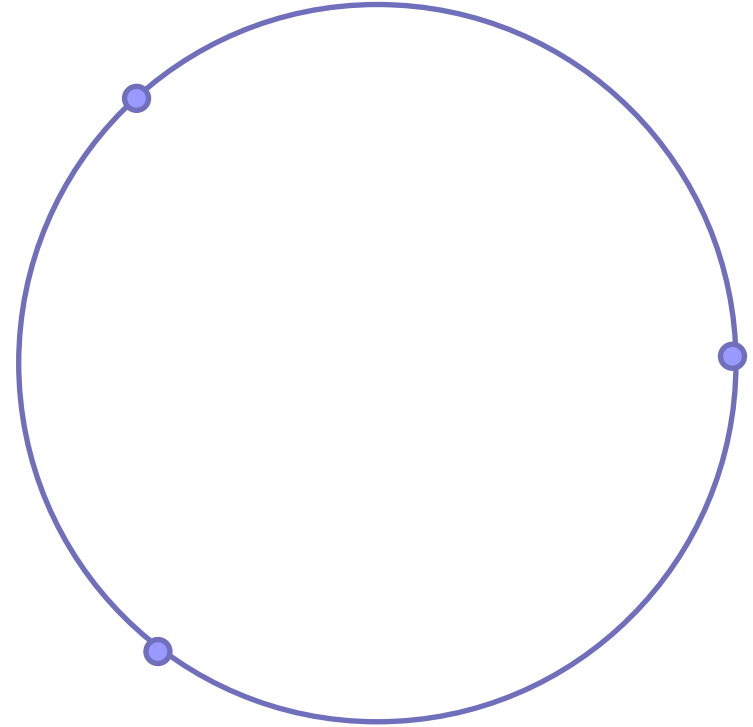
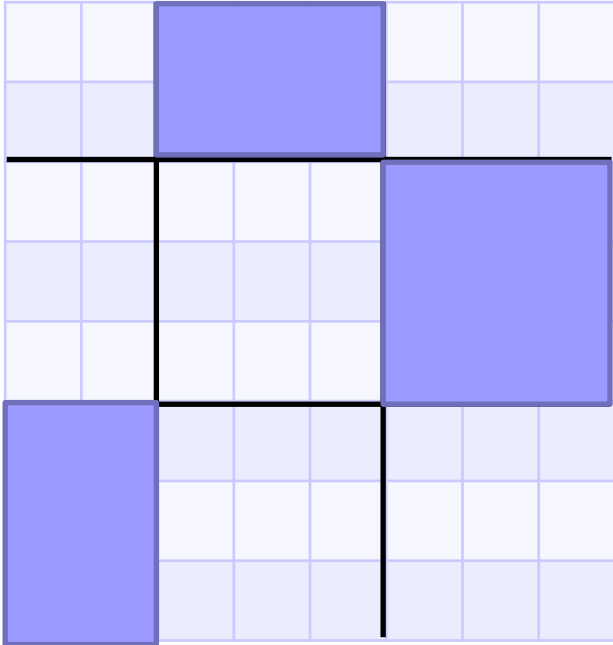
Schur always exists

➔ Schur is not unique

Schur values of P become Schur values of P_C

$$T = \begin{pmatrix} 5/12 & 5/12 & 1/6 \\ 1/4 & 1/4 & 1/2 \\ 1/3 & 1/3 & 1/3 \end{pmatrix}$$

P



P_C

0	1	0
0	0	1
1	0	0

Non-reversibility $n = \|DP_C - (DP_C)^T\|_{F,\mu}$ of P_C is a consequence of the non-diagonality of the Schur matrix and the „separability / orthogonality“ of the clusters.

$$n = \sqrt{2 \sum_{i>j} |\Lambda_{ij} - \Lambda_{ji}| \cdot \|A^{(i)} \times A^{(j)}\|_{\mu}^2}$$

Djurdjevac-Conrad, Schütte, Weber, 2016

Λ

+	+	+	+
	+	+	+
		+	+
		+	+

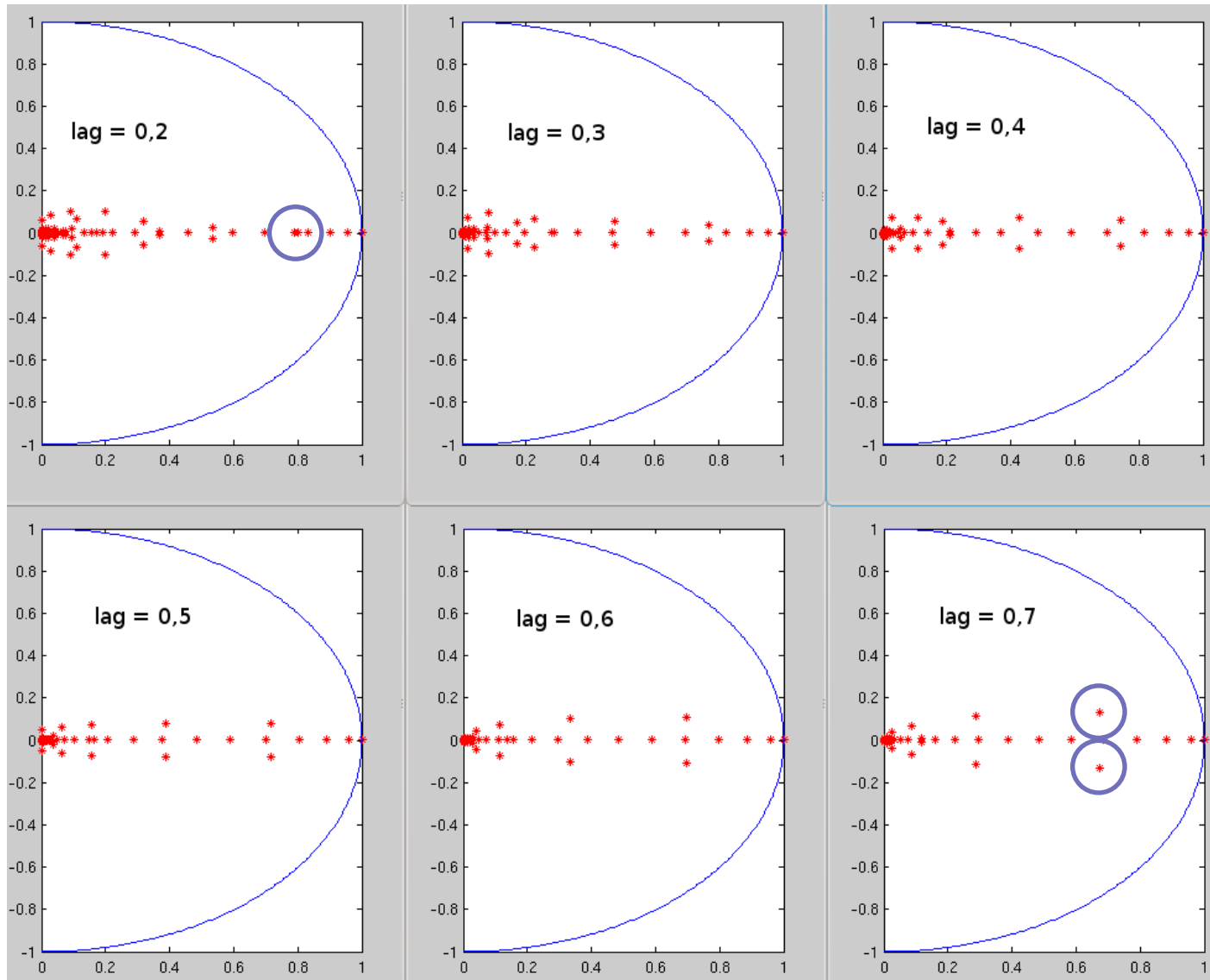
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Djurdjevac-Conrad, Schütte, Weber, 2016

Λ

+	+	+	+
	+	+	+
		+	+
		+	+



SDE Hindemarsch-Rose (neuronal excitation)

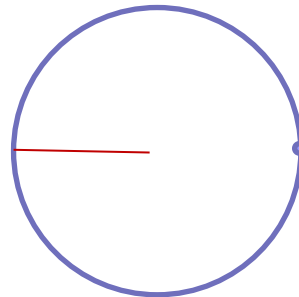
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Preparation of the invariant space:

```
Pd=diag(sqrt(sd))*P*diag(1./sqrt(sd));
[Q, R]=schur(Pd);
[Q, R]=SRSchur(Q, R);
X=diag(1./sqrt(sd))*Q;
```

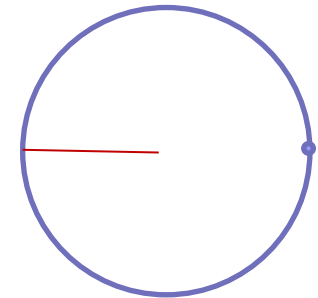
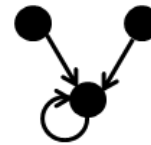
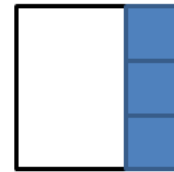
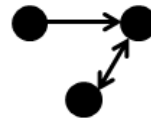
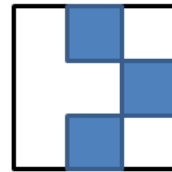
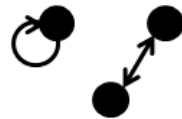
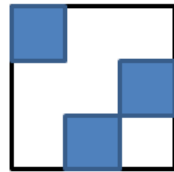
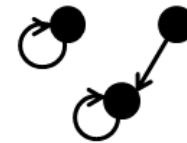
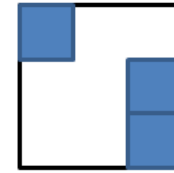
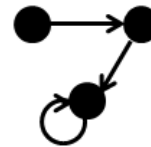
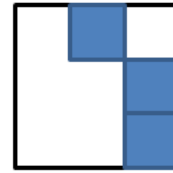
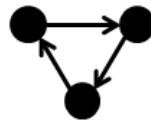
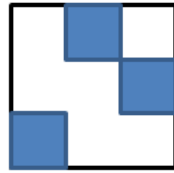
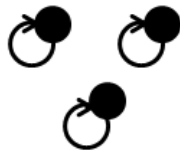
X orthogonal with regard to the stationary distribution
SRSchur sorts the eigenvalues

```
function [val,pos] = select(r)
%[val pos] = min(abs(1-r)); %Metastability
[val, pos]=max(abs(r)); %Permutation Matrices
%[val, pos]=max(abs(r-(real(r)<0).*real(r))); % LOG
%...
```



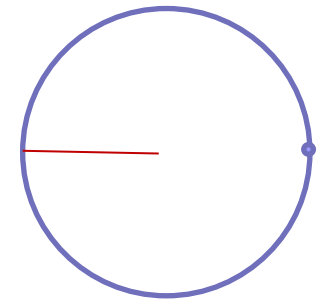
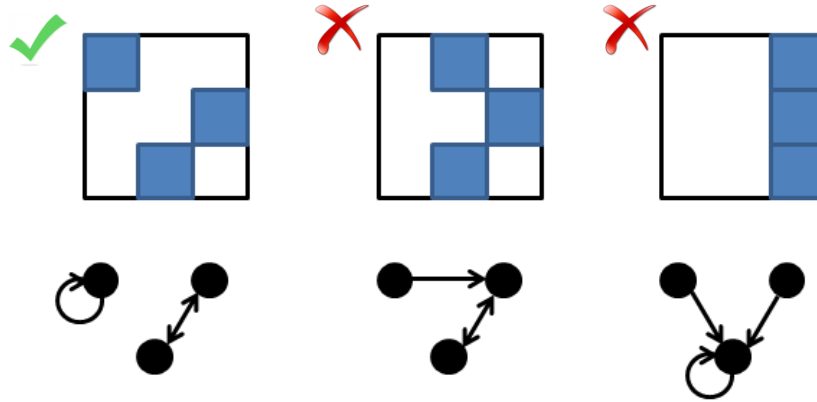
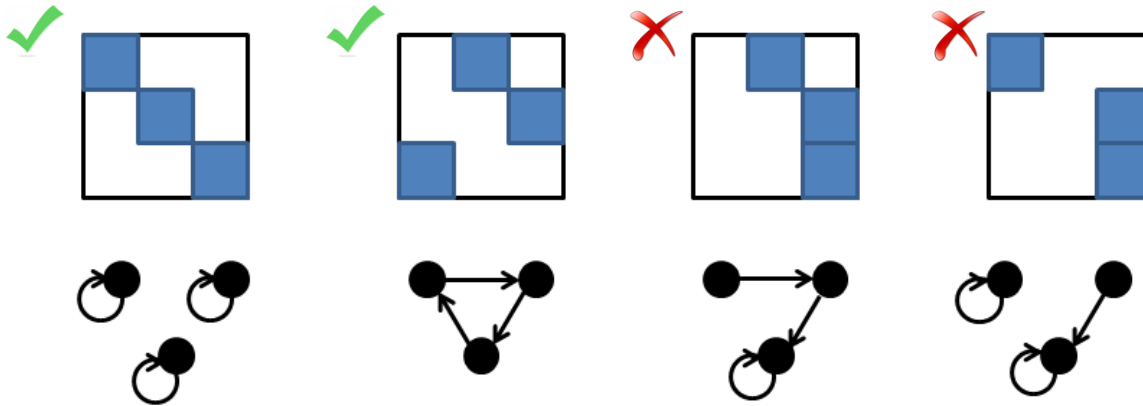
Why taking the absolute value?

sources/sinks = redundancy

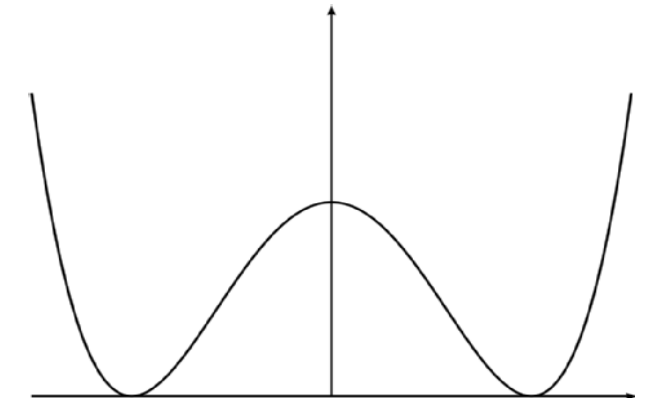
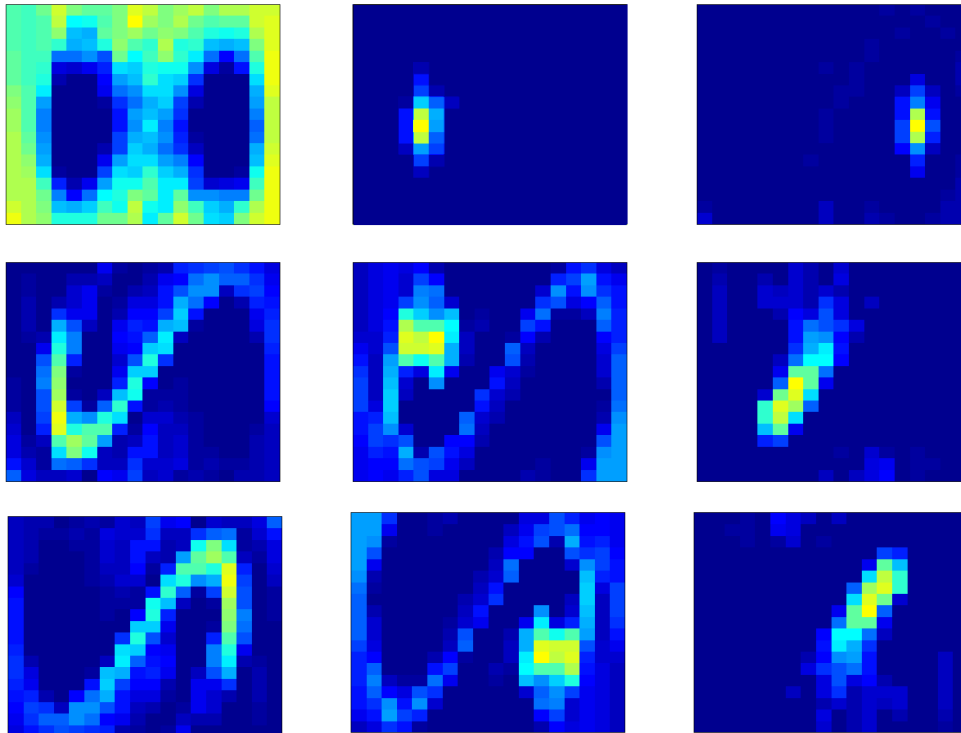


Why taking the absolute value?

sources/sinks = redundancy



momentum
↑
space →



low-friction Langevin dynamics

1	0	0	0	0	0	0	0	0
0	1	0	0	0	0	0	0	0
0	0	1	0	0	0	0	0	0
0	0	0	0	1	0	0	0	0
0	0	0	0	0	1	0	0	0
0	0	0	1	0	0	0	0	0
0	0	0	0	0	0	0	1	0
0	0	0	0	0	0	0	0	1
0	0	0	0	0	0	1	0	0

Djurdjevac-Conrad, Schütte, Weber, 2016

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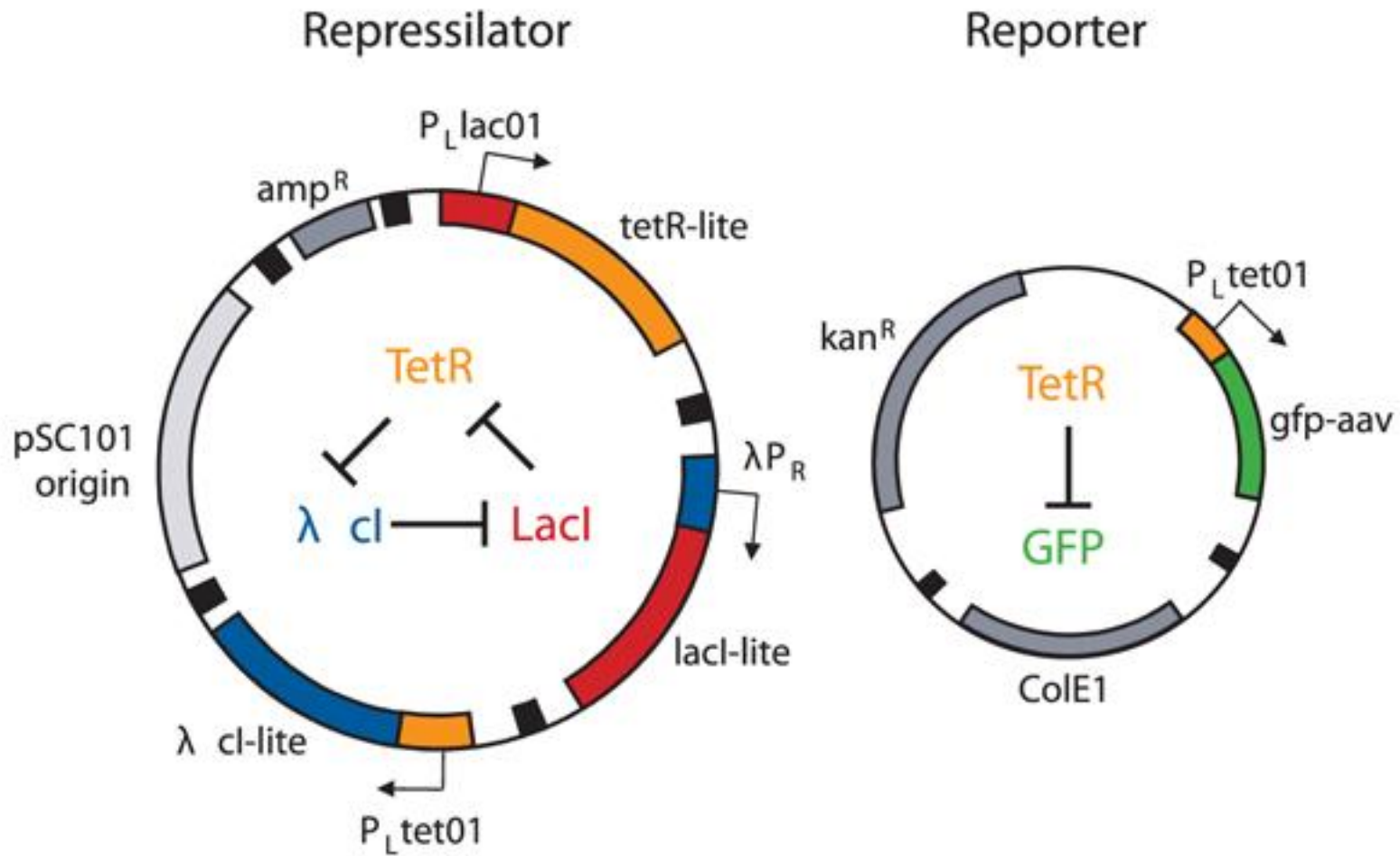
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Djurdjevac-Conrad, Schütte, Weber, 2016

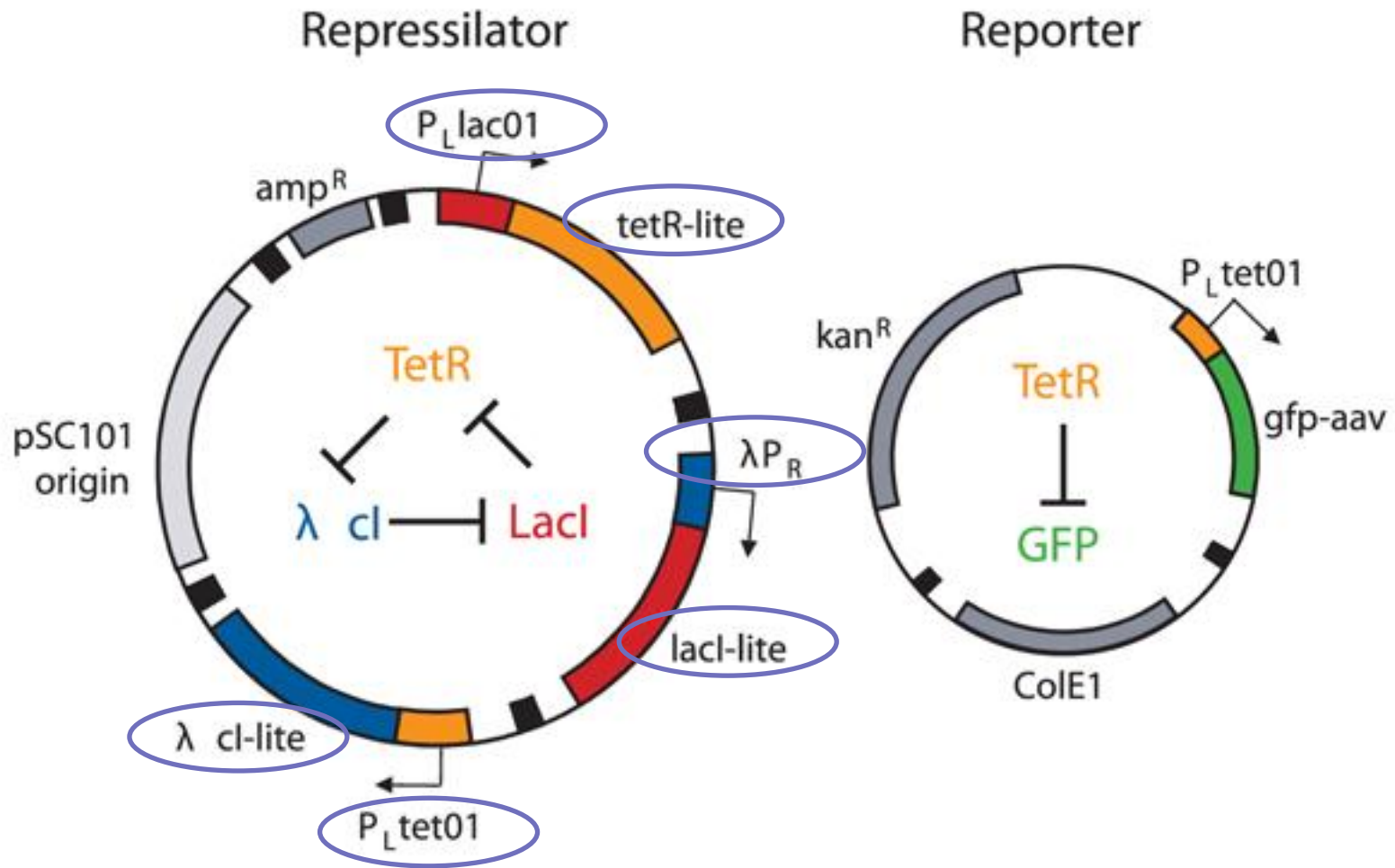
Λ

+	+	+	+
	+	+	+
		+	+
		+	+

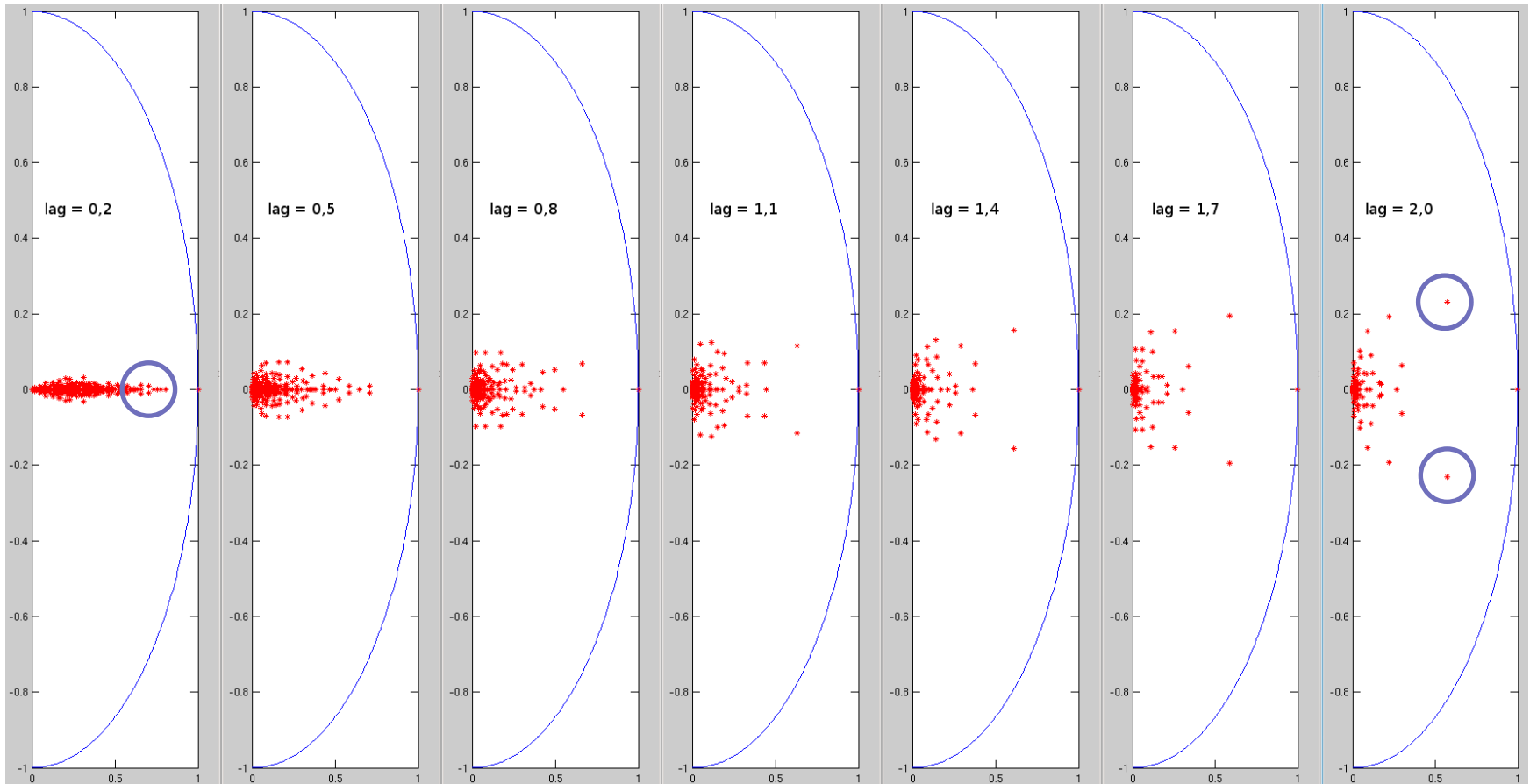
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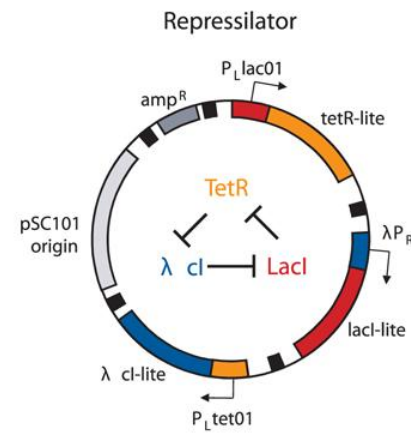
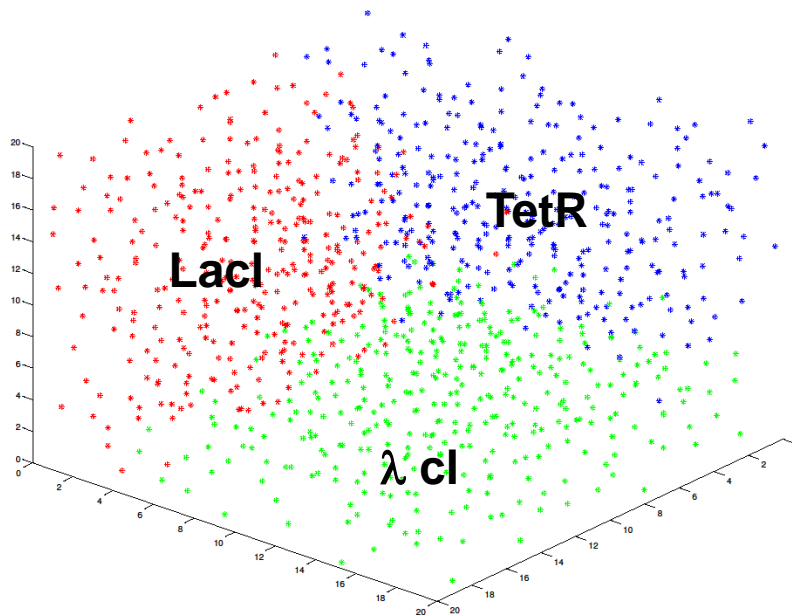
A Synthetic Oscillatory Network of Transcriptional Regulators;
[Michael Elowitz](#) and [Stanislas Leibler](#); Nature. 2000 Jan
 20;403(6767):335-8.



A Synthetic Oscillatory Network of Transcriptional Regulators;
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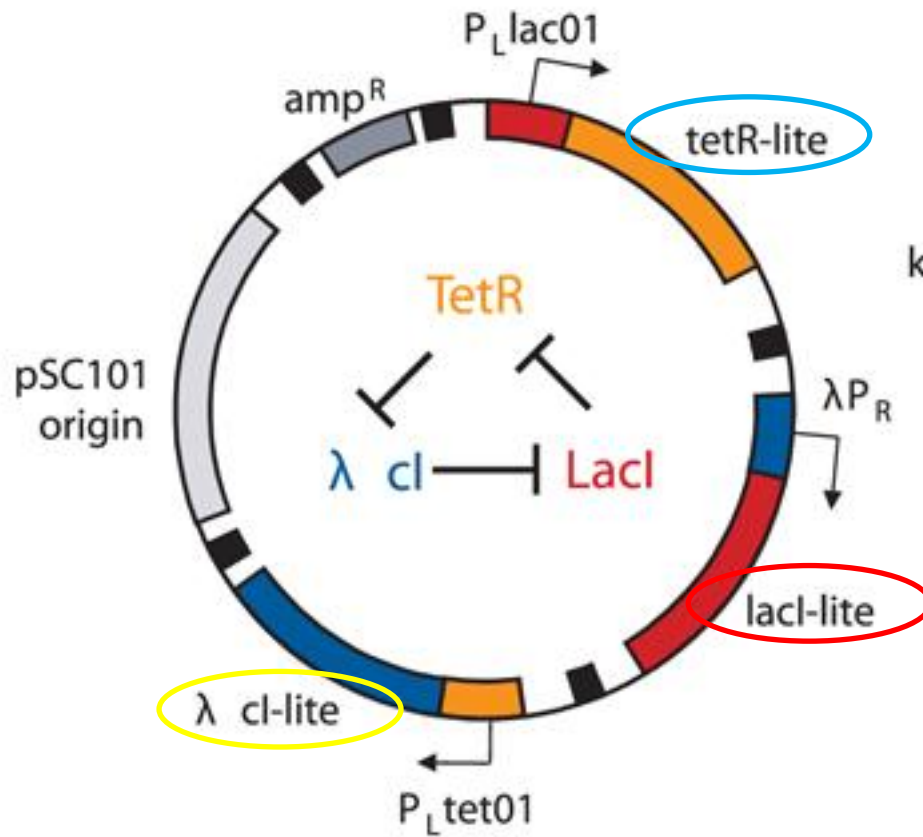


Generalized PCCA

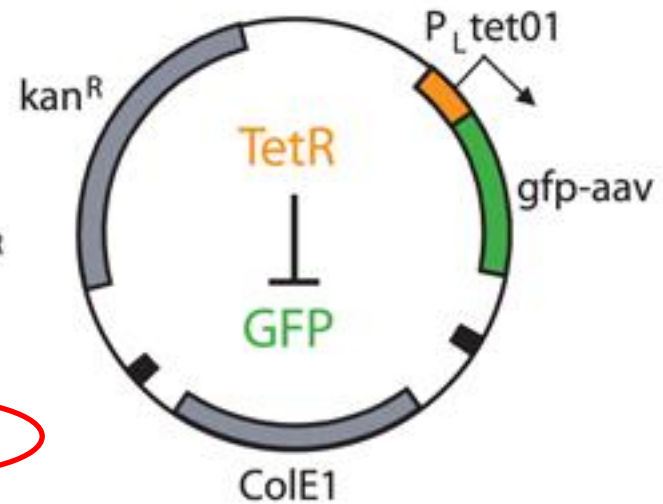


Reinmiedl, 2016
 Beckenbach, Eifler, Fackeldey, Gleixner, Grever, Weber,
 Witzig, 2016 (subm.)

Repressilator

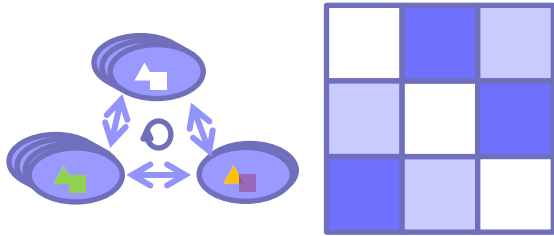


Reporter



Generalized PCCA
($n=0.005$)

Information „to go“

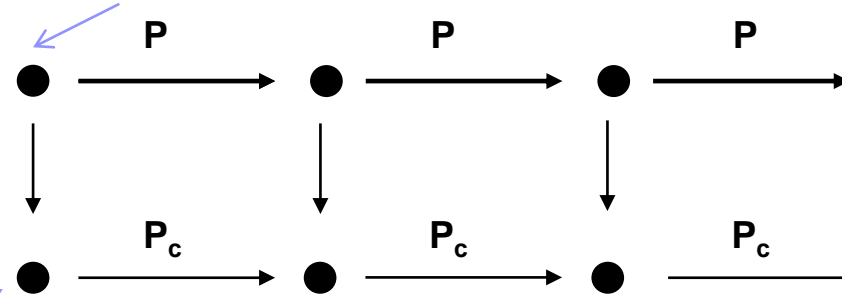


efficiency = non-reversibility

membership vectors

$$\chi = XA$$

$$PX = X\Lambda$$



linear coefficients of the membership vectors

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