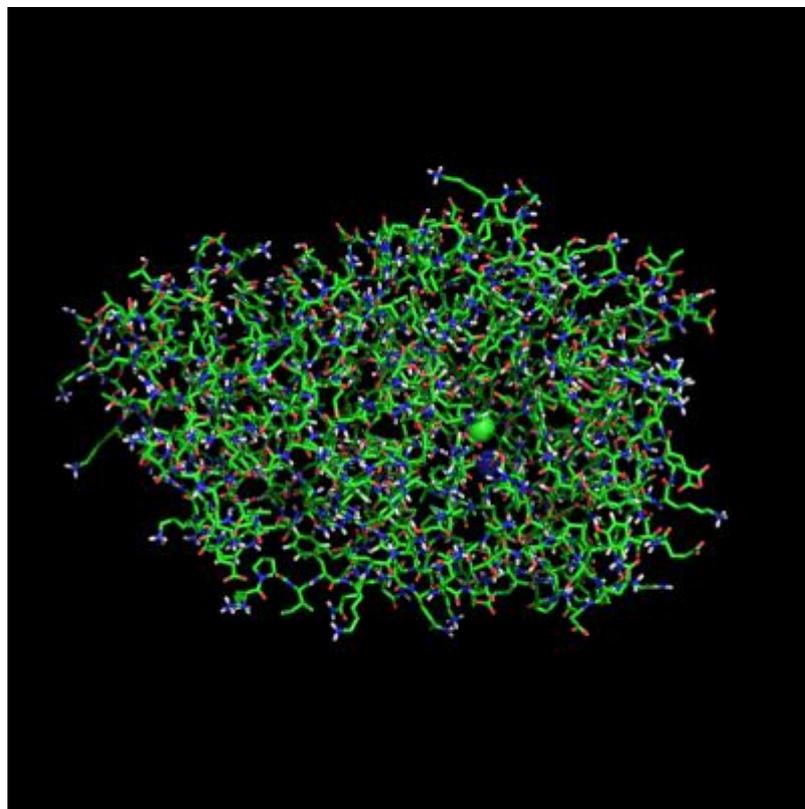


# Sparse Matrices in Molecular Simulation and in Physiological Models



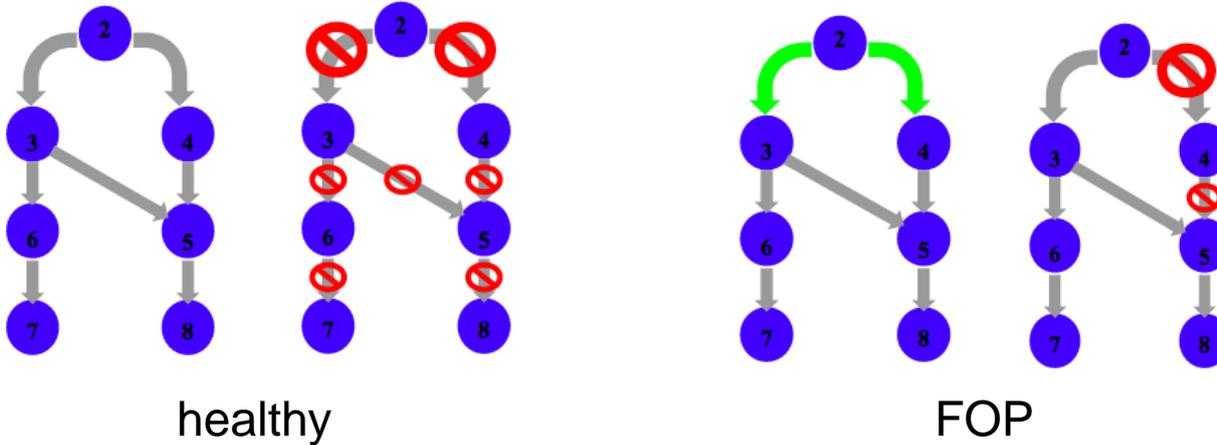
When will the ball escape?



Zuse Institute  
Berlin

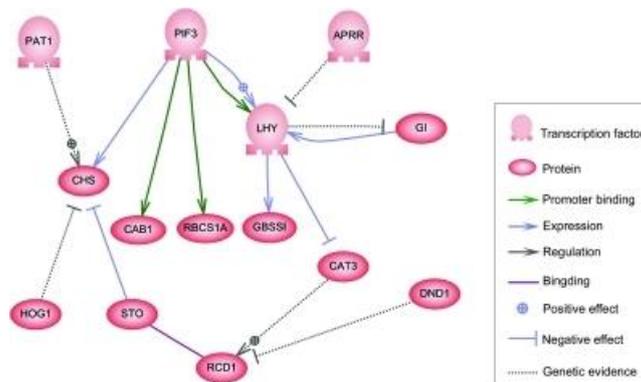
PD Dr. Marcus Weber  
Head of  
Computational Molecular Design  
[www.zib.de/weber](http://www.zib.de/weber)

# Sparse Transition Networks

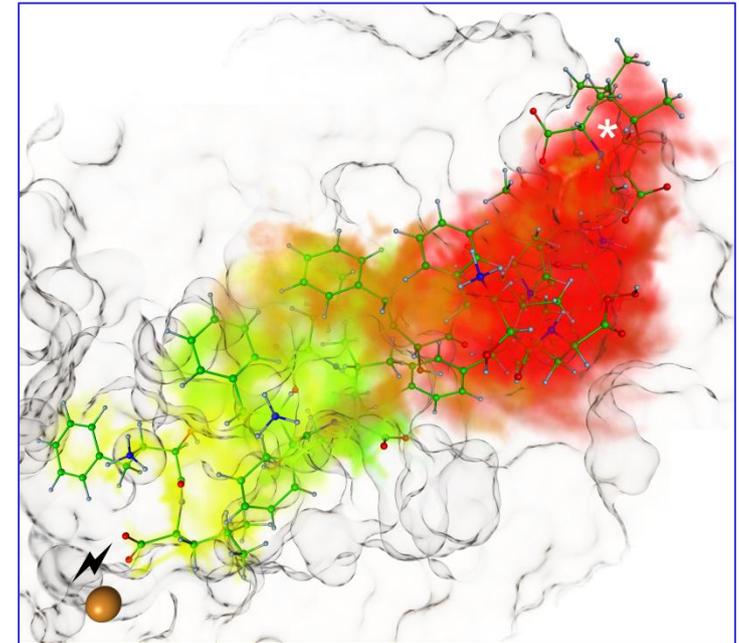
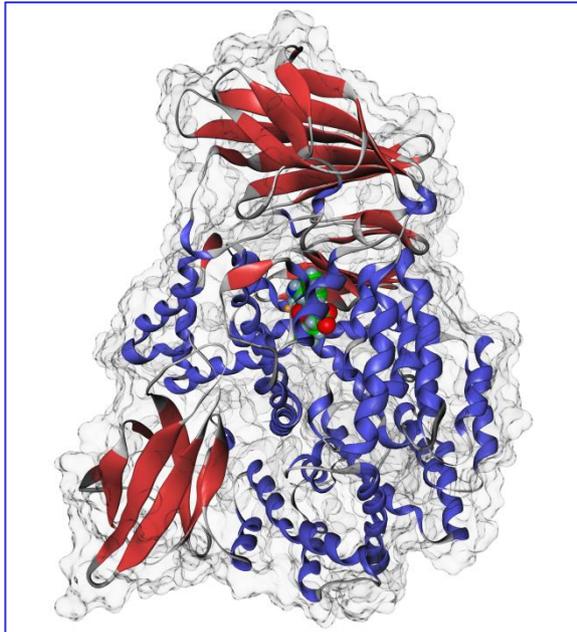


| State | Description                    |
|-------|--------------------------------|
| 2     | Receptor activation            |
| 3     | Smad1/5/8 phosphorylation      |
| 4     | p38 MAPK phosphorylation       |
| 5     | ID1/ID3 upregulation           |
| 6     | MSX2 upregulation              |
| 7     | MSX2 physiological response    |
| 8     | ID1/ID3 physiological response |

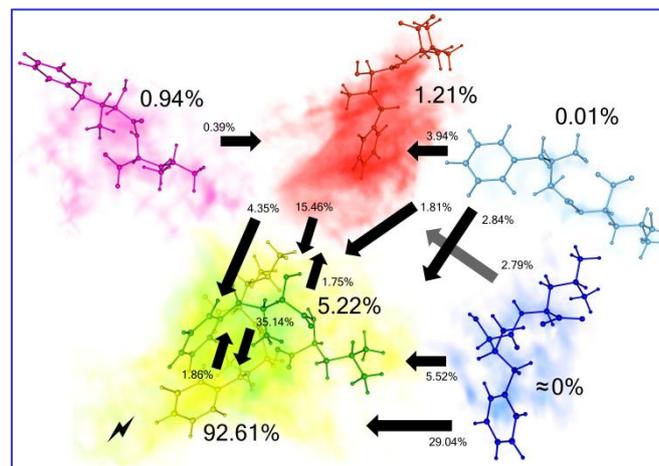
DeSouza, Quaranto, Weber (unpublished 2019)

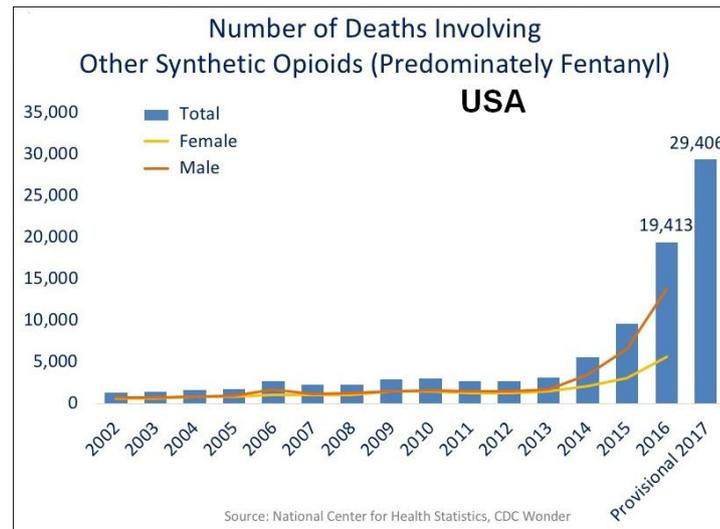


[http://openi.nlm.nih.gov/detailedresult.php?img=2993235\\_mplantssq046f07\\_4c&req=4](http://openi.nlm.nih.gov/detailedresult.php?img=2993235_mplantssq046f07_4c&req=4)



**A. Bujotzek, M. Weber:** Efficient Simulation of Ligand-Receptor Binding Processes Using the Conformation Dynamics Approach. *Journal of Bioinformatics and Computational Biology*, 7(5):811-831, April 2009

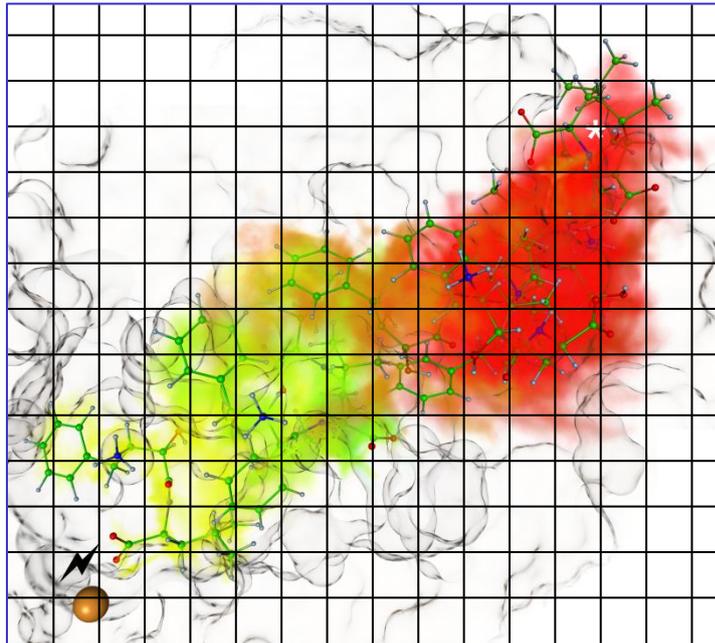




**V. Spahn, G. Del Vecchio, D. Labuz, A. Rodriguez-Gaztelumendi, N. Massaly, J. Temp, V. Durmaz, P. Sabri, M. Reidelbach, H. Machelka, M. Weber, C. Stein:** A nontoxic pain killer designed by modeling of pathological receptor conformations. *Science*, 355(6328):966-969, March 2017

# Algorithm (Sparse)

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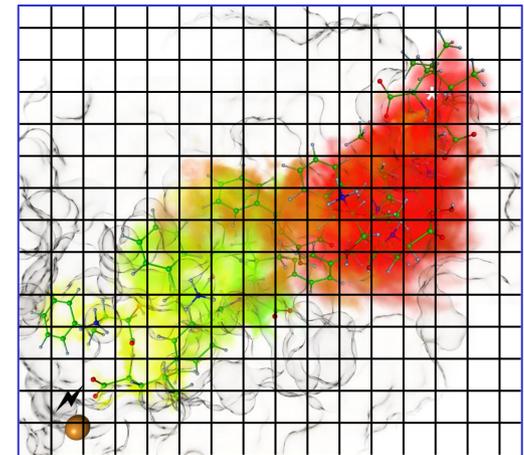
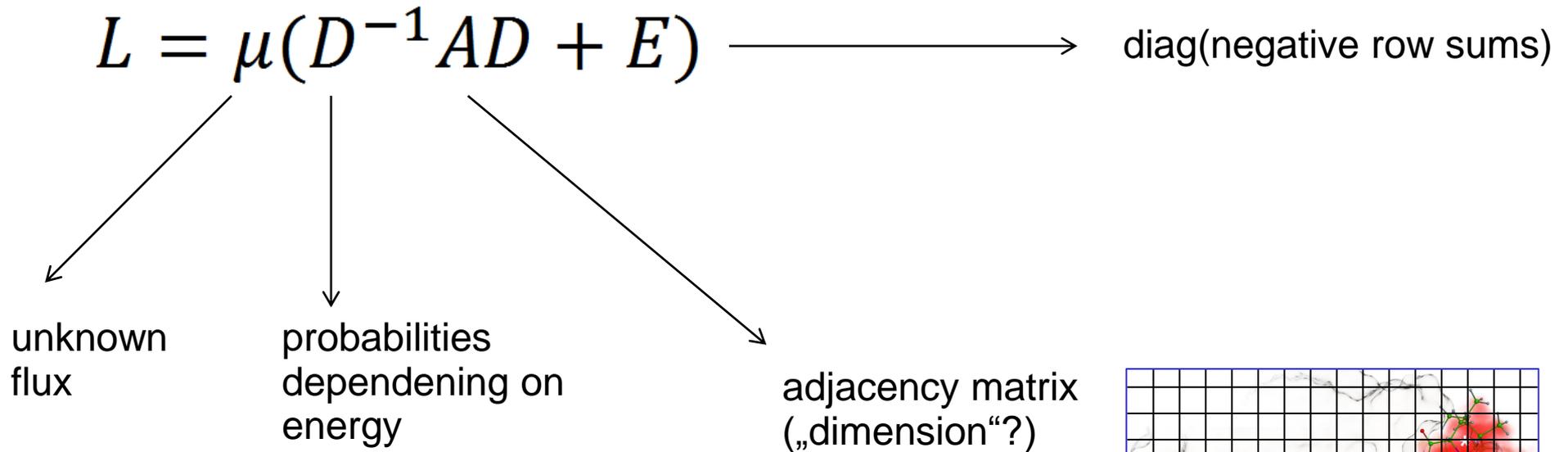


dimension:  $3N$

alternative: reaction coordinates

# Algorithm (Sparse)

L. Donati, M. Heida, M. Weber, B. Keller: Estimation of the infinitesimal generator by square-root approximation, Weierstraß Report Nr. 2416, 2018.



project B05 in CRC 1114

$P =$

|     |     |     |     |
|-----|-----|-----|-----|
| P11 | P12 | P13 | P14 |
| P21 | P22 | P23 | P24 |
| P31 | P32 | P33 | P34 |
| P41 | P42 | P43 | P44 |

$$P_{ij} = \exp(\tau L_{ij})$$

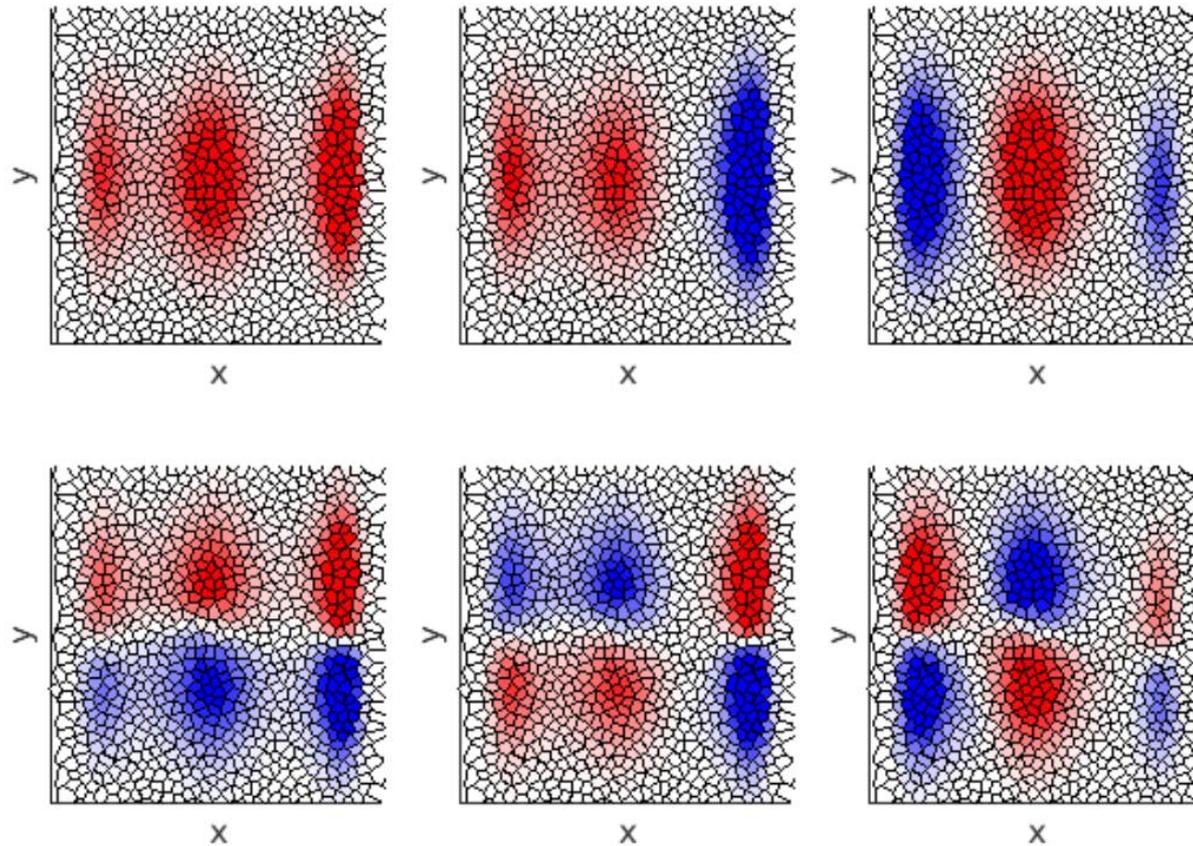
$$P = X\Lambda \longrightarrow \begin{array}{ccccc} \text{real Schur matrix} & & & & \\ 1 & * & * & * & * \\ 0 & \lambda_1 & * & * & * \\ 0 & 0 & \lambda_2 & * & * \\ 0 & 0 & * & \lambda_2 & * \\ 0 & 0 & 0 & 0 & \lambda_3 \end{array}$$

Schur vectors  
 $X^T D^2 X = I$

- slowest timescales
- rate-determining steps
- reaction coordinates
- reversibility (entropy production)

# Intepretation of Solution

For one point in time:



$P =$

|     |     |     |     |
|-----|-----|-----|-----|
| P11 | P12 | P13 | P14 |
| P21 | P22 | P23 | P24 |
| P31 | P32 | P33 | P34 |
| P41 | P42 | P43 | P44 |

$$P_{ij} = \exp(\tau L_{ij})$$

**K. Fackeldey, P. Koltai, P. Nevir, H. Rust, A.Schild, M. Weber:** From Metastable to Coherent Sets -- time-discretization schemes. *Chaos*, 29:012101, 2019.

# Computational Alternatives (Missing Data)

$$P = \begin{pmatrix} \mathbf{x} & \dots & \dots & \mathbf{x} \\ \mathbf{u} & \dots & \dots & \mathbf{u} \\ \vdots & & & \vdots \\ \mathbf{x} & \dots & \dots & \mathbf{x} \\ \mathbf{u} & \dots & \dots & \mathbf{u} \\ \mathbf{x} & \dots & \dots & \mathbf{x} \\ \mathbf{x} & \dots & \dots & \mathbf{x} \\ \mathbf{u} & \dots & \dots & \mathbf{u} \\ \vdots & & & \vdots \\ \vdots & \dots & \dots & \vdots \end{pmatrix} \rightsquigarrow P_R = \begin{pmatrix} \mathbf{x} & \dots & \dots & \mathbf{x} \\ \mathbf{x} & \dots & \dots & \mathbf{x} \\ \mathbf{x} & \dots & \dots & \mathbf{x} \\ \vdots & & & \vdots \\ \mathbf{x} & \dots & \dots & \mathbf{x} \end{pmatrix}$$

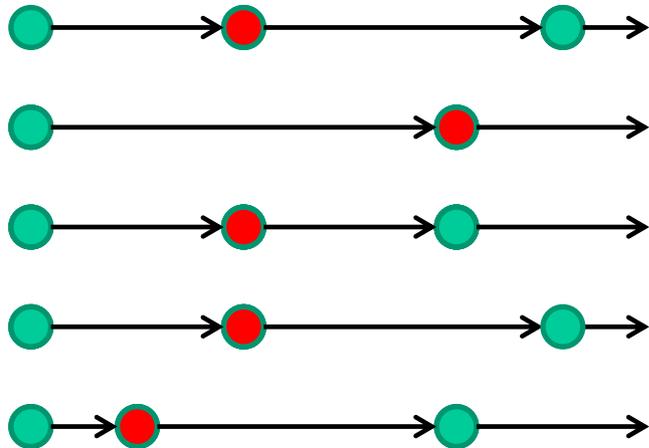
**Figure 2:** Given the matrix  $P$  we delete the unknown rows and obtain  $P_R$  which is rectangular. The matrix  $P_R$  has  $k$  rows and  $n$  columns.

$$\tilde{P} = \begin{pmatrix} * & \dots & \dots & \mathbf{x} \\ \mathbf{x} & \ddots & \dots & \mathbf{x} \\ \mathbf{x} & \dots & \ddots & \mathbf{x} \\ \mathbf{x} & \dots & \dots & * \end{pmatrix}$$

**Figure 3:** The matrix  $\tilde{P}$  is constructed by skipping the corresponding columns  $\ell \in [n - k]$  and summing the skipped values row-wise onto the diagonal, \* - summed diagonal elements according to (5).

**K. Fackeldey, A. Niknejad, M. Weber:** Finding Metastabilities in Reversible Markov Chains based on Incomplete Sampling: Case of Molecular Simulation. *Spec. Matrices*, 5:73–81, 2017

# Computational Alternatives (Concept)



|     |     |     |     |
|-----|-----|-----|-----|
| L11 | L12 | L13 | L14 |
| L21 | L22 | L23 | L24 |
| L31 | L32 | L33 | L34 |
| L41 | L42 | L43 | L44 |

# Research Question

Algorithms to compute (partial, real) Schur decompositions of matrices which are not sparse, but which are constructed block-wise from exponentials of sparse matrices.



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