

# TENSOR TRAINS FOR HIGHLY MULTIDIMENSIONAL DYNAMICS SIMULATIONS

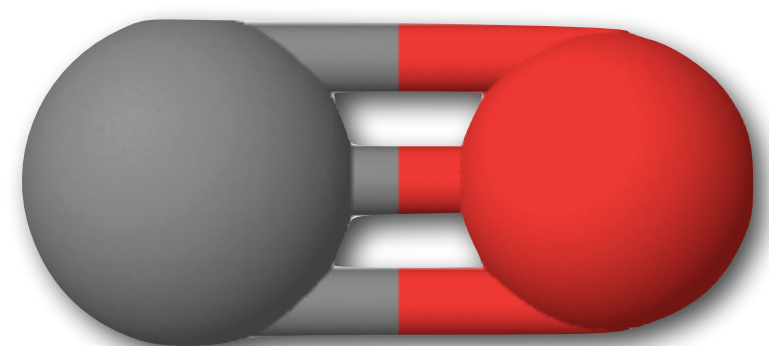
**MICHELINE B. SOLEY**

DEPT. OF CHEMISTRY AND DEPT. OF PHYSICS-AFFILIATE,  
UNIVERSITY OF WISCONSIN-MADISON

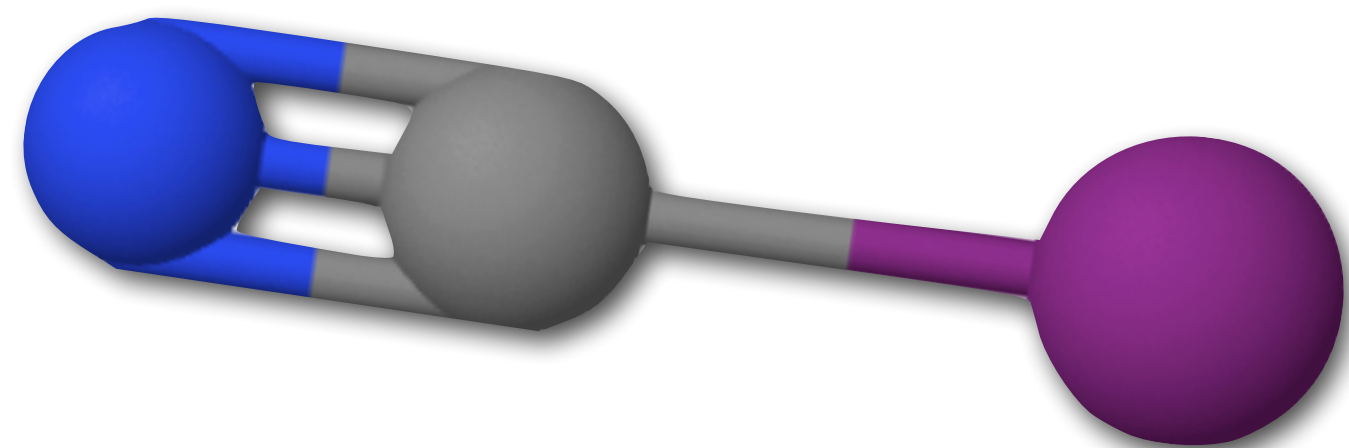
VIRTUAL INTERNATIONAL SEMINAR ON THEORETICAL ADVANCEMENTS (VISTA)

# CURSE OF DIMENSIONALITY

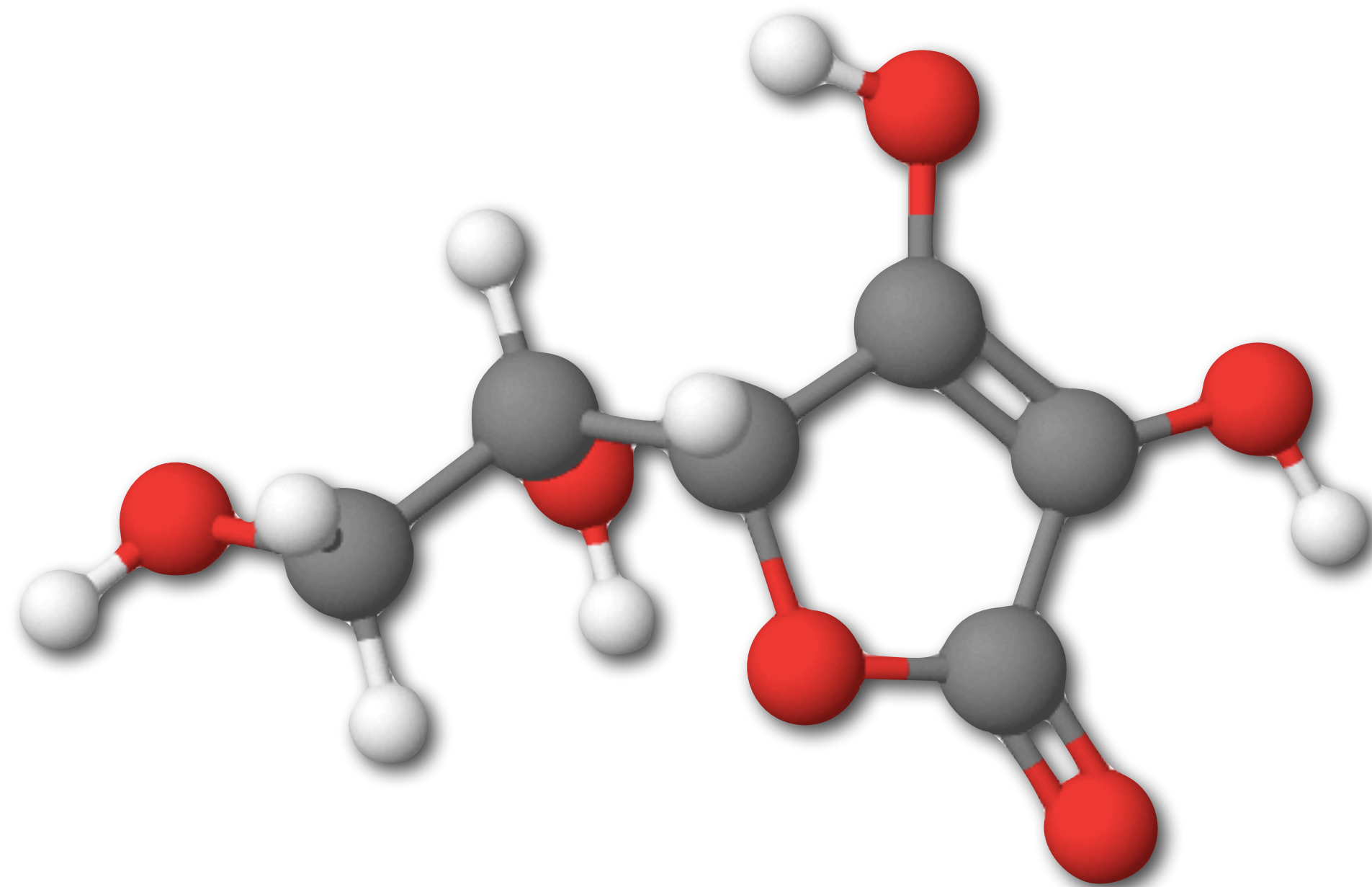
Many classical computer approaches to quantum mechanics simulations become computationally intractable for large molecular systems.



2 Atoms:  
2 KB



3 Atoms:  
100 MB



20 Atoms:  
 $10^{119}$  TB

# LOW-RANK TENSOR-TRAIN DECOMPOSITION REDUCES COST

Original Image



100%

Tensor Train



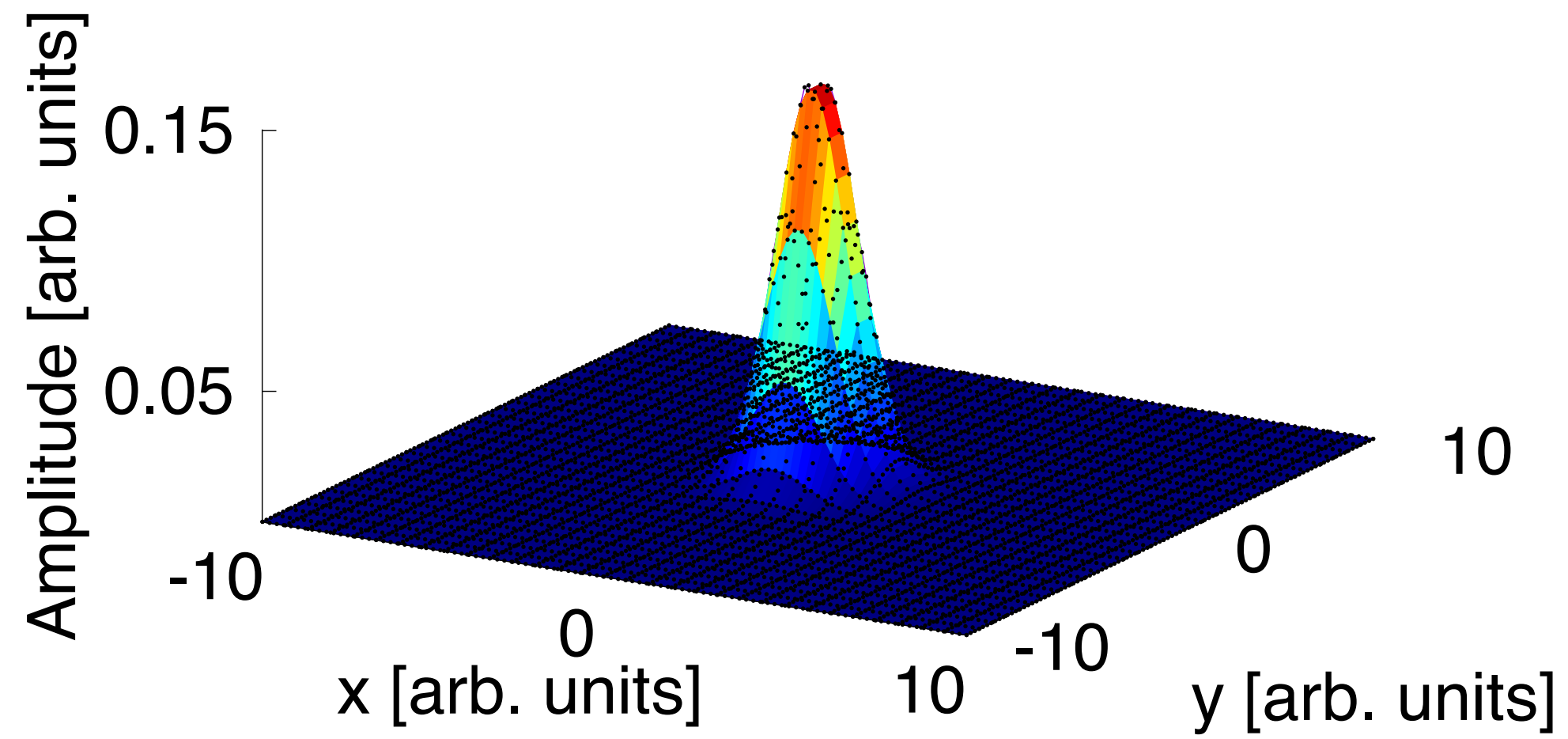
15%

N. Lyu\*, E. Mulvihill\*, **Micheline B. Soley**, E. Geva, V. S. Batista, (2023) JCTC, in press.  
T. H. Kyaw\*, **Micheline B. Soley**,\* B. Allen, P. Bergold, C. Sun, V. S. Batista, A. Aspuru-Guzik, (2022) arXiv:2208.10470v1.  
**Micheline B. Soley**,\* P. E. Videla,\* E. T. J. Nibbering, V. S. Batista, J. Phys. Chem. Lett., 13 (2022) 8354.  
**Micheline B. Soley**, P. Bergold, A. A. Gorodetsky, V. S. Batista, JCTC, 18 (2022) 25.

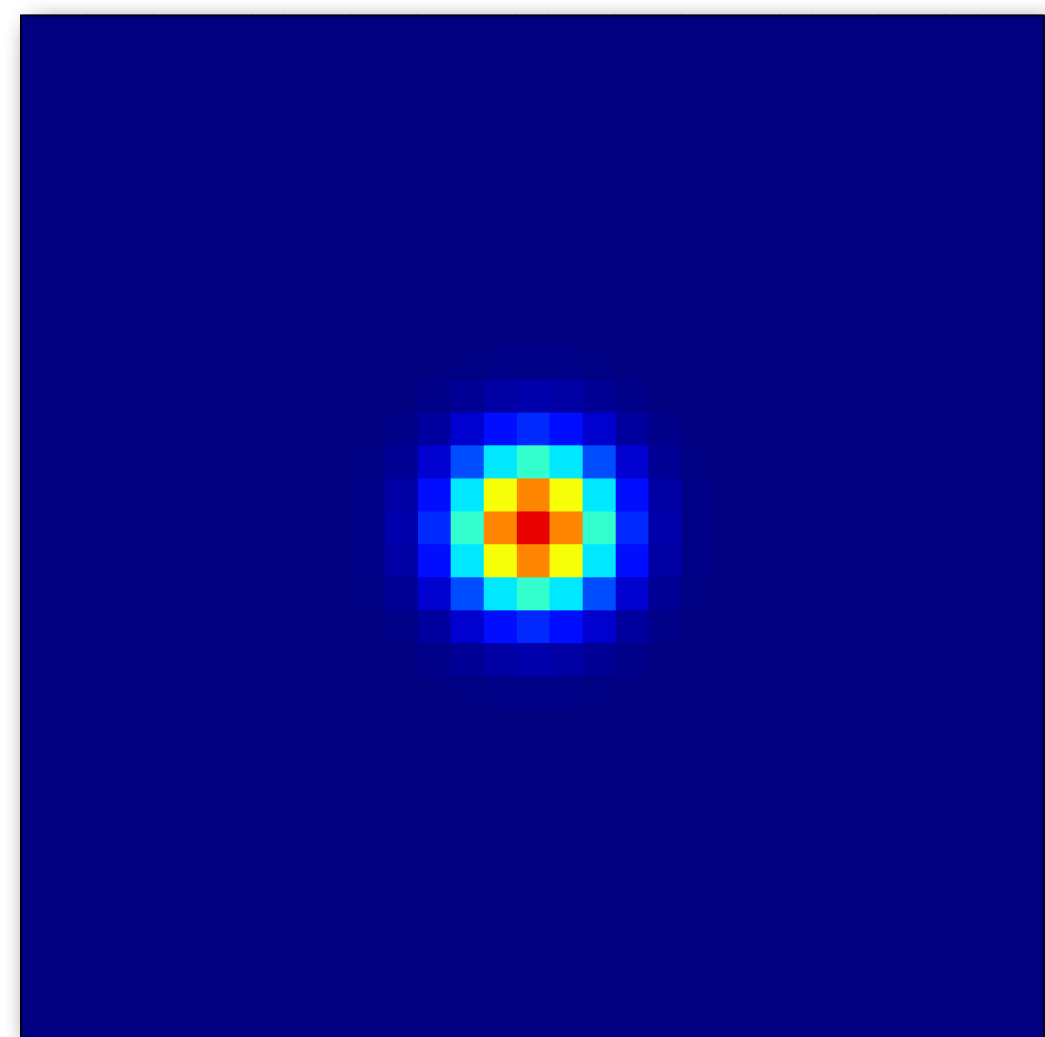
Y. Wang, E. Mulvihill, Z. Hu, N. Lyu, S. Shivpuje, Y. Liu, **Micheline B. Soley**, E. Geva, V. S. Batista, S. Kais, 2022, arXiv:2209.04956.  
N. Lyu, **Micheline B. Soley**, V. S. Batista, JCTC, 18 (2022) 3327.  
**Micheline B. Soley**, P. Bergold, V. S. Batista, JCTC 17 (2021) 3280.  
B. N. Khoromskij, Constr. Approx. 34 (2011) 257.  
I. Oseledets, E. Tyryshnikov, Linear Algebra Appl. 432 (2010) 70.  
J. C. Napp, et al. Phys. Rev. X 12 (2022) 021021.

## 2D Gaussian

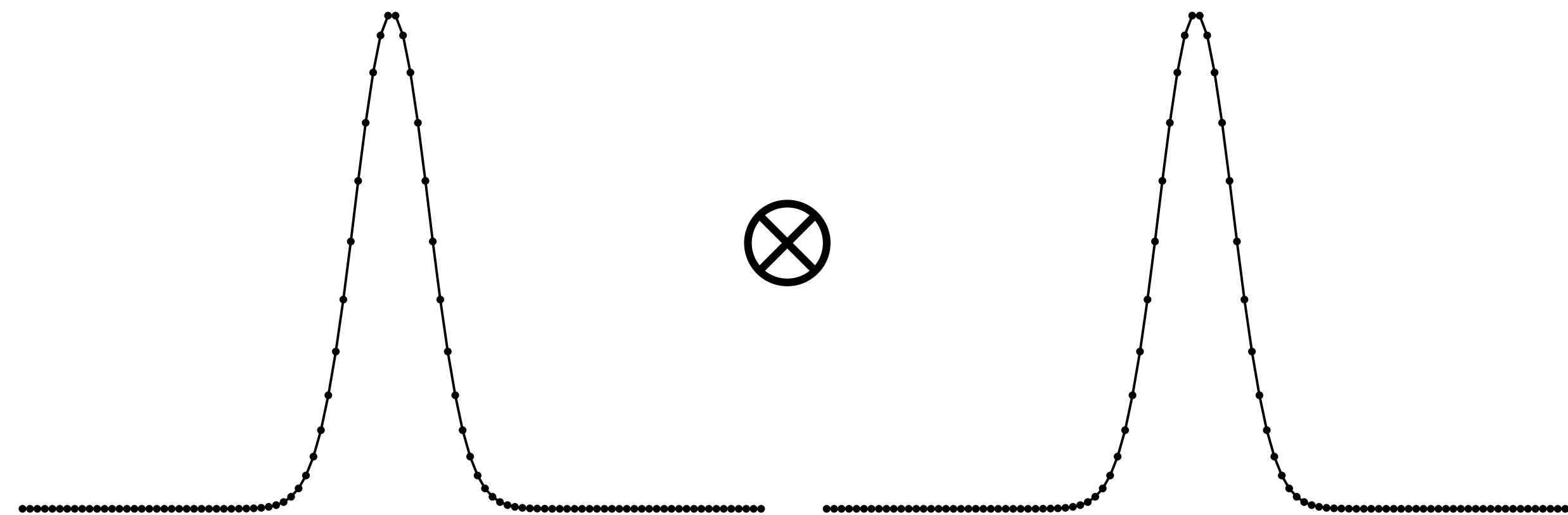
## 2D TENSOR TRAIN (RANK 1)



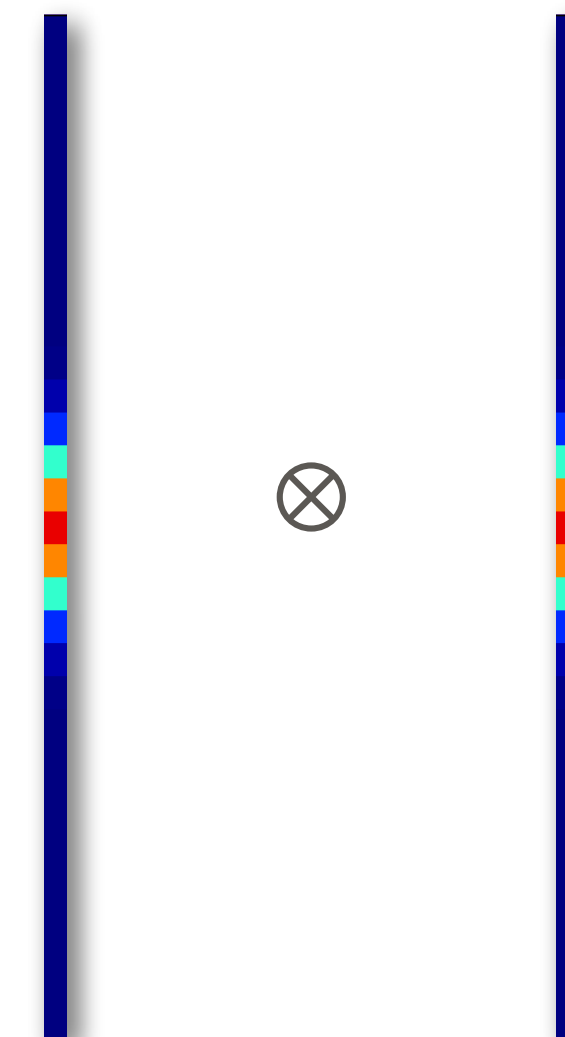
$$\Psi(x, y) = \frac{1}{2\pi} \exp\left(-\frac{x^2}{2} - \frac{y^2}{2}\right)$$



**vs.**



$$\begin{aligned} \psi(x)\psi(y) &= \left(\frac{1}{\sqrt{2\pi}} \exp\left(-\frac{x^2}{2}\right)\right) \left(\frac{1}{\sqrt{2\pi}} \exp\left(-\frac{y^2}{2}\right)\right) \\ &= \sum_{\alpha=1}^r \psi_1(1, x, \alpha) \psi_2(\alpha, y, 1) \end{aligned}$$

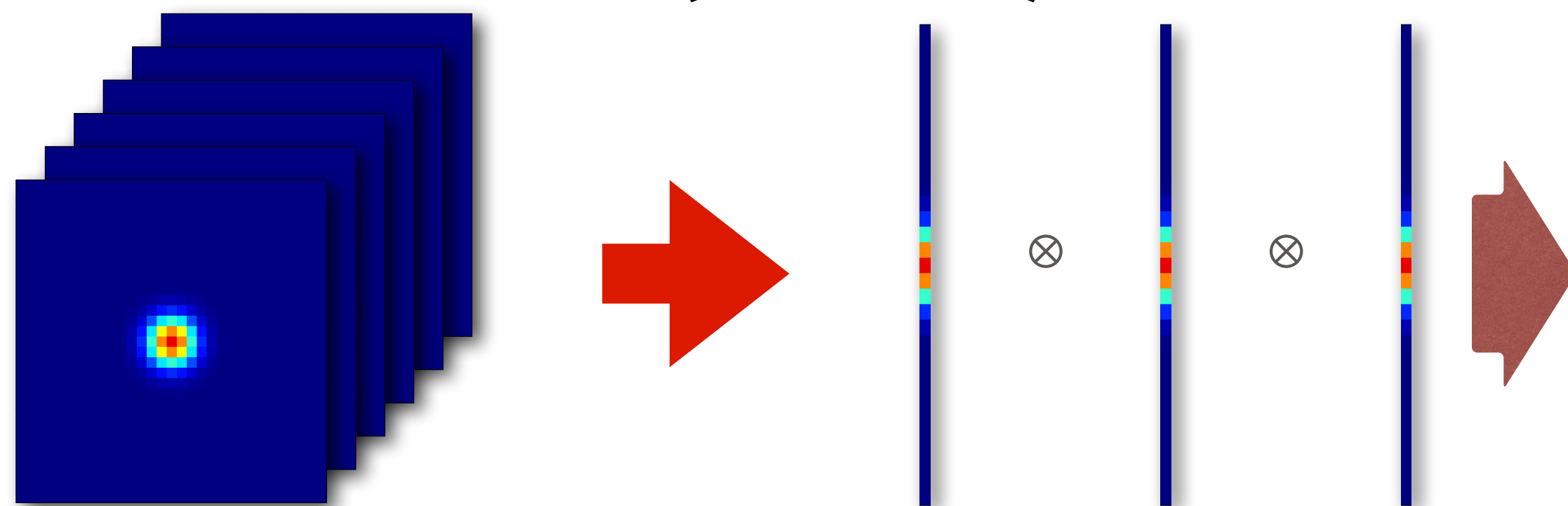


⊗

Cost of evaluating reduces to

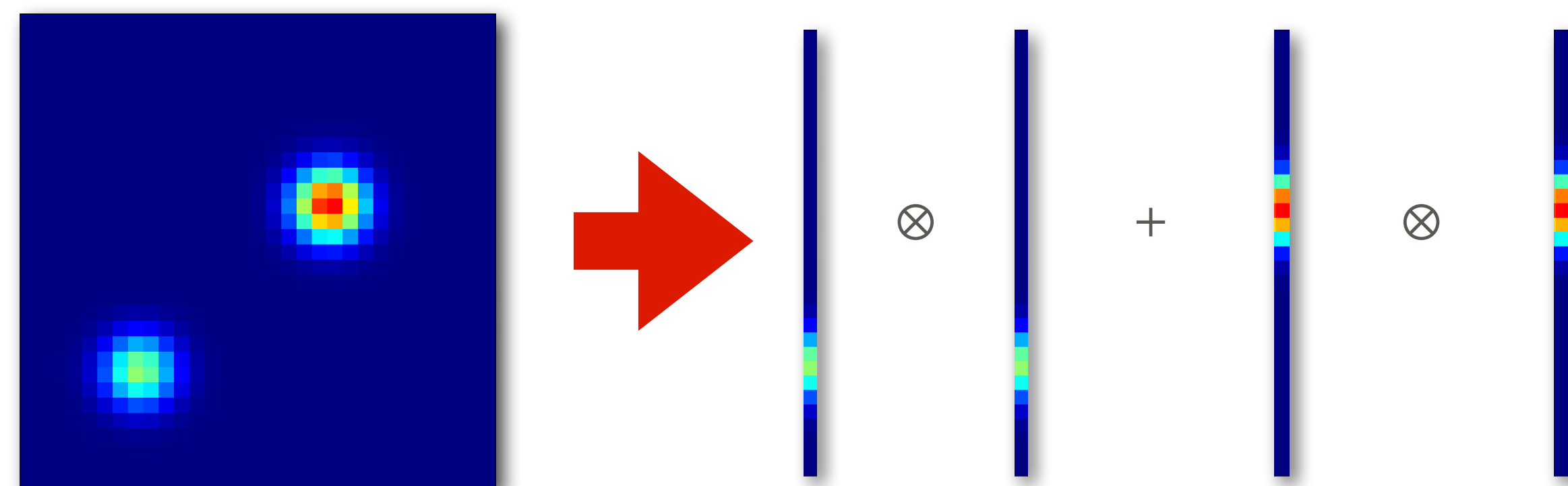
$$n^2 \rightarrow 2n$$

## ND TENSOR TRAIN (RANK 1)



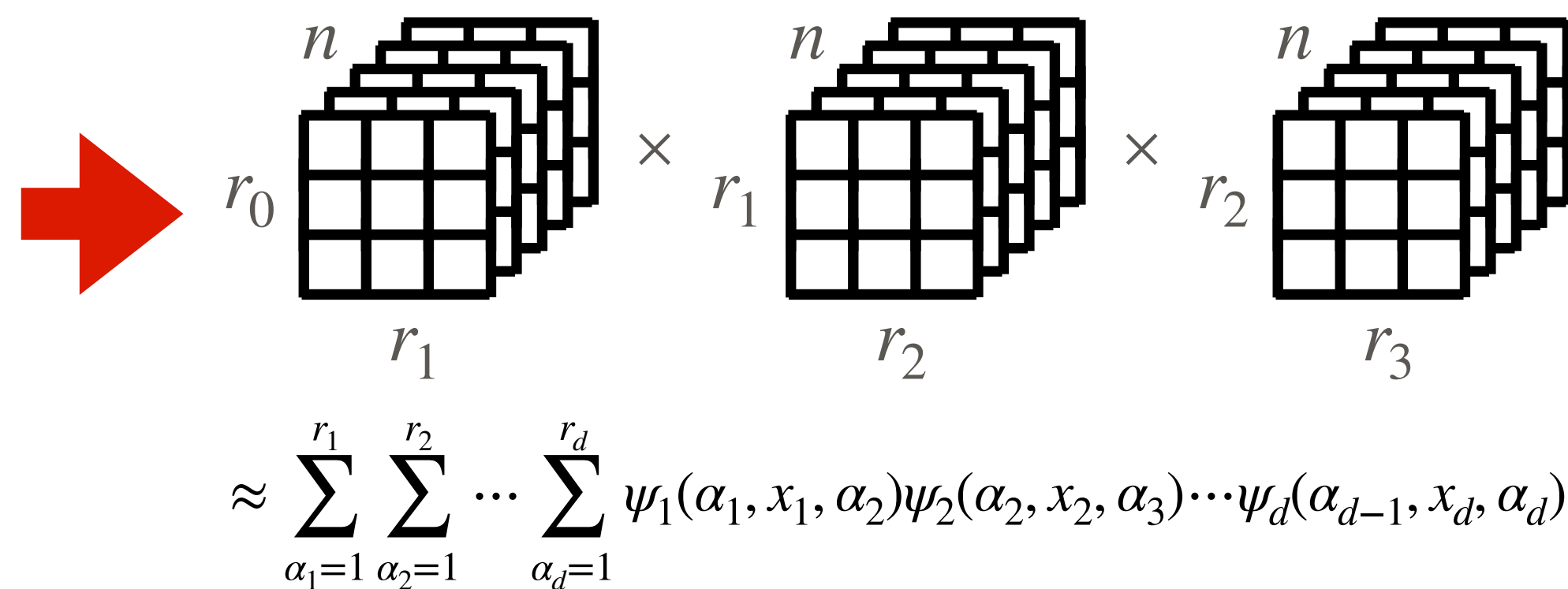
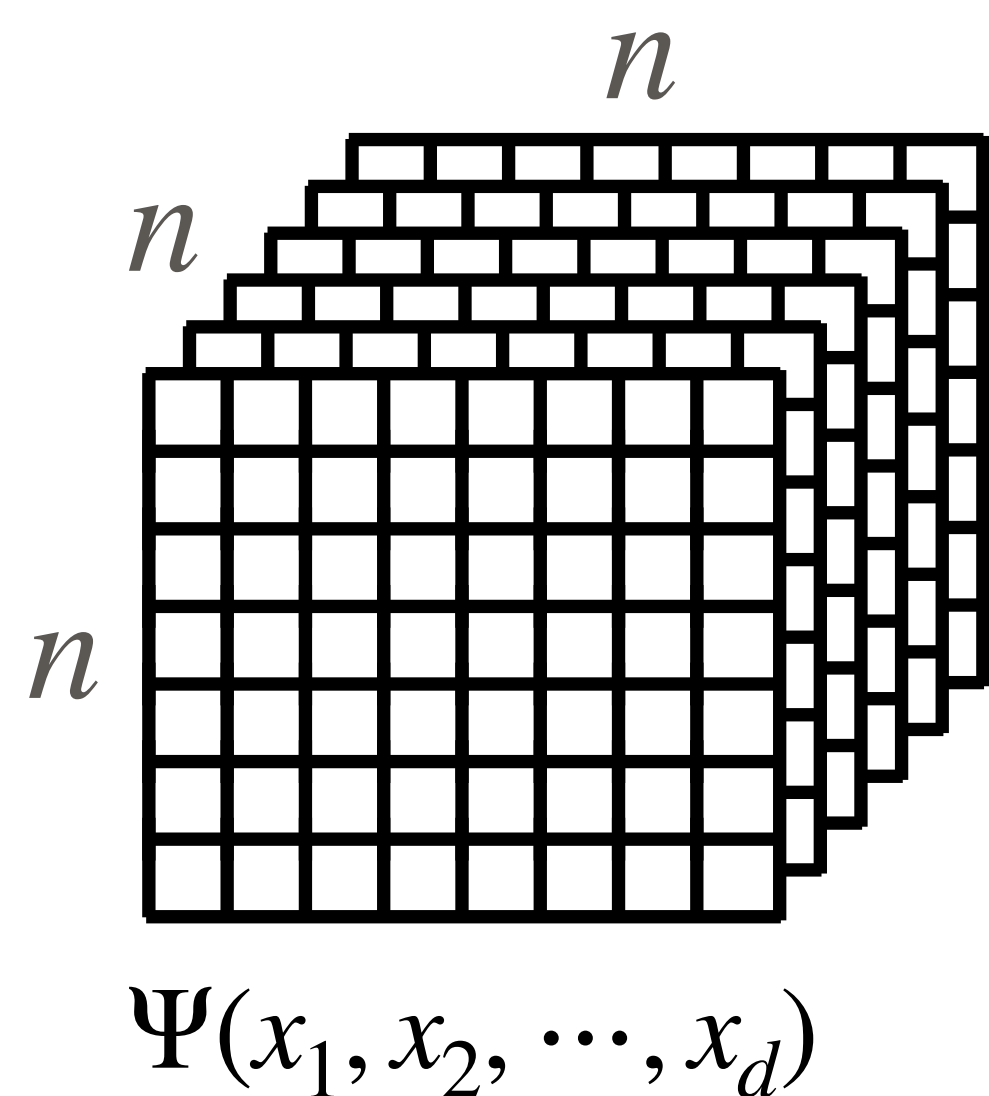
Cost Reduction:  $n^3 \rightarrow 3n$

## 2D TENSOR TRAIN (RANK 2)



Cost Reduction:  $n^2 \rightarrow 2 \times 2n$

## ND TENSOR TRAIN (RANK R)

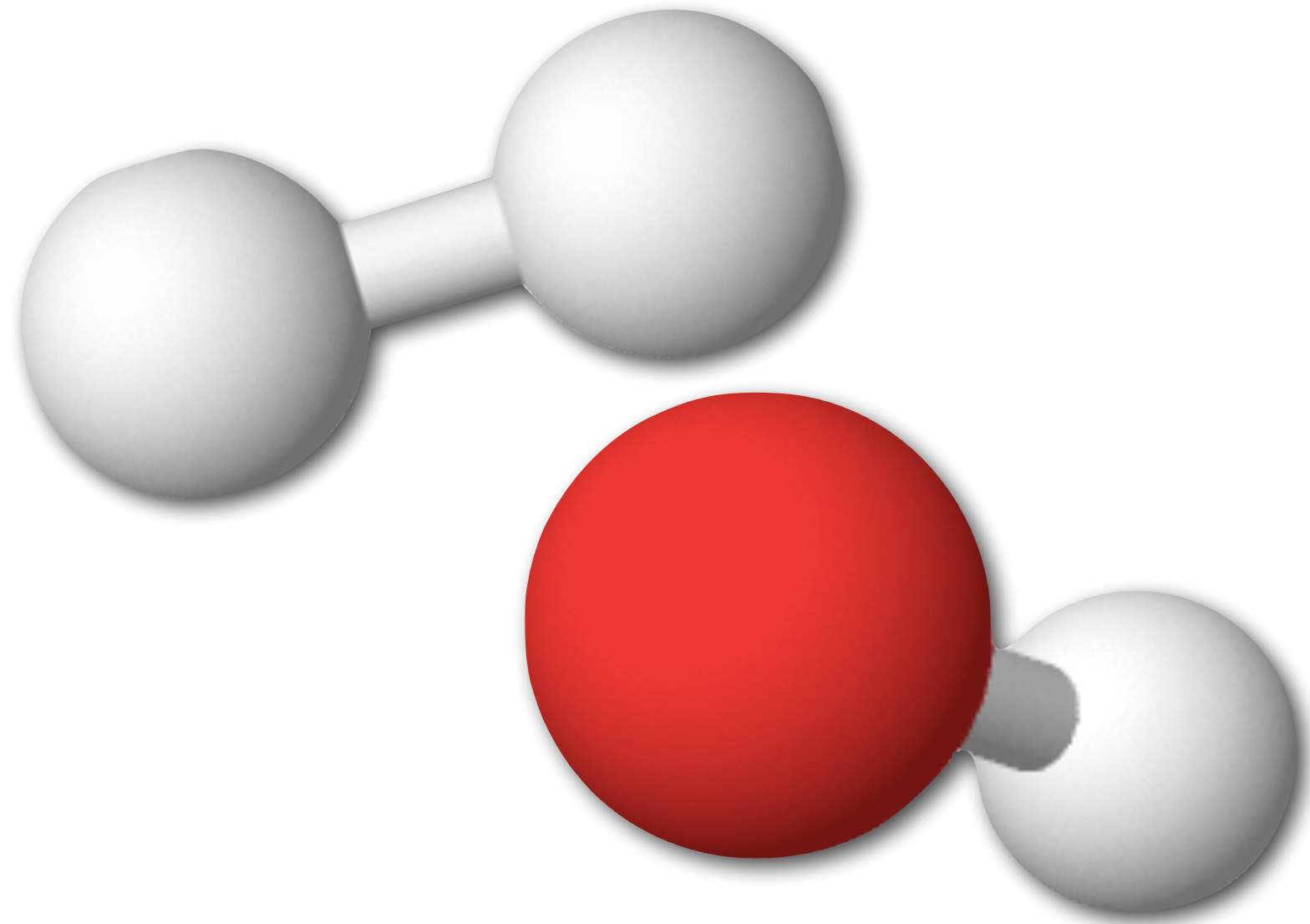


Cost Reduction  
 $n^d \rightarrow dnr^2$

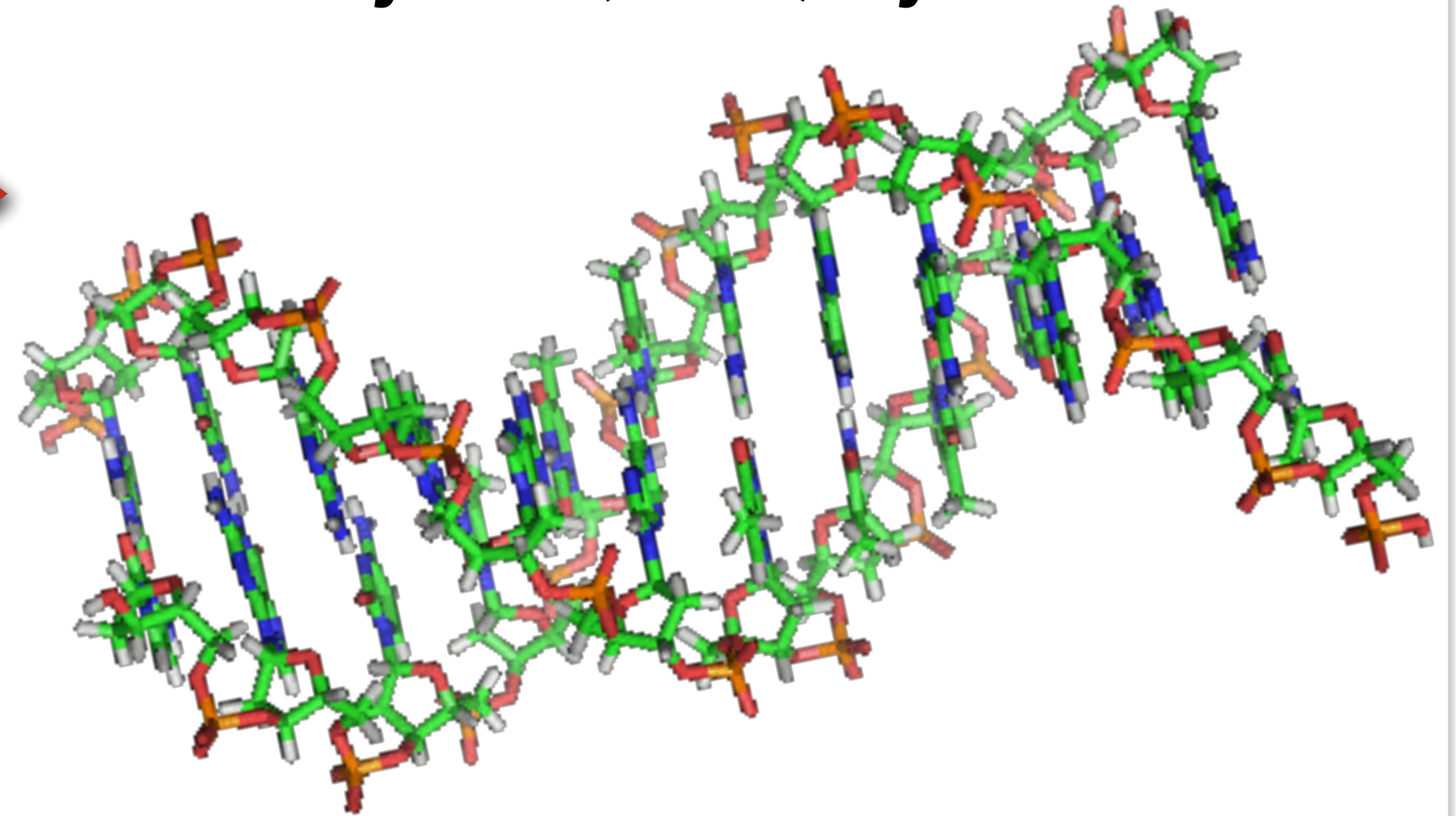
M.-L. Li, K. S. Candan, M. L. Salino, "GTT: Guiding the tensor train decomposition." International Conference on Similarity Search and Applications. Springer, Cham, 2020.  
 I. Oseledets, E. Tyryshnikov, Linear Algebra Appl. 432 (2010) 70.

# TENSOR TRAINS FOR HIGHLY MULTIDIMENSIONAL QUANTUM DYNAMICS

Largest System Investigable with Standard Chebyshev Dynamics



Functional Tensor-Train Chebyshev (FTTC) Dynamics



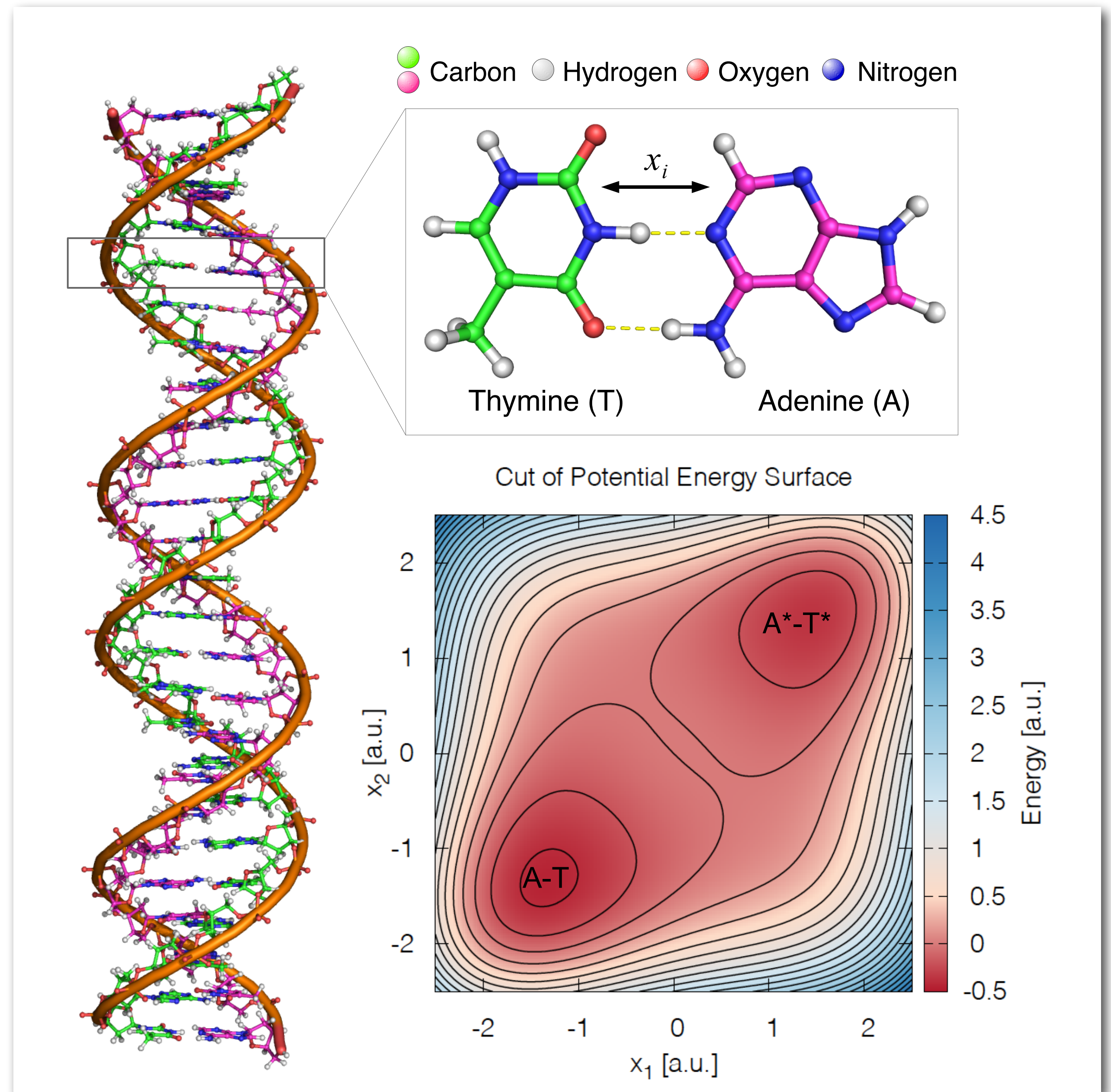
M. T. Cvitaš, S. C. Althorpe, J. Chem. Phys. 139 (2013) 064307.  
E. M. Goldfield, S. K. Gray, J. Chem. Phys. 117 (2002) 1604.  
H. Tal-Ezer, R. Kosloff, J. Chem. Phys. 81 (1984) 3967.

[Micheline B. Soley, P. Bergold, A. A. Gorodetsky, V. S. Batista, JCTC, 18 \(2022\) 25.](#)

# FUNCTIONAL TENSOR TRAIN CHEBYSHEV (FTTC) DYNAMICS

$$\Psi(t) = e^{-it\hat{H}}\Psi(0)$$

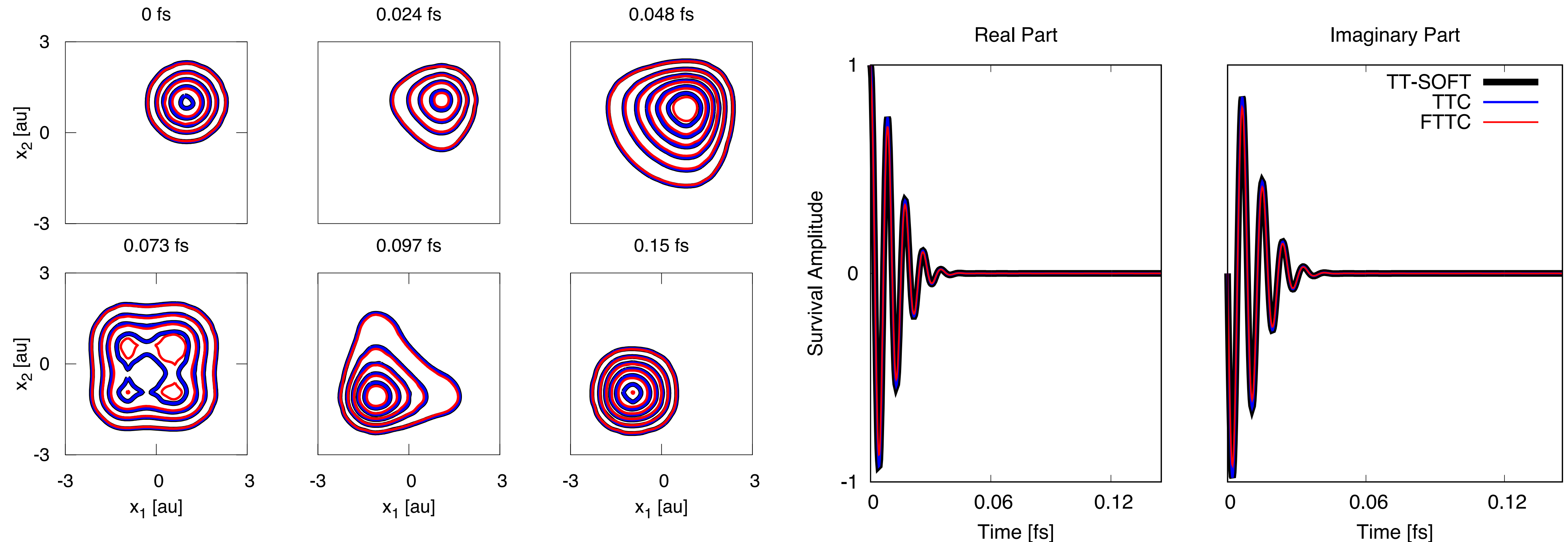
$$\approx e^{-it^+} \sum_{k=0}^{N-1} (2 - \delta_{k,0})(-i)^k J_k(t^-) T_k(\hat{H}_0) \mathbf{W}_0$$



# APPLICATION: HYDROGEN BONDING IN DNA

Probability Density Dynamics, Uncoupled Bath

Survival Amplitude

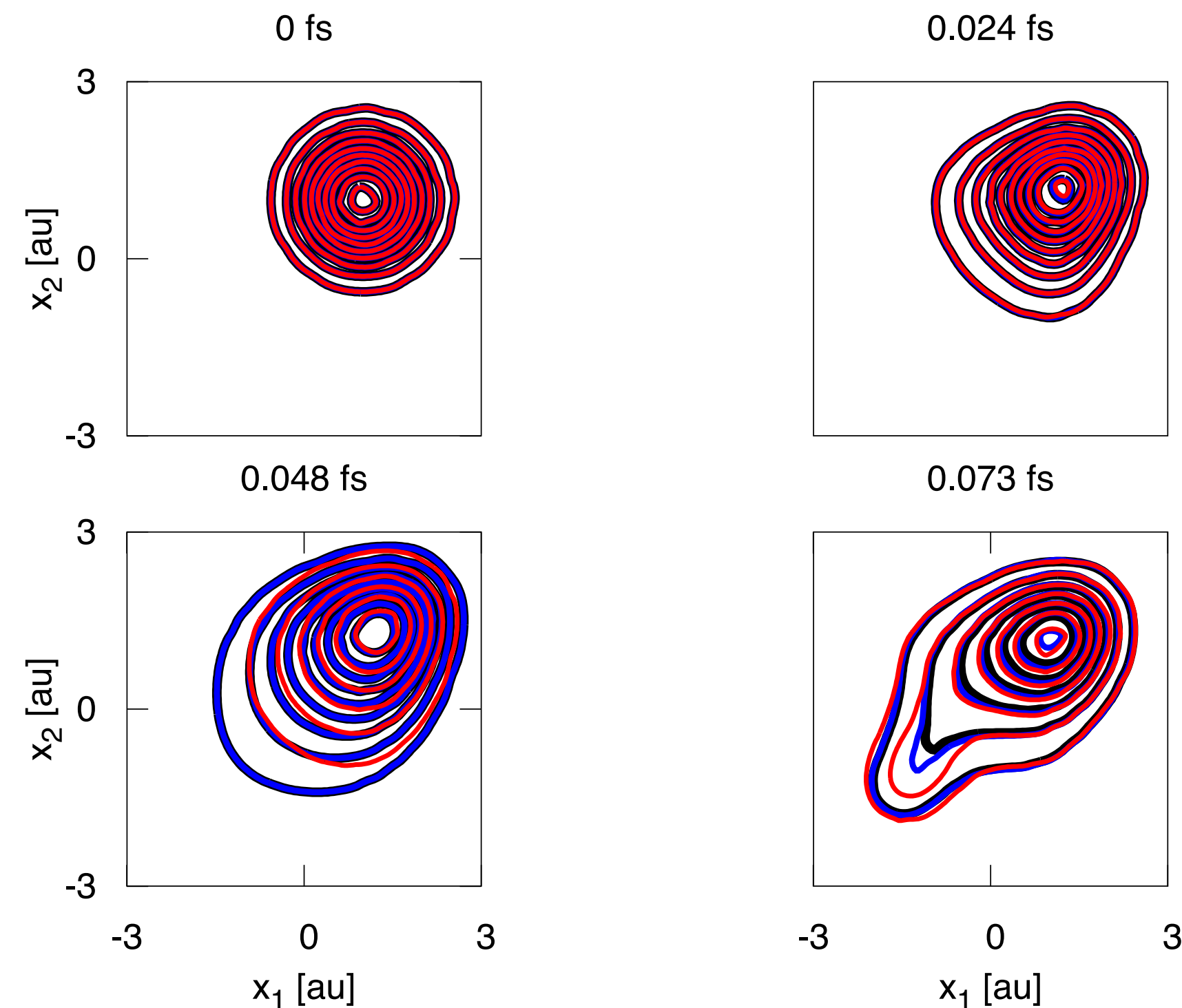


FTTC extends the Chebyshev method from simulation of four-atom systems to molecular systems in ***50 dimensions.***

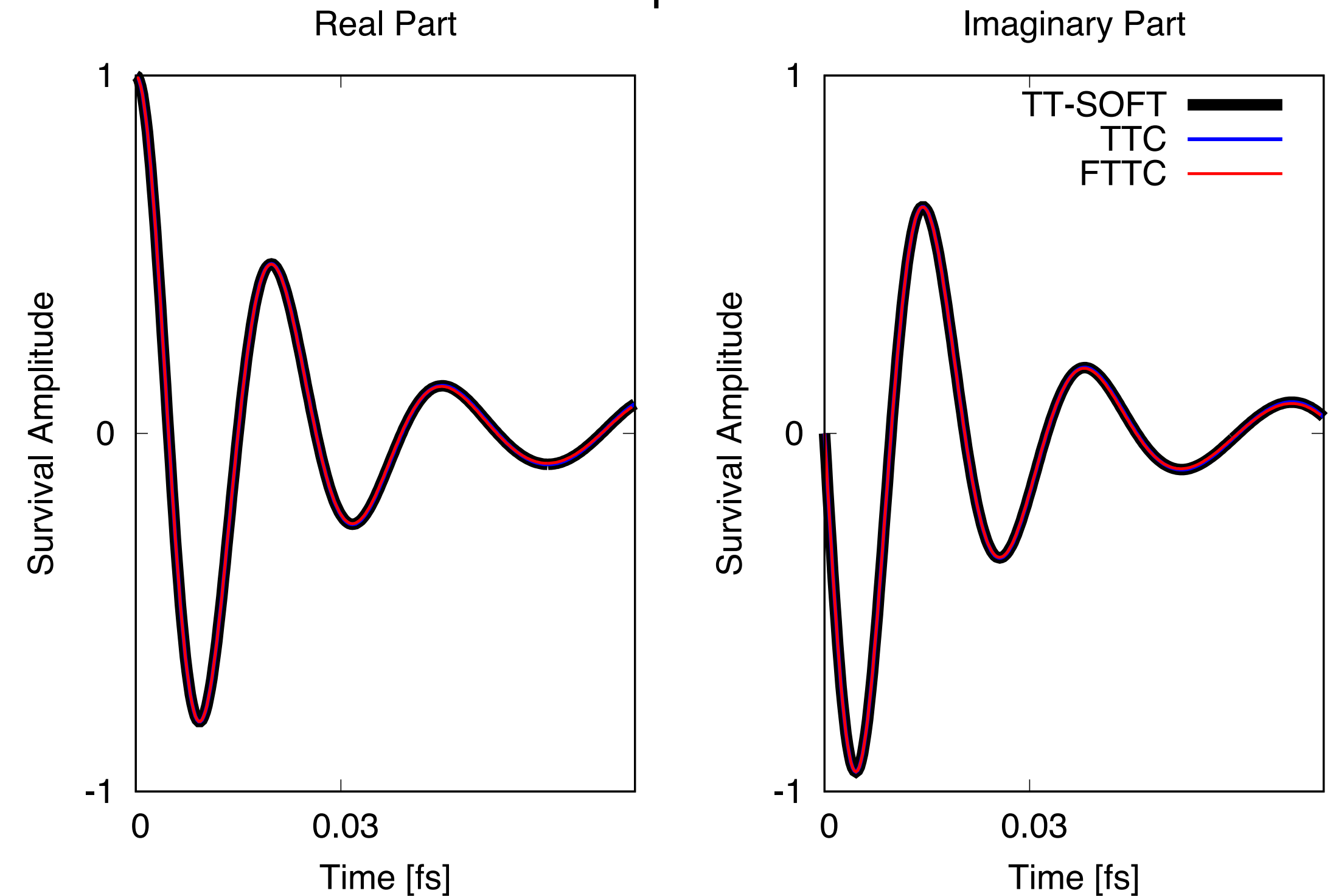


# COUPLED HYDROGEN BONDING IN DNA

Probability Density Dynamics, Anharmonically Coupled Bath



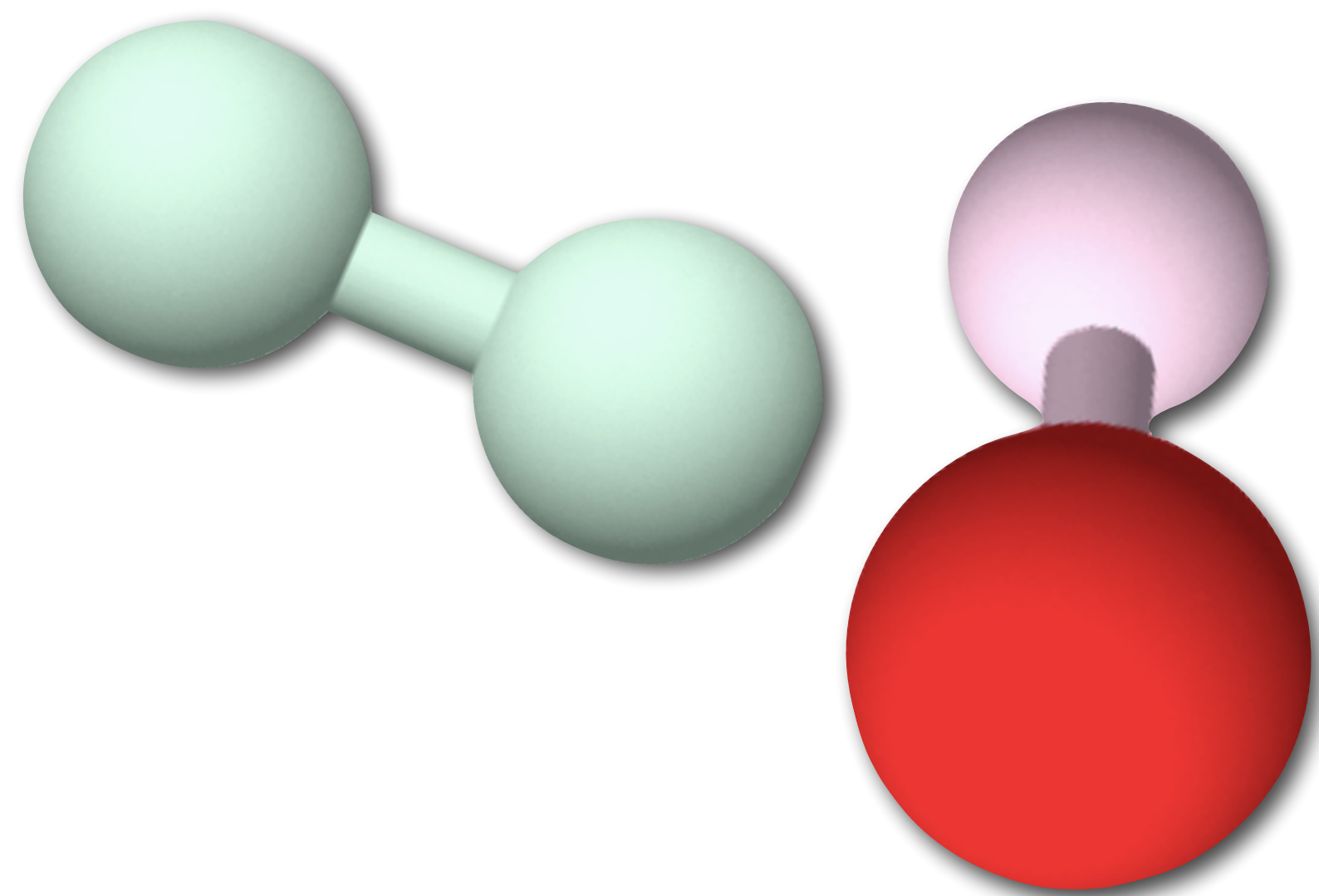
Survival Amplitude



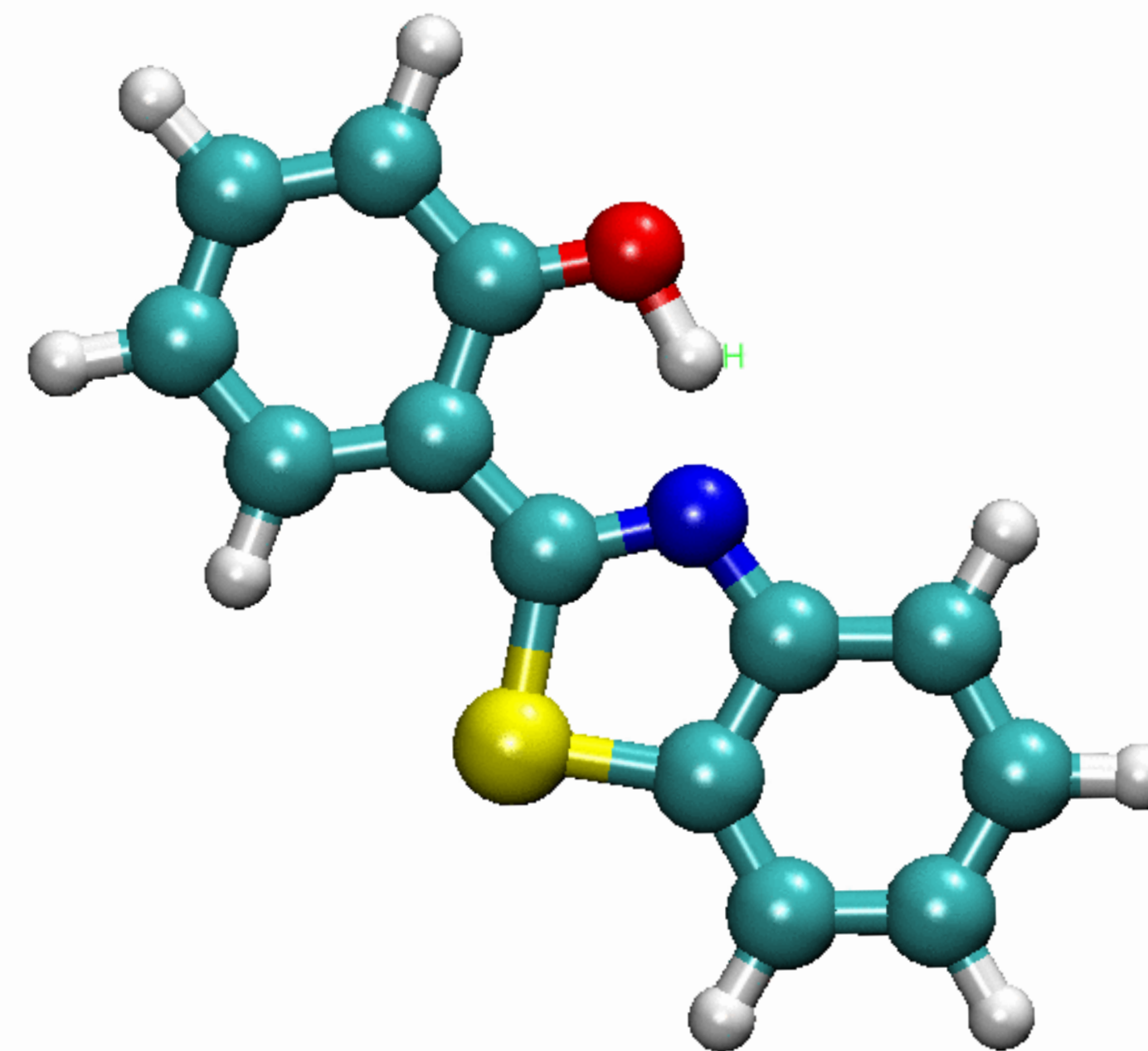
**FTTC successfully determines molecular dynamics  
even with significant coupling of atomic motion between modes.**

# TENSOR-TRAINS FOR UV-PUMP/X-RAY PROBE SPECTROSCOPY

Largest System Investigable with Standard Fixed-Grid SOFT Dynamics



Tensor-Train Split-Operator Fourier Transform (TT-SOFT) Dynamics

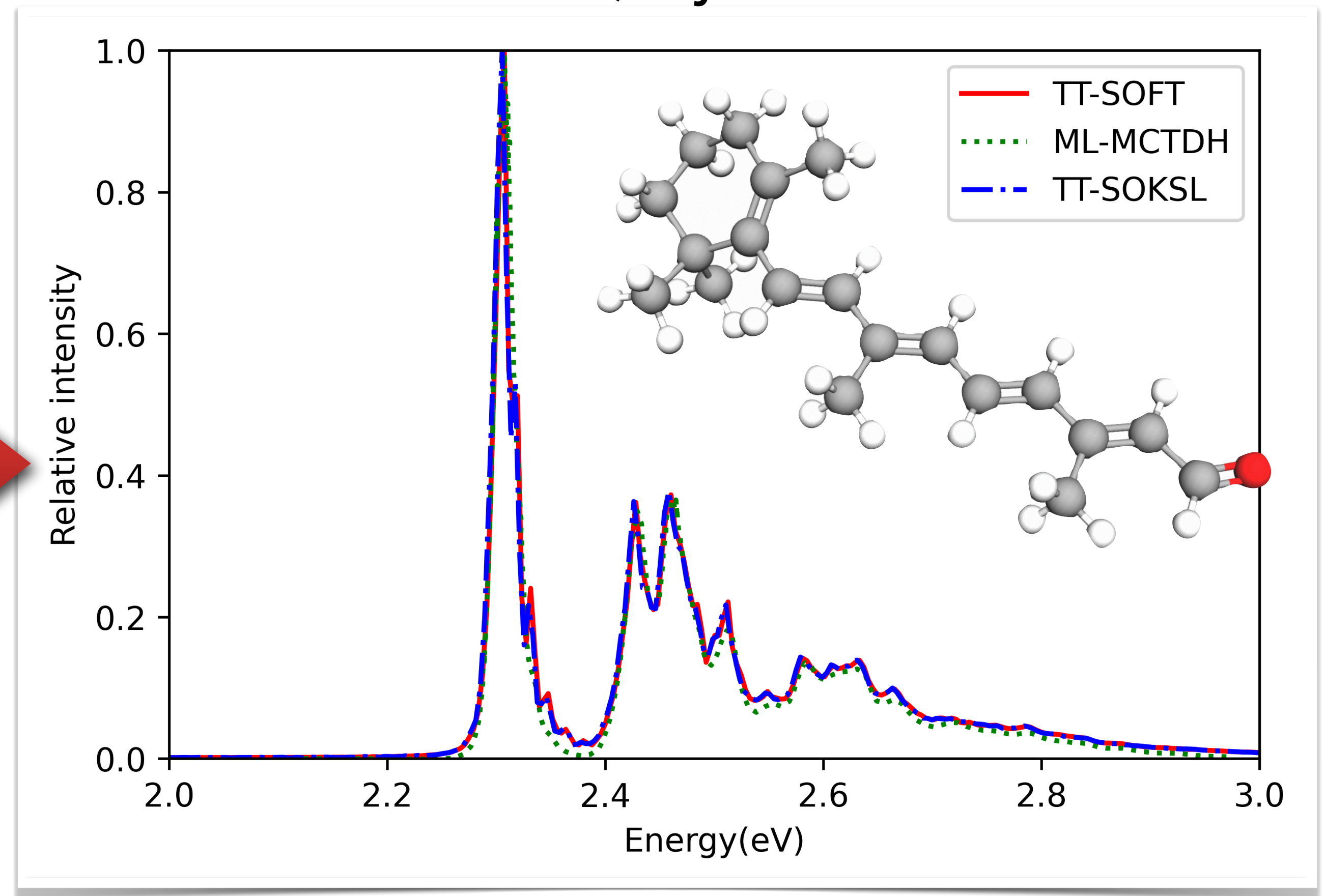
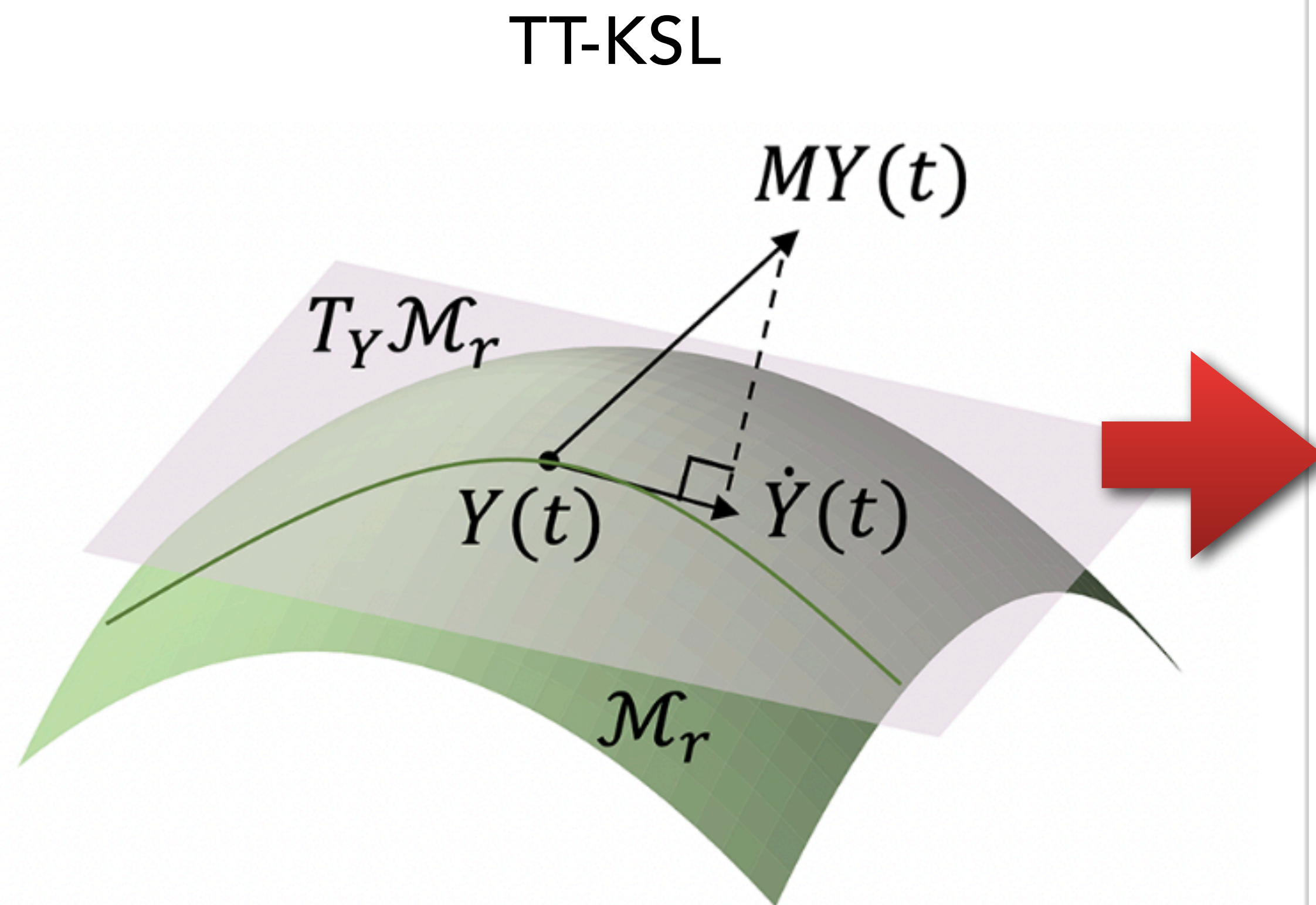


J. A. Fleck Jr., A. Steiger, J. Comput. Phys. 47 (1982) 412.  
S. M. Greene, V. S. Batista, JCTC 13 (2017) 4034.

**Micheline B. Soley**,\* P. E. Videla,\* E. T. J. Nibbering, V. S. Batista, (2022) J. Phys. Chem. Lett., 18 (2022) 8254.

# NOVEL TENSOR-TRAIN MOLECULAR METHODS

## Tensor-Train Split-Operator KSL (TT-SOKSL) Dynamics

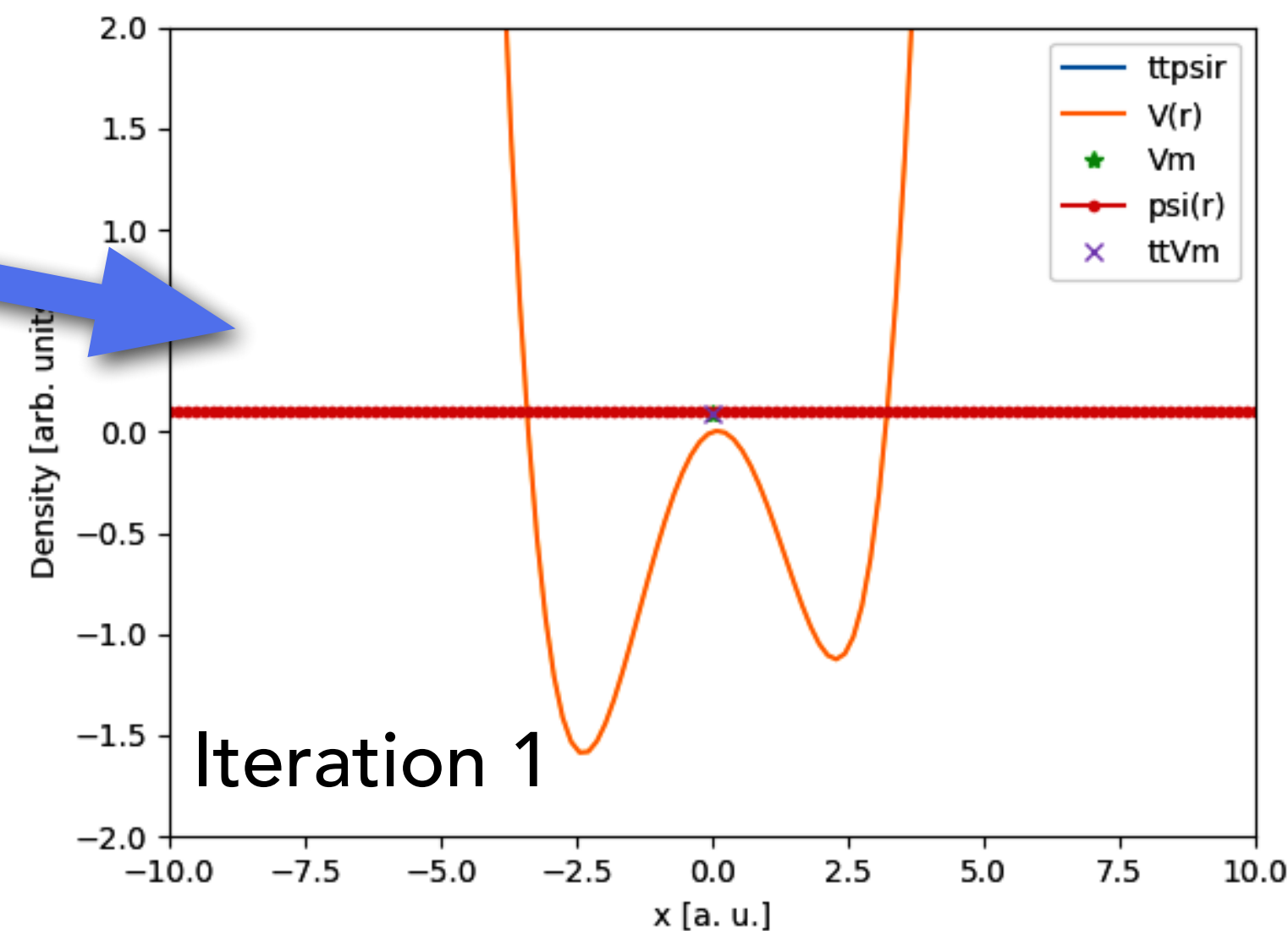


Micheline B. Soley, N. Lyu, V. S. Batista, JCTC, 18 (2022) 3327.

C. Lubich, From Quantum to Classical Molecular Dynamics: Reduced Models and Numerical Analysis, European Mathematical Society, 2009.

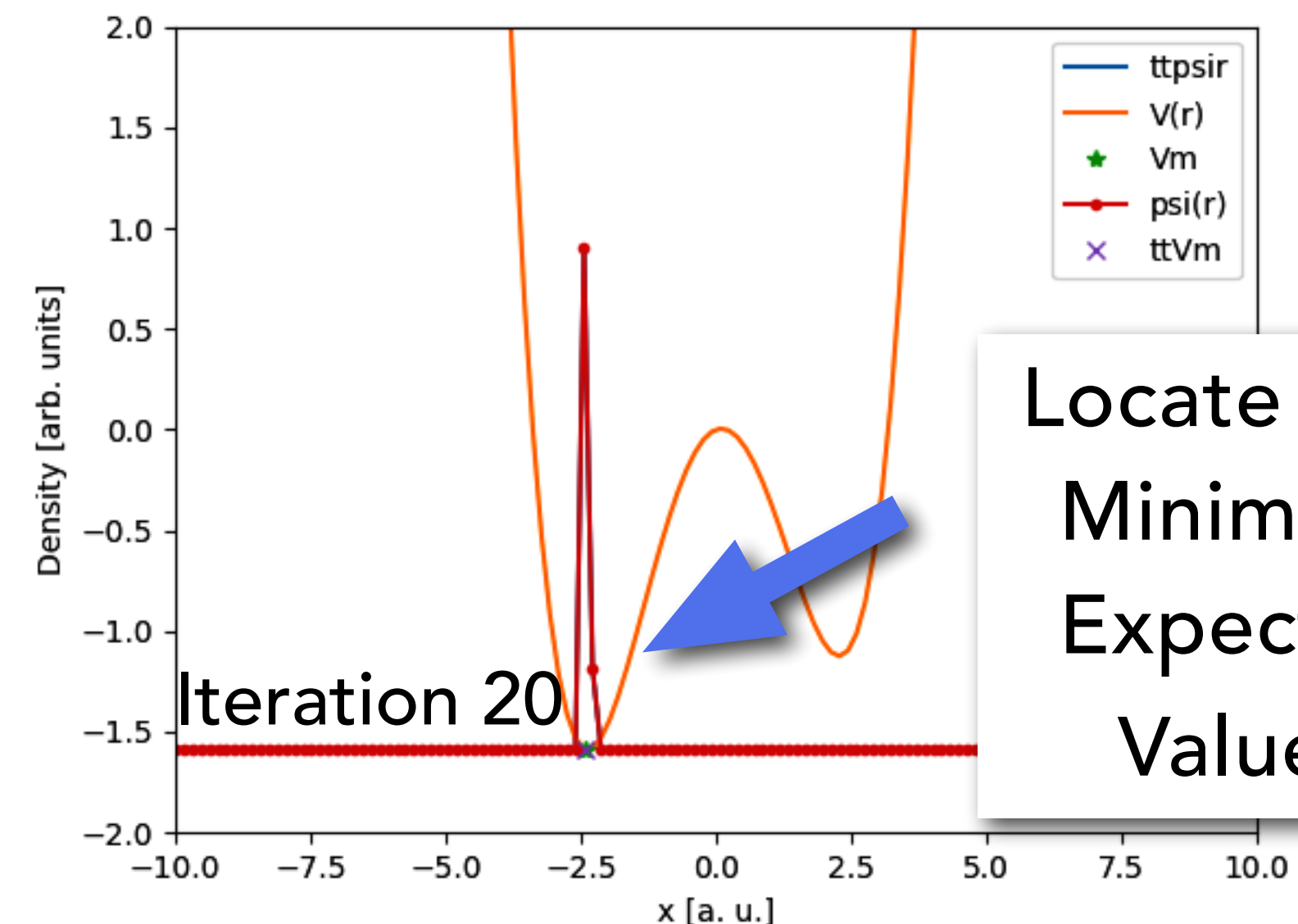
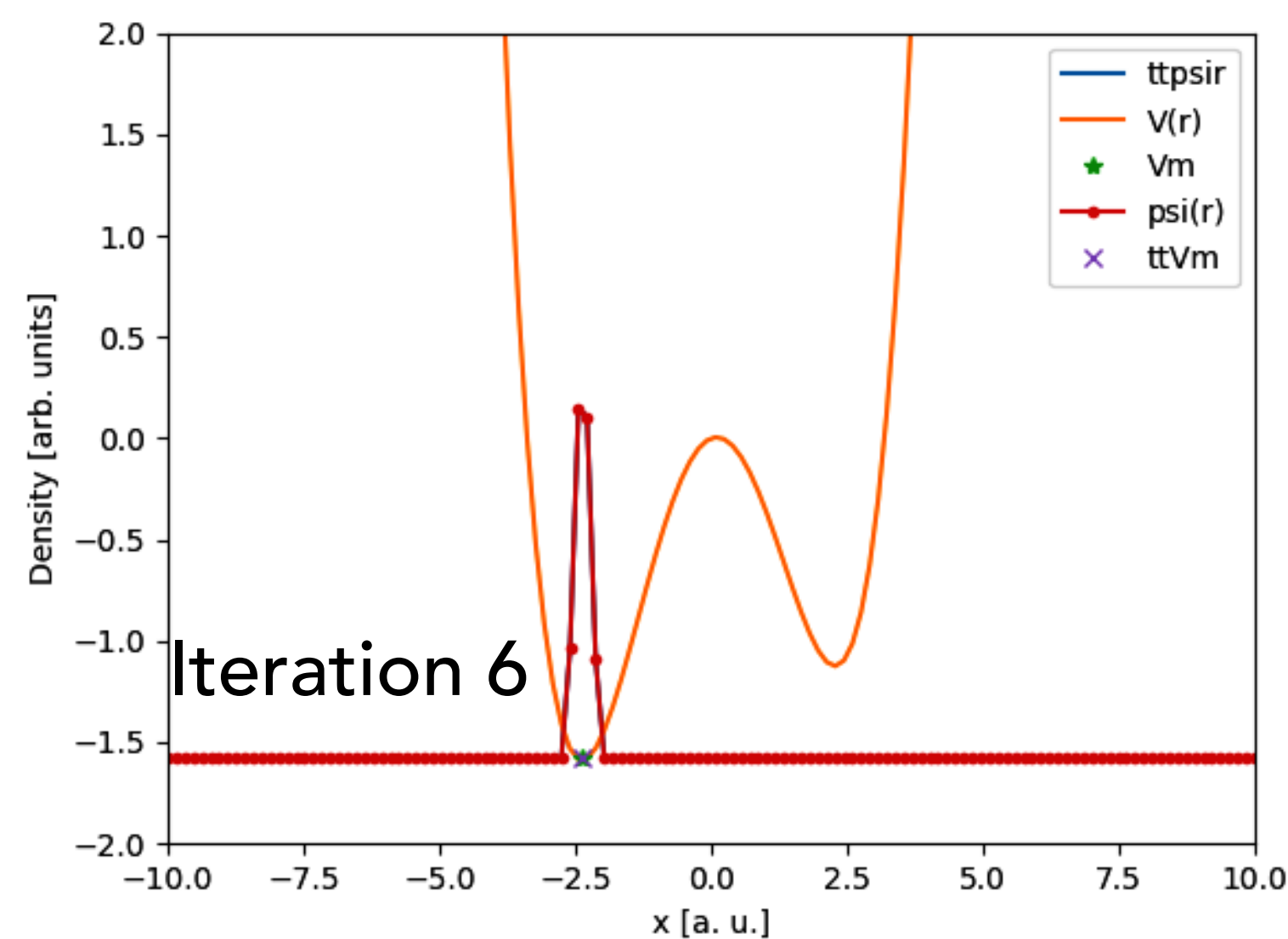
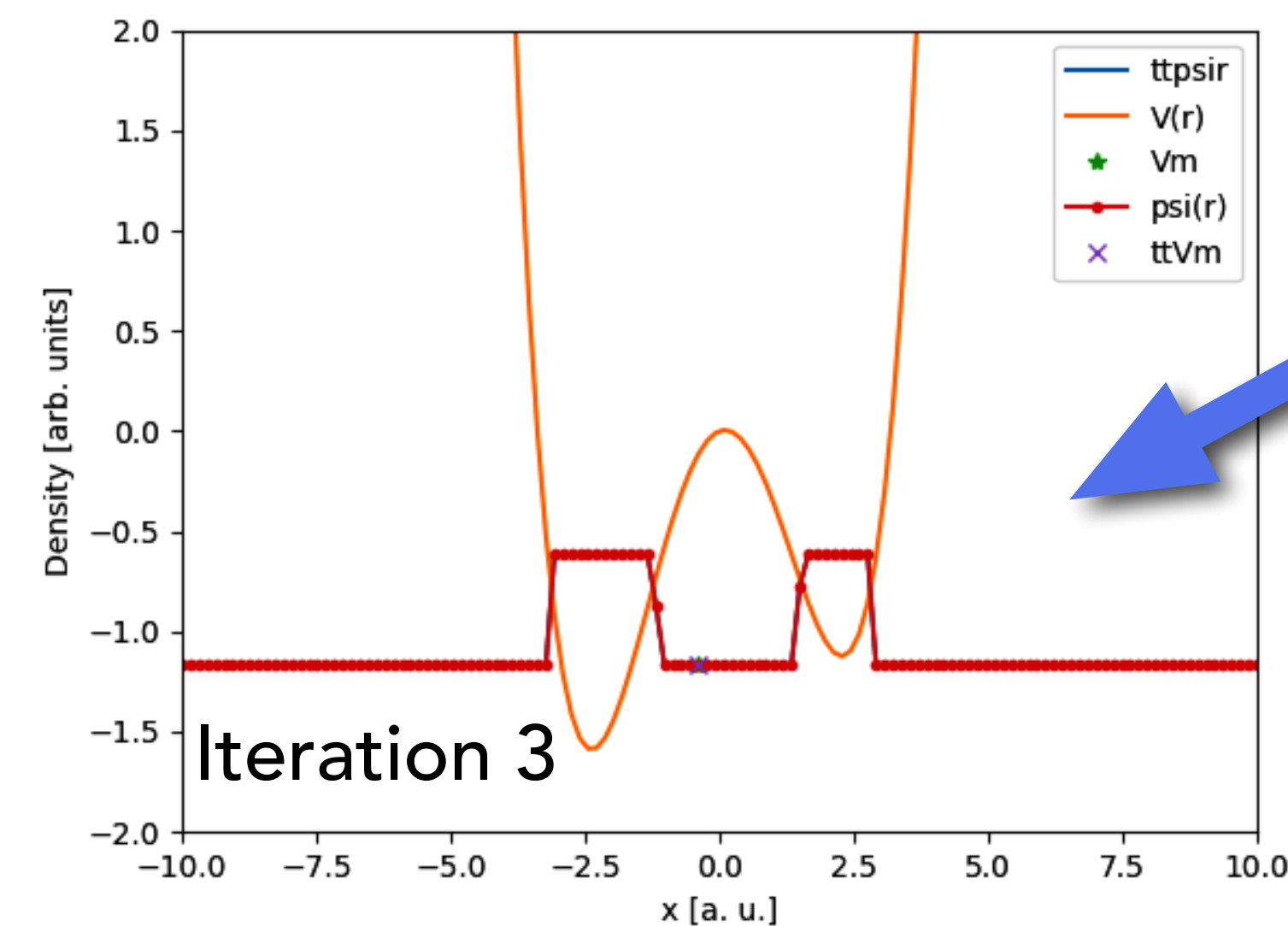
# TENSOR-TRAIN ITERATIVE POWER ALGORITHM (IPA)

Initialize  $\rho$  and  $V$  as Tensor Trains



Iteratively Apply Oracle

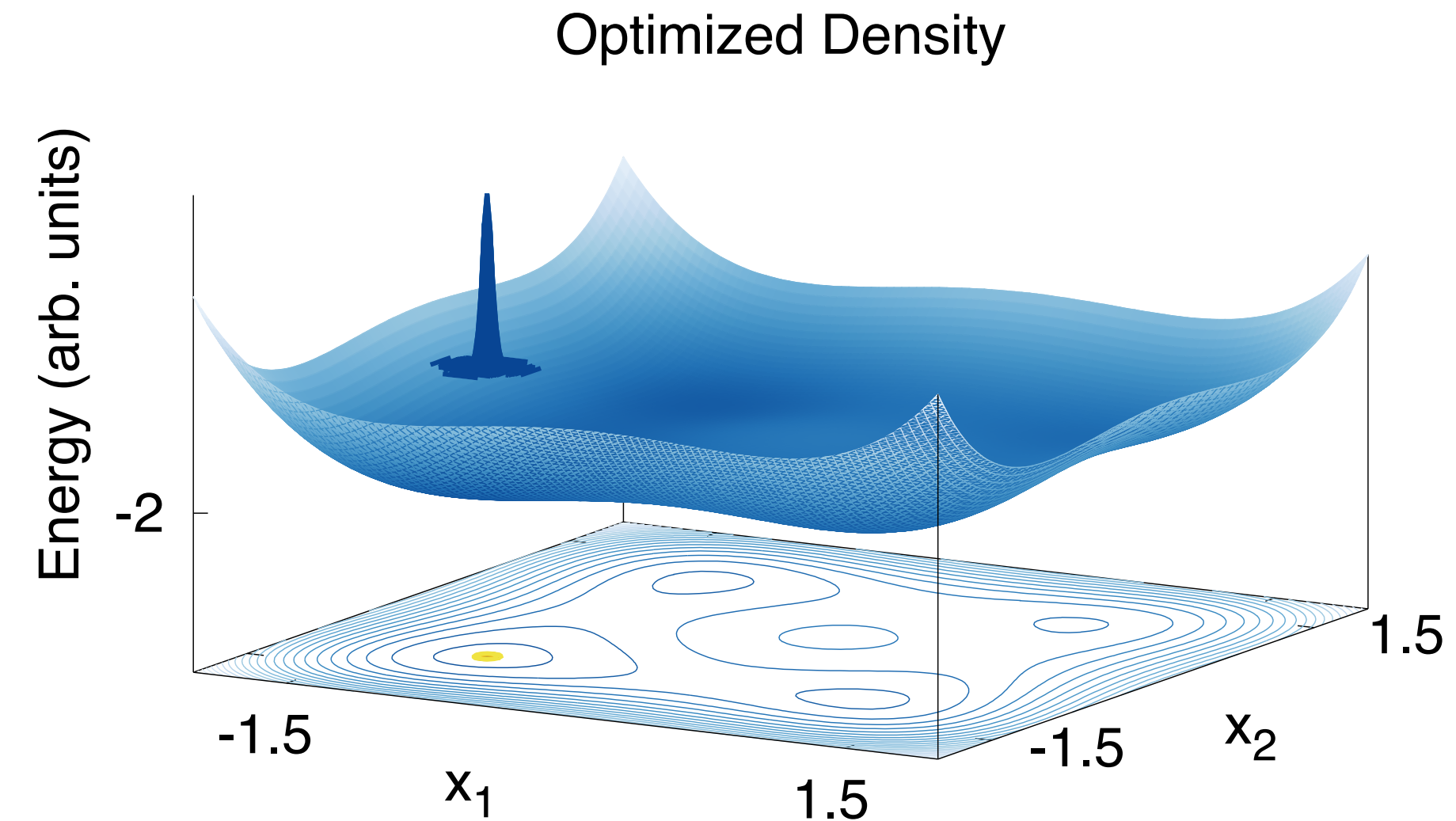
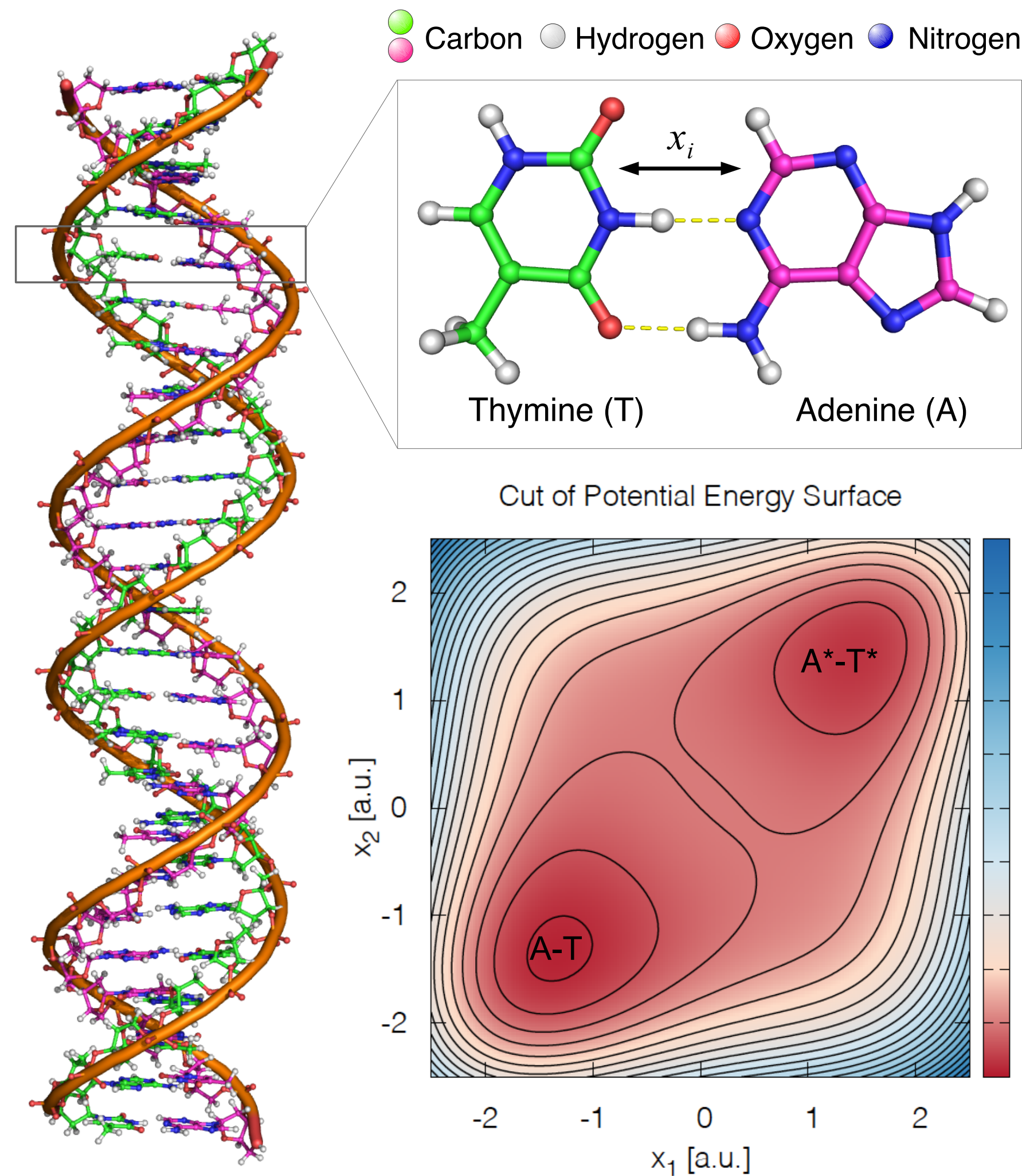
$$\rho(x) \rightarrow \mathcal{N}U(x)\rho(x)$$
$$U(x) = e^{-cV(x)}$$



Locate Global Minimum as Expectation Value  $\langle x \rangle$



# TENSOR-TRAIN ITERATIVE POWER ALGORITHM (IPA) FOR HYDROGEN BOND CONFIGURATIONS



IPA enables identification of minimum energy molecular structure in up to **400 dimensions** beyond the capabilities of traditional grid-based approaches.

# TENSOR-TRAIN IPA FOR PRIME FACTORIZATION OF

N=706851632784678312266808500466226101921669464385547527413569690763875796535401970800670478642897888271976359744482  
562359245255871481739623795579868353636314461966874443074041408313128692006363959785724202562228073836999106855285692  
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2583939955697859360862764310036106686693705353525252656142323642791967328180835792001

**2,773 DIGITS (MORE THAN 9,212 BITS)**

# NUMBER OF ITERATIONS

required to reach optimal result with 50% certainty (à la Grover's algorithm):

$$U = \text{diag}(\lambda_2, \dots, \lambda_2, \lambda_1, \lambda_2, \dots, \lambda_2) \in \mathbb{R}^{n \times n}, \quad 0 < \lambda_2 < \lambda_1$$

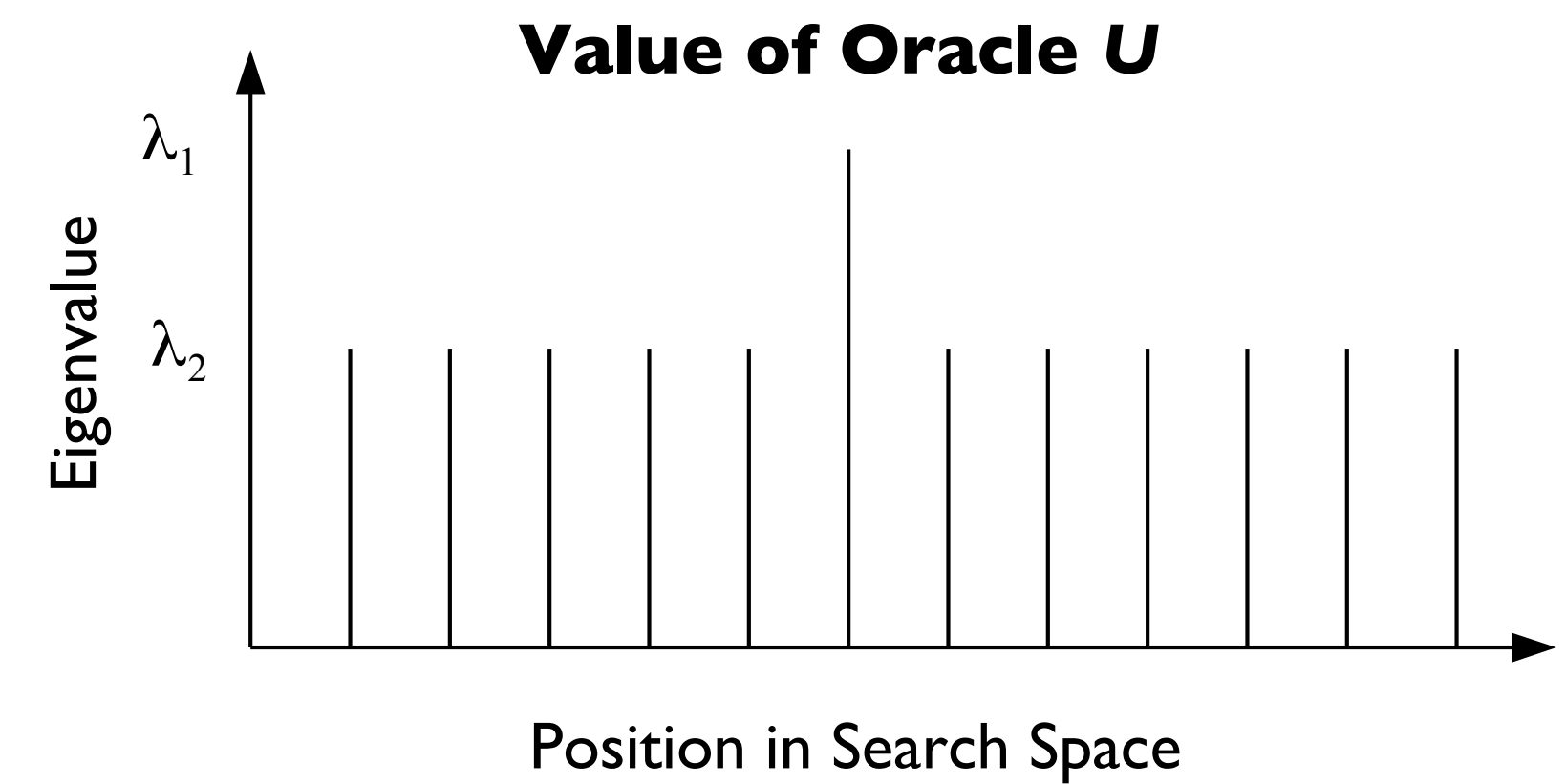
$$\rho_0 = \frac{1}{n}(1, \dots, 1) \in \mathbb{R}^n$$

$$\rho_k = \frac{U^k \rho_0}{\|U^k \rho_0\|_1} \quad \frac{\rho_{k, \min}}{\rho_{k, \max}} = \left( \frac{\lambda_2}{\lambda_1} \right)^k$$

$$1 = \|\rho_k\| = \rho_{k, \max} + (n - 1)\rho_{k, \min}$$

$$\rho_{k, \max} = \frac{1}{1 + (n - 1) \cdot (\lambda_2/\lambda_1)^k}$$

$$\frac{1}{2} \leq \frac{1}{1 + (n - 1) \cdot (\lambda_2/\lambda_1)^k}$$



$$k \geq \frac{\log(n - 1)}{\log(\lambda_1/\lambda_2)}$$

IPA requires fewer iterations than foremost quantum approach.

Micheline B. Soley, P. Bergold, V. S. Batista, JCTC 17 (2021) 3280.

L. K. Grover, Proceedings, 28th Annual ACM Symposium on the Theory of Computing, May 1996, 212.

# TENSOR-TRAIN EFFICIENT QUANTUM COMPUTING ALGORITHMS FOR MOLECULAR SYSTEMS

This computational efficiency raises the question — can IPA be used to develop more efficient quantum computing algorithms for molecular systems?

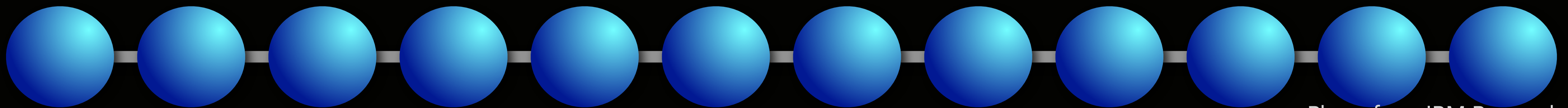
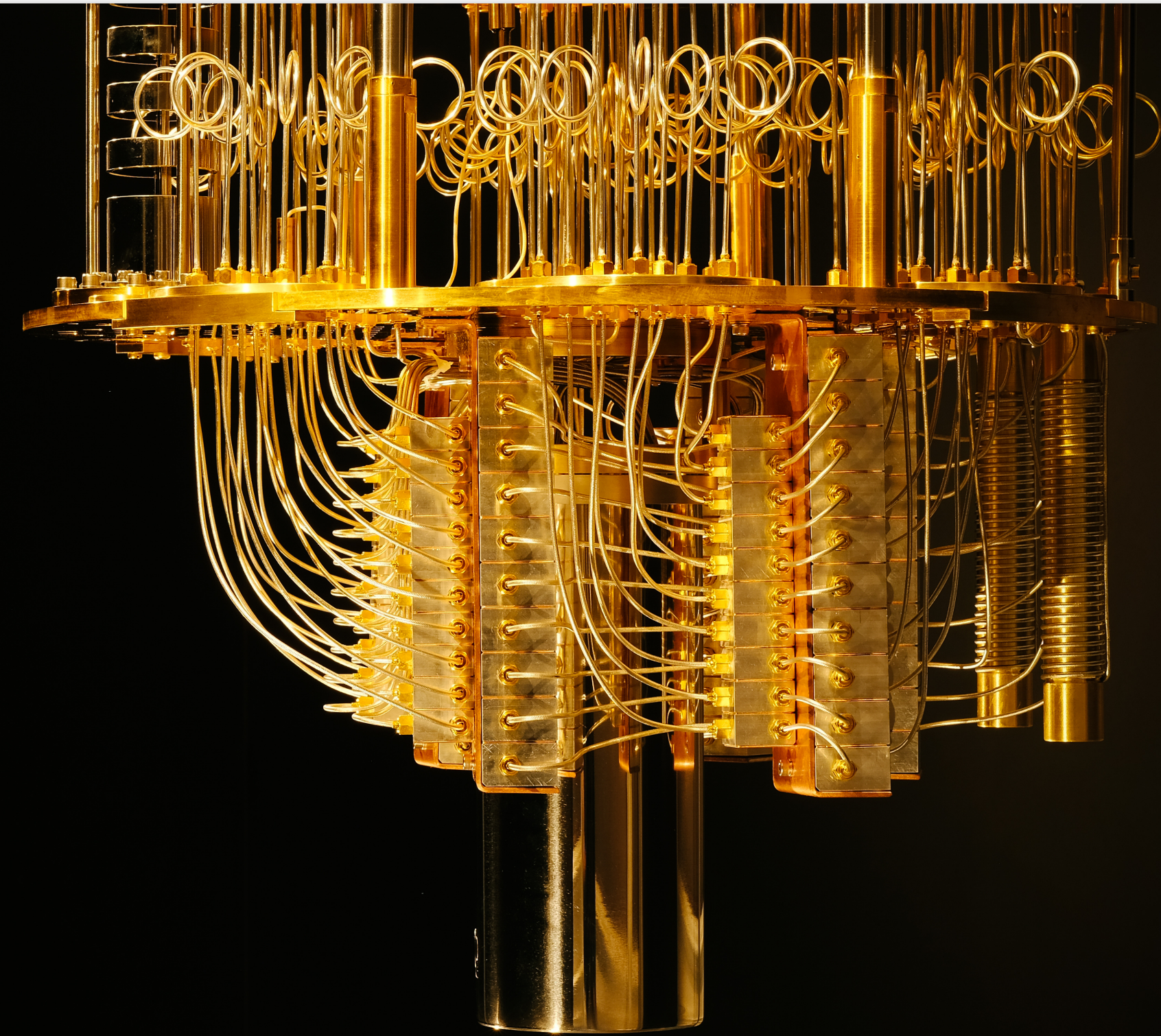
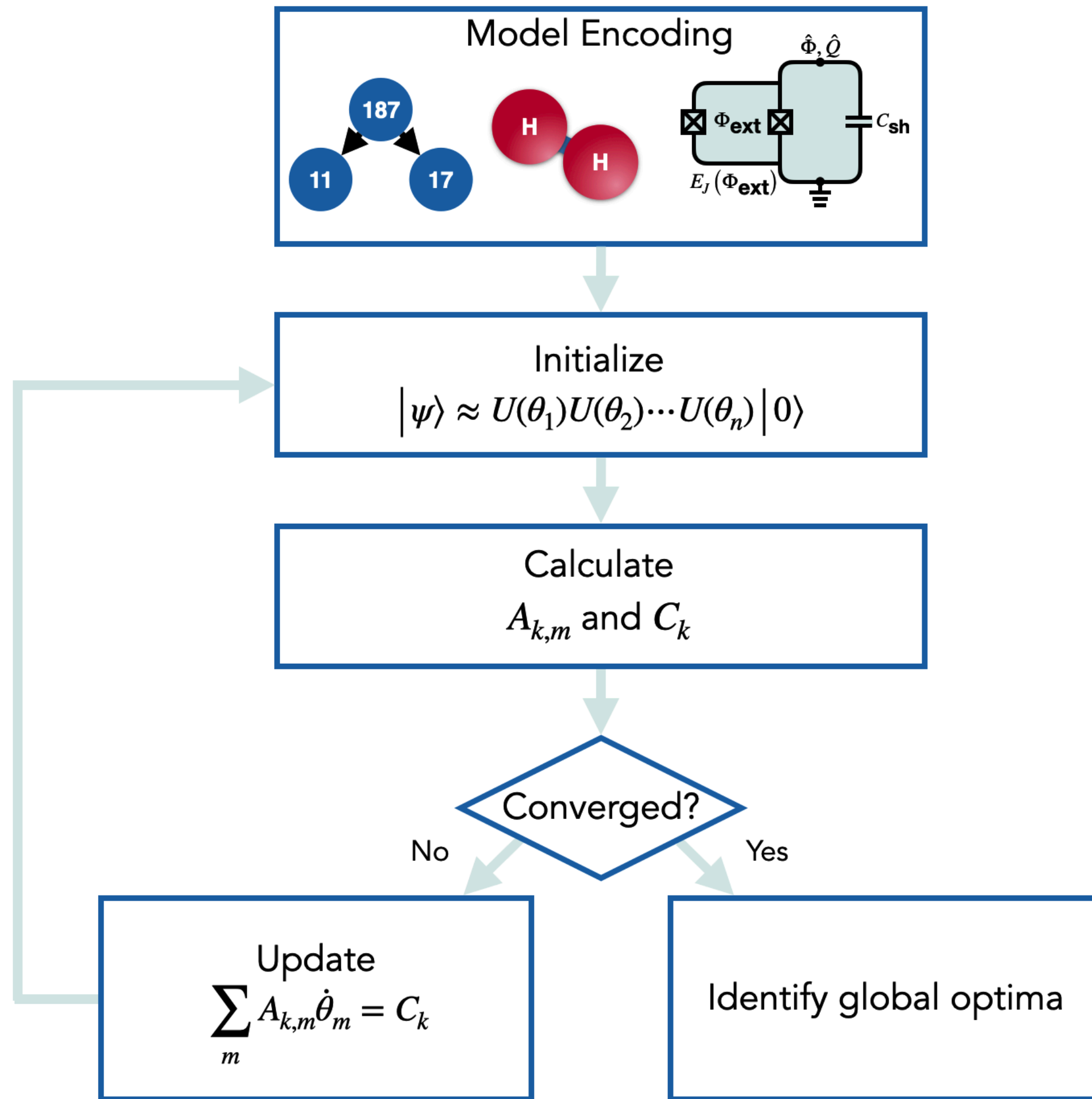


Photo from IBM Research.

T. H. Kyaw\*, **Micheline B. Soley**\*, B. Allen, P. Bergold, C. Sun, V. S. Batista, A. Aspuru-Guzik, (2022) arXiv:2208.10470v1.  
F. Arute, et al. Science 369 (2020) 1084.



# QUANTUM ITERATIVE POWER ALGORITHM (QIPA)



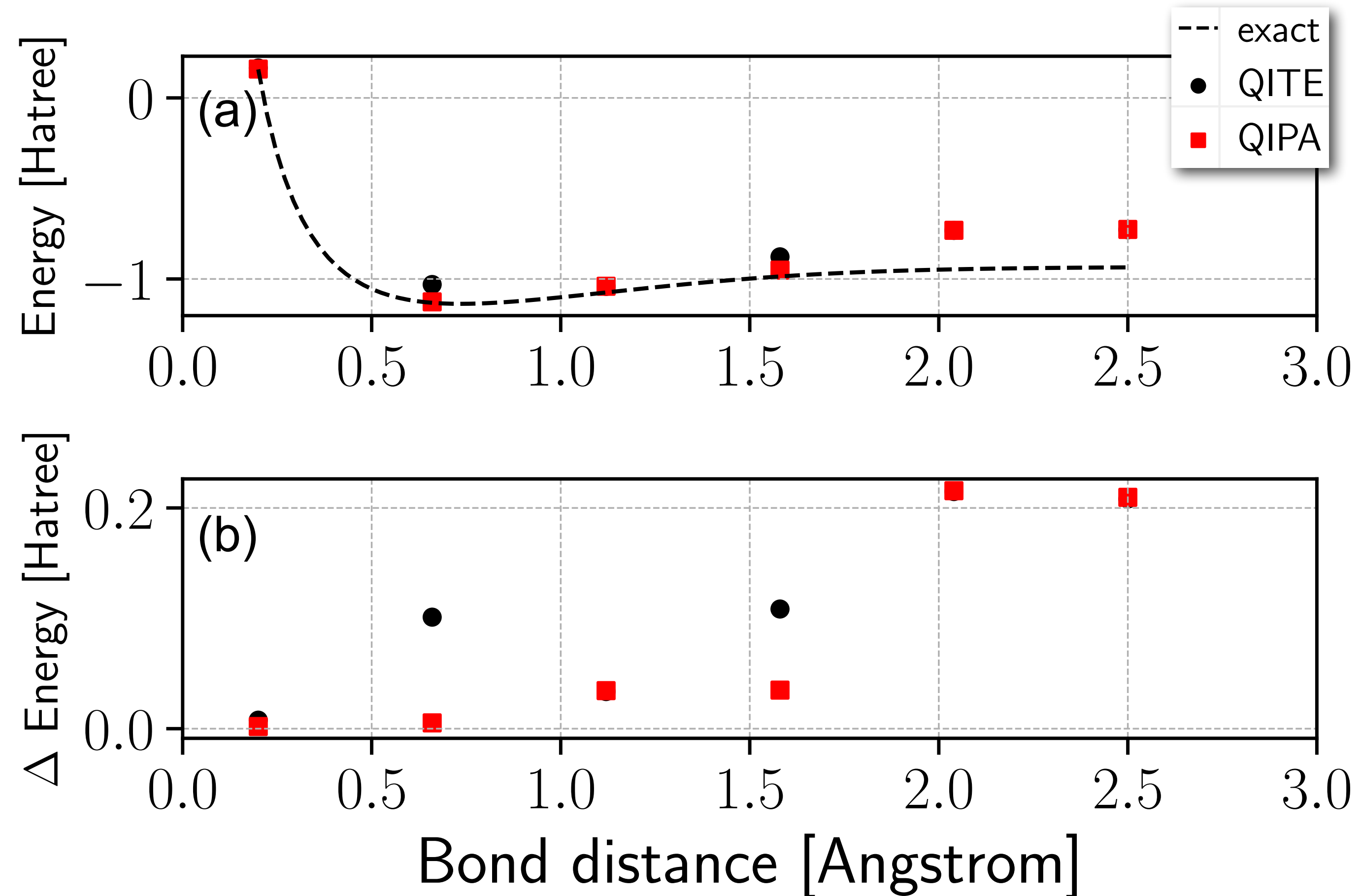
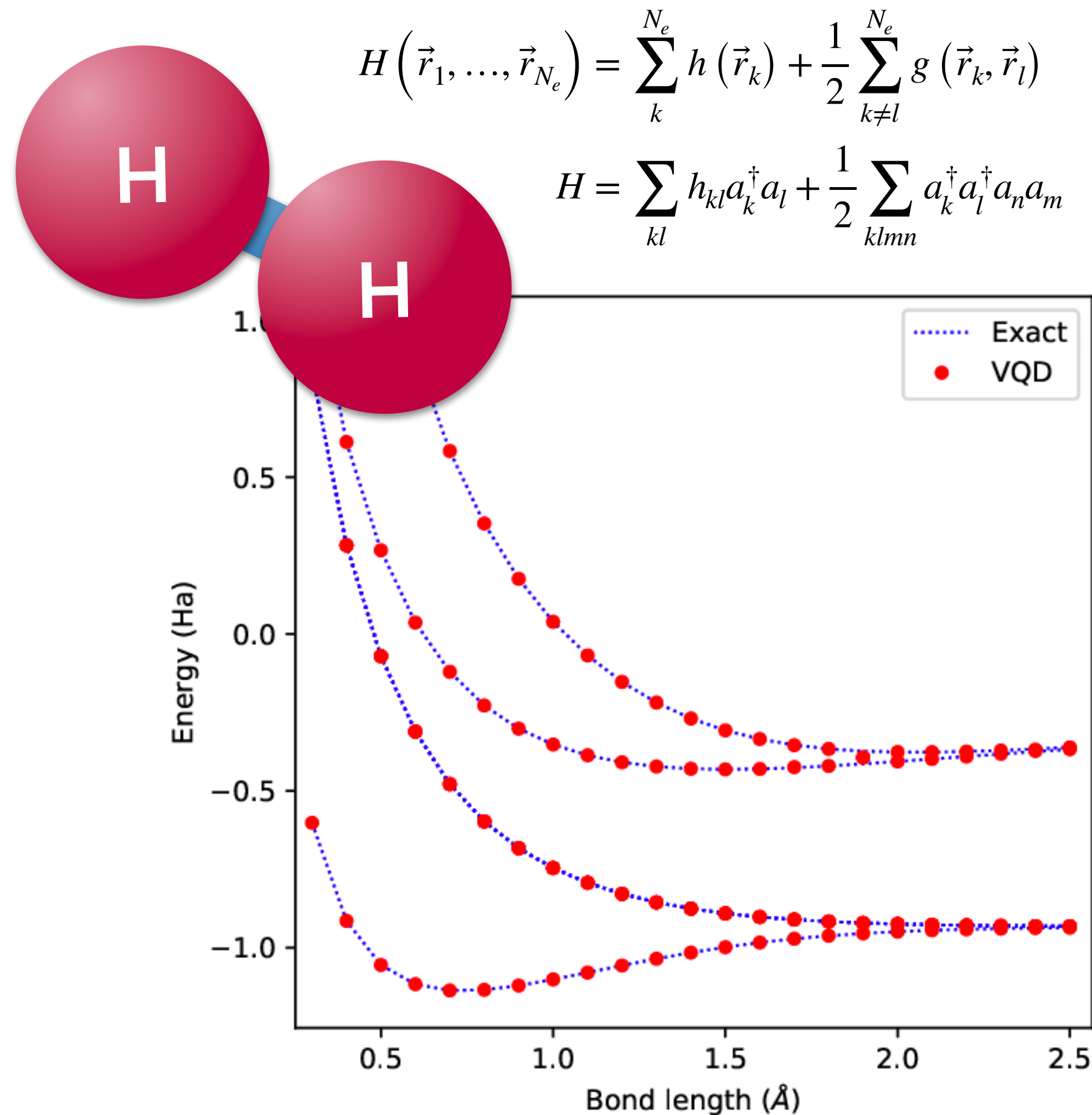
Connection to Cooling Schemes in Quantum Annealing

Accelerated Determination of Global Minima

First Instance of IPA on Quantum Computer

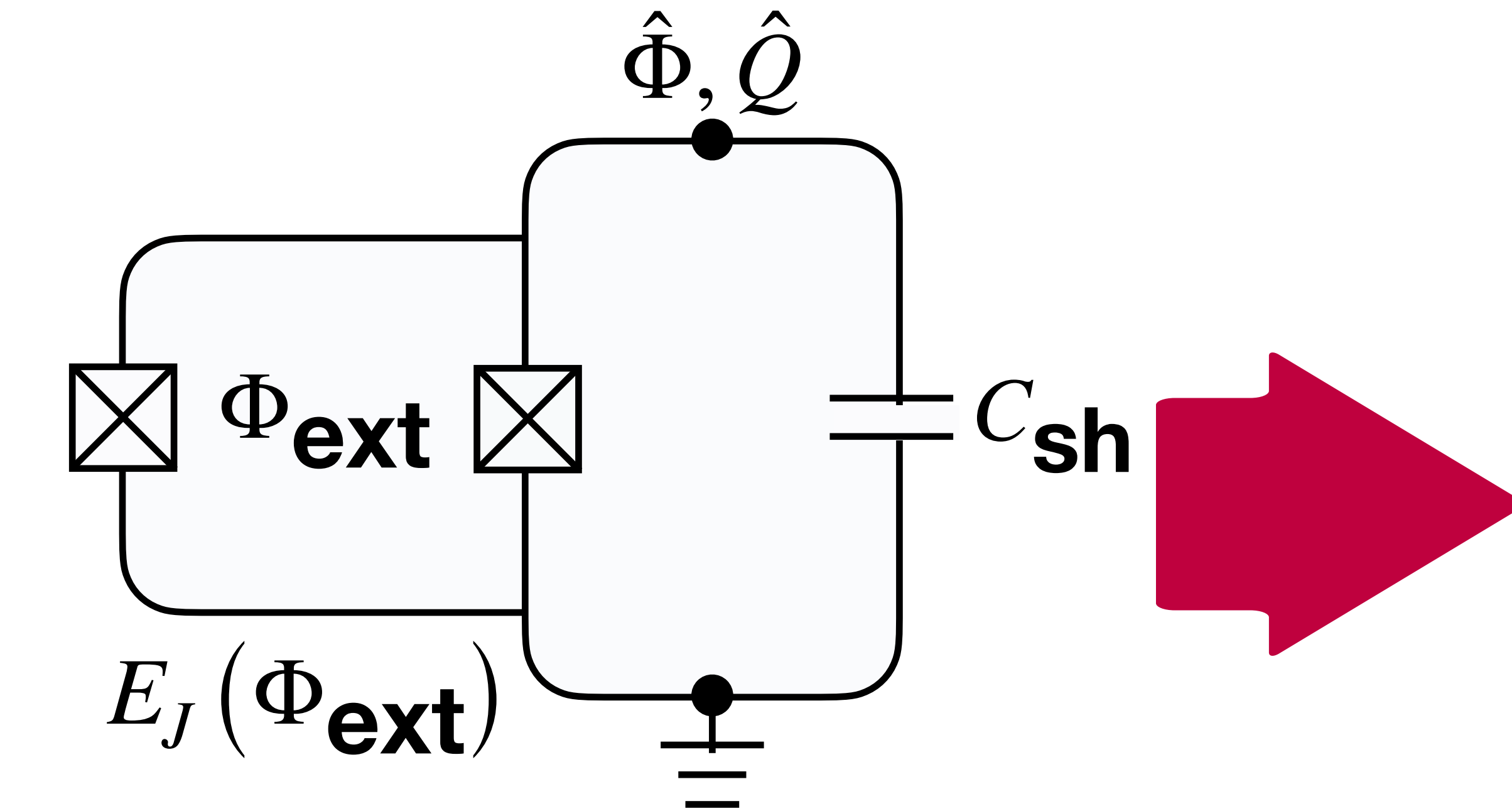
# QIPA FOR ELECTRONIC STRUCTURE THEORY

QIPA **successfully** identifies the ground state energies of H<sub>2</sub> to high accuracy

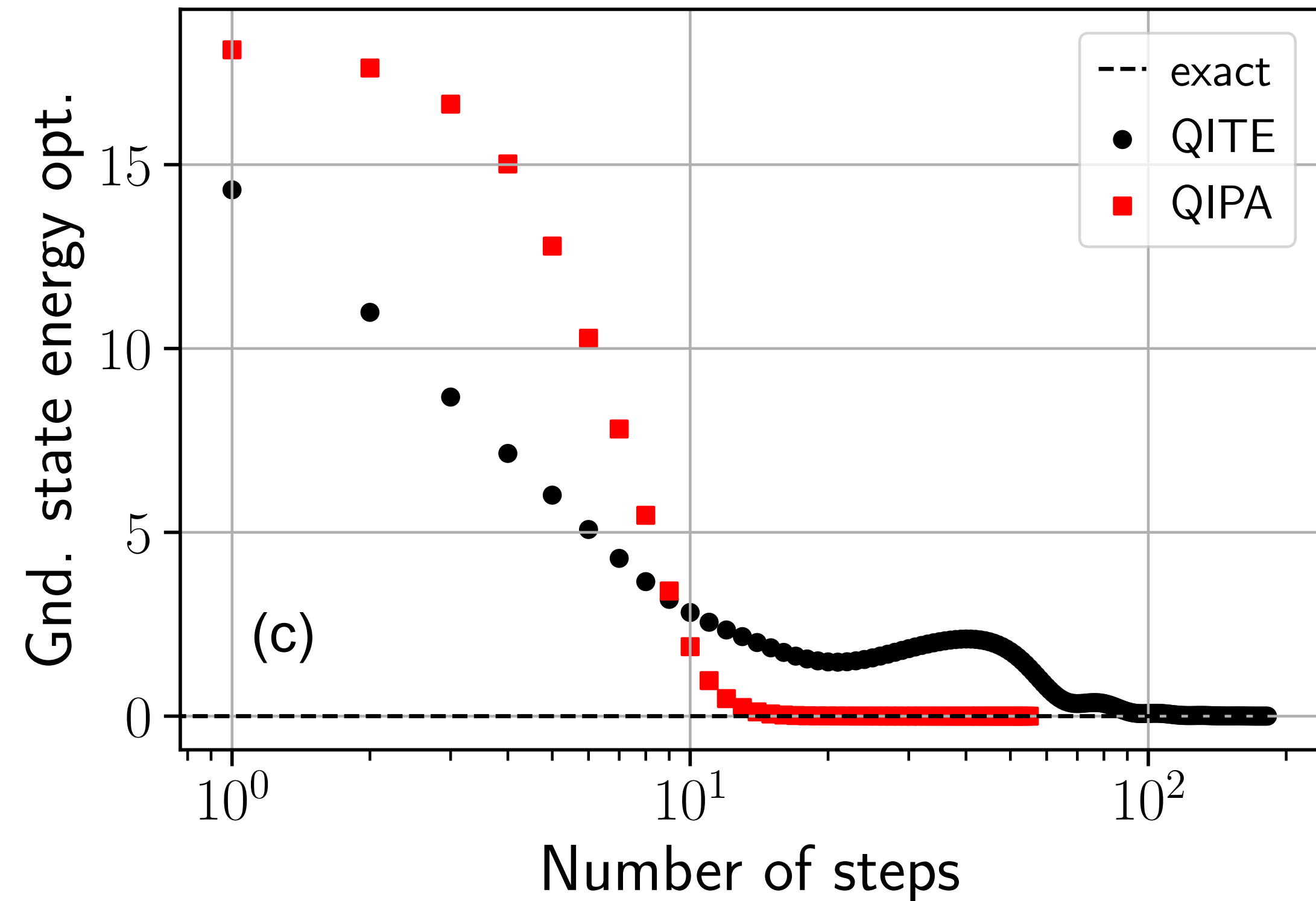


# QIPA FOR QUANTUM COMPUTER DESIGN

First dynamical optimization of quantum processor design via quantum computing



$$\hat{H}_{M\text{-transmon}} = 2e^2 \sum_{i,j=1}^M (\mathbf{C}^{-1})_{ij} \hat{N}_i \hat{N}_j - 2 \sum_{i=1}^M E_{J,i} |\cos(2\pi f_i)| \cos \hat{\phi}_i$$



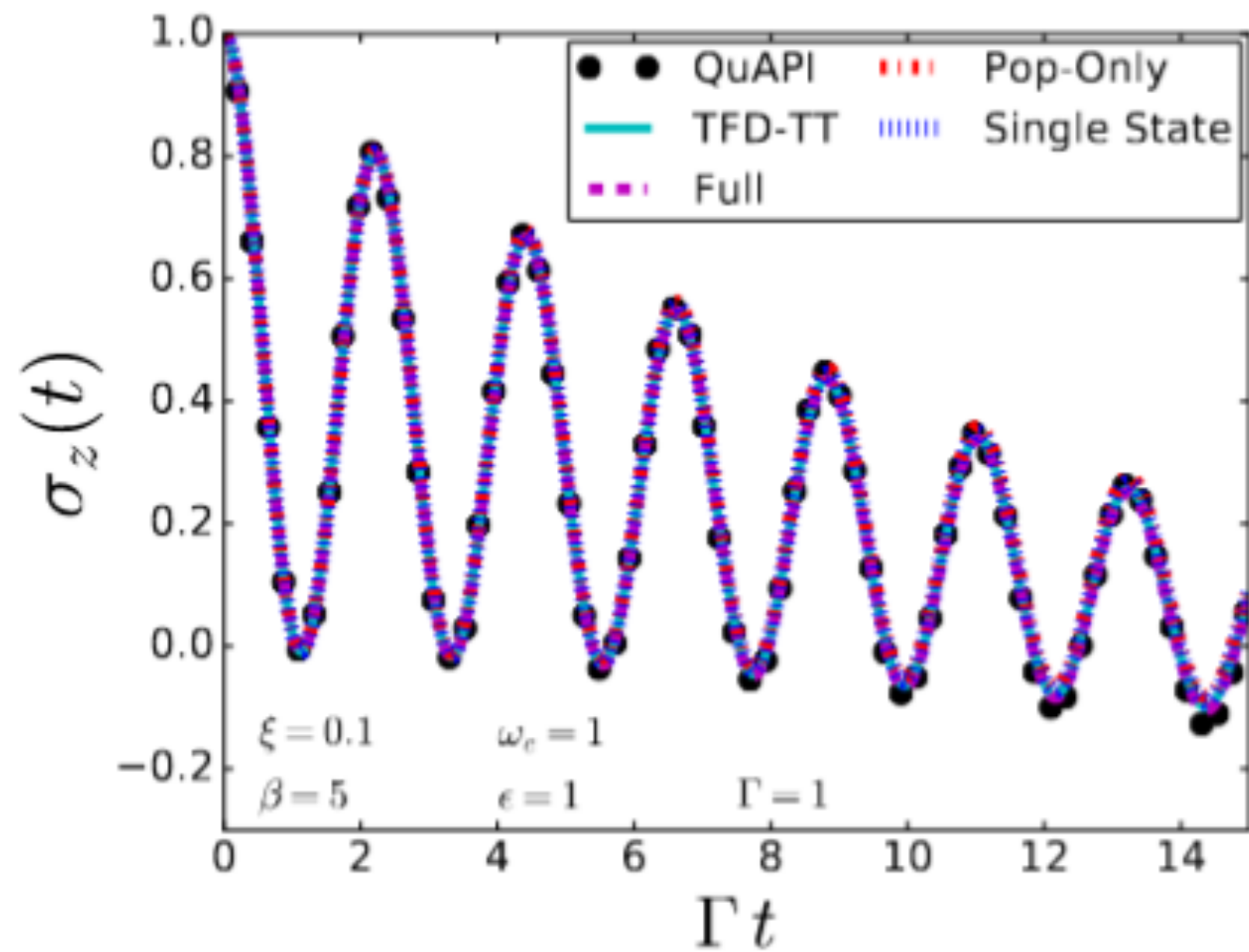
**Advantage** of QIPA over QITE for early steps over a broad parameter range

T. H. Kyaw\*, [Micheline B. Soley](#)\*, B. Allen, P. Bergold, C. Sun, V. S. Batista, A. Aspuru-Guzik, (2022) arXiv:2208.10470v1.

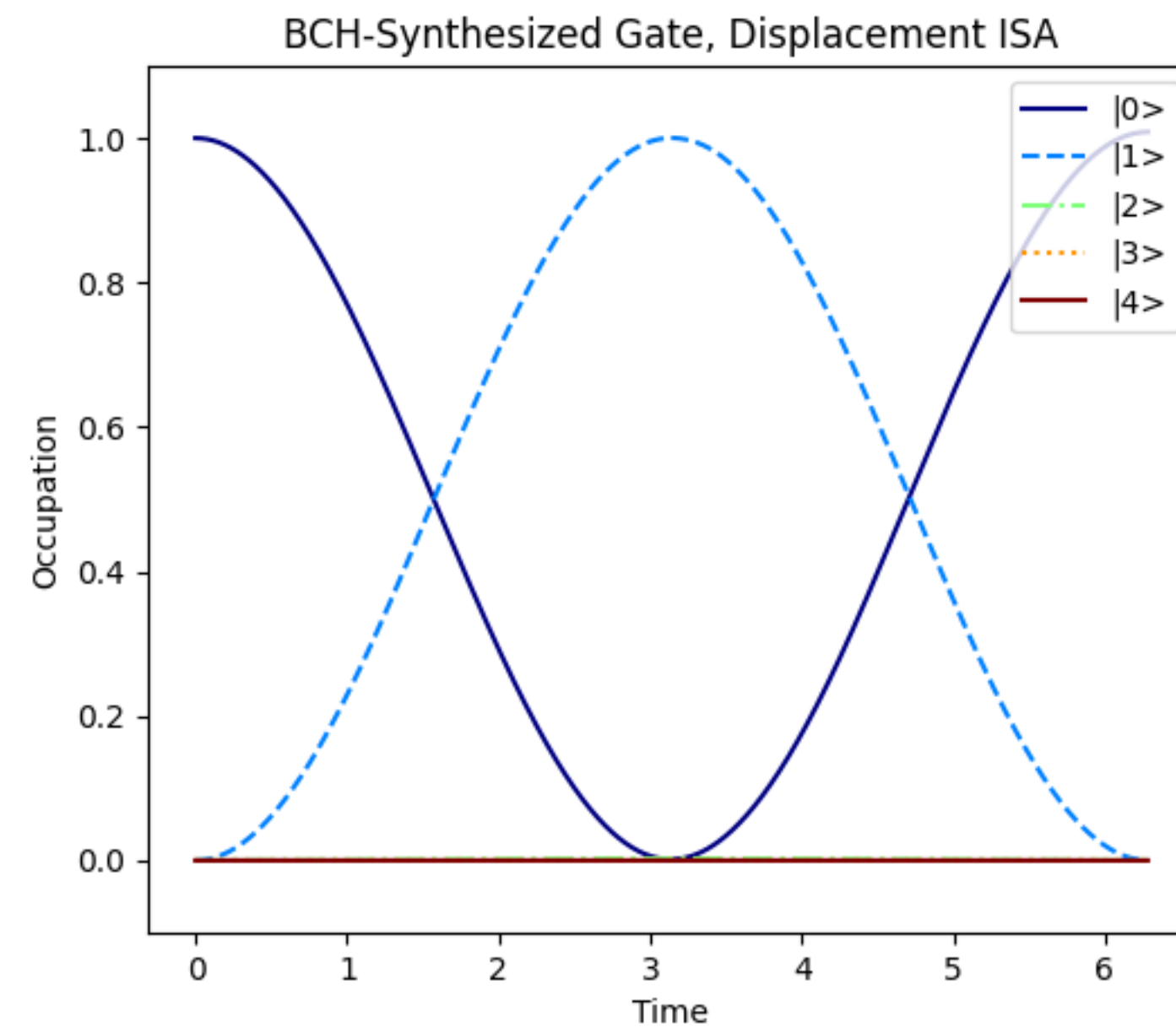
T. H. Kyaw\*, Ti. Menke\*, et al. (2021) arXiv:2006.03070v3.

# APPLICATIONS OF TENSOR-NETWORK AND QUANTUM COMPUTING APPROACHES

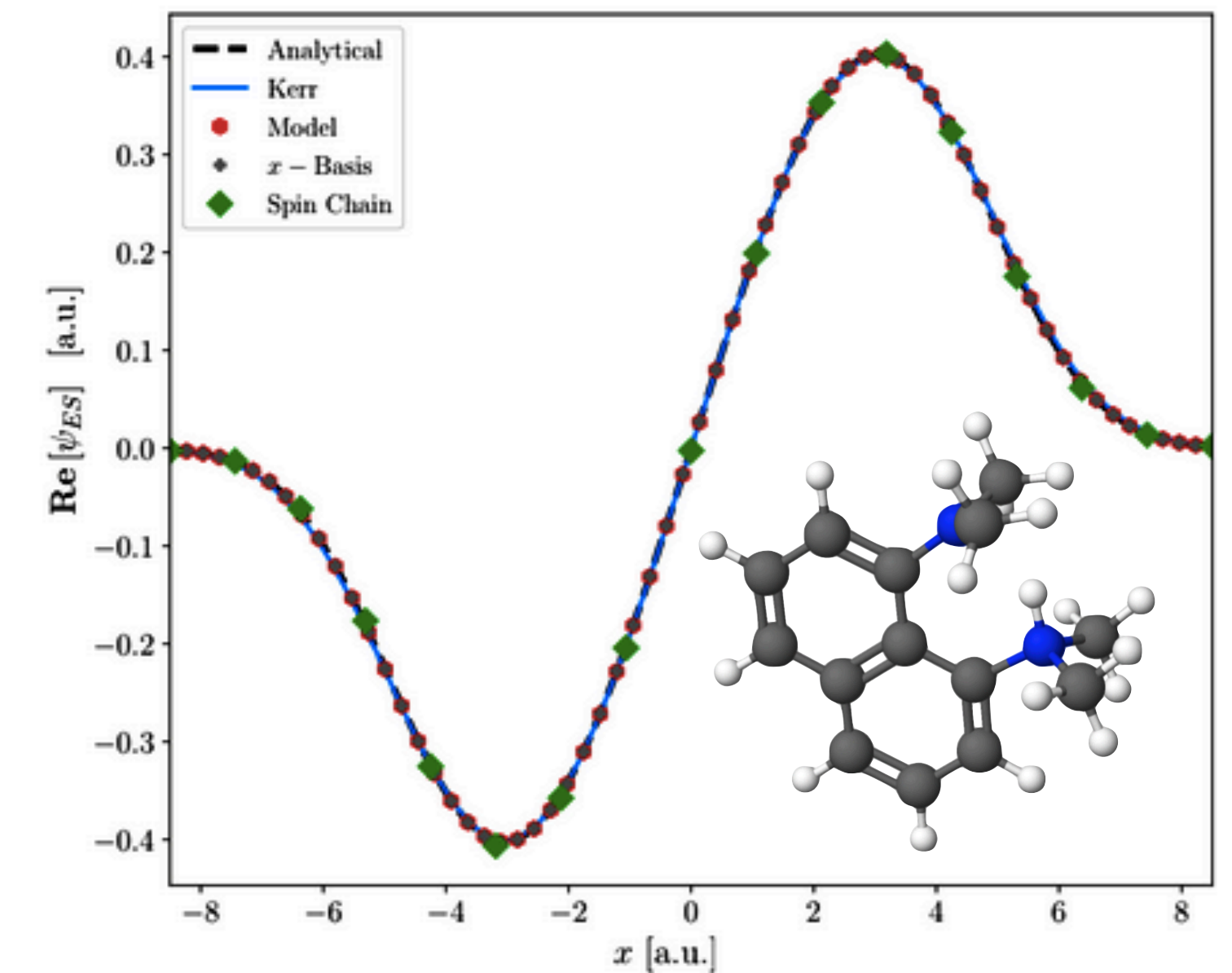
## OPEN QUANTUM DYNAMICS



## INSTRUCTION SET ARCHITECTURE



## EIGENSTATE DETERMINATION METHOD



C. Kang,\* **Micheline B. Soley**,\* E. Crane, S. M. Girvin, N. Wiebe, (2023) arXiv:2303.15542.

N. Lyu\*, E. Mulvihill\*, **Micheline B. Soley**, E. Geva, V. S. Batista, JCTC, 19 (2023) 1111.

T. H. Kyaw\*, **Micheline B. Soley**,\* B. Allen, P. Bergold, C. Sun, V. S. Batista, A. Aspuru-Guzik, (2022) arXiv:2208.10470v1.

Y. Wang, E. Mulvihill, Z. Hu, N. Lyu, S. Shivpuje, Y. Liu, **Micheline B. Soley**, E. Geva, V. S. Batista, S. Kais, 2022, arXiv:2209.04956.

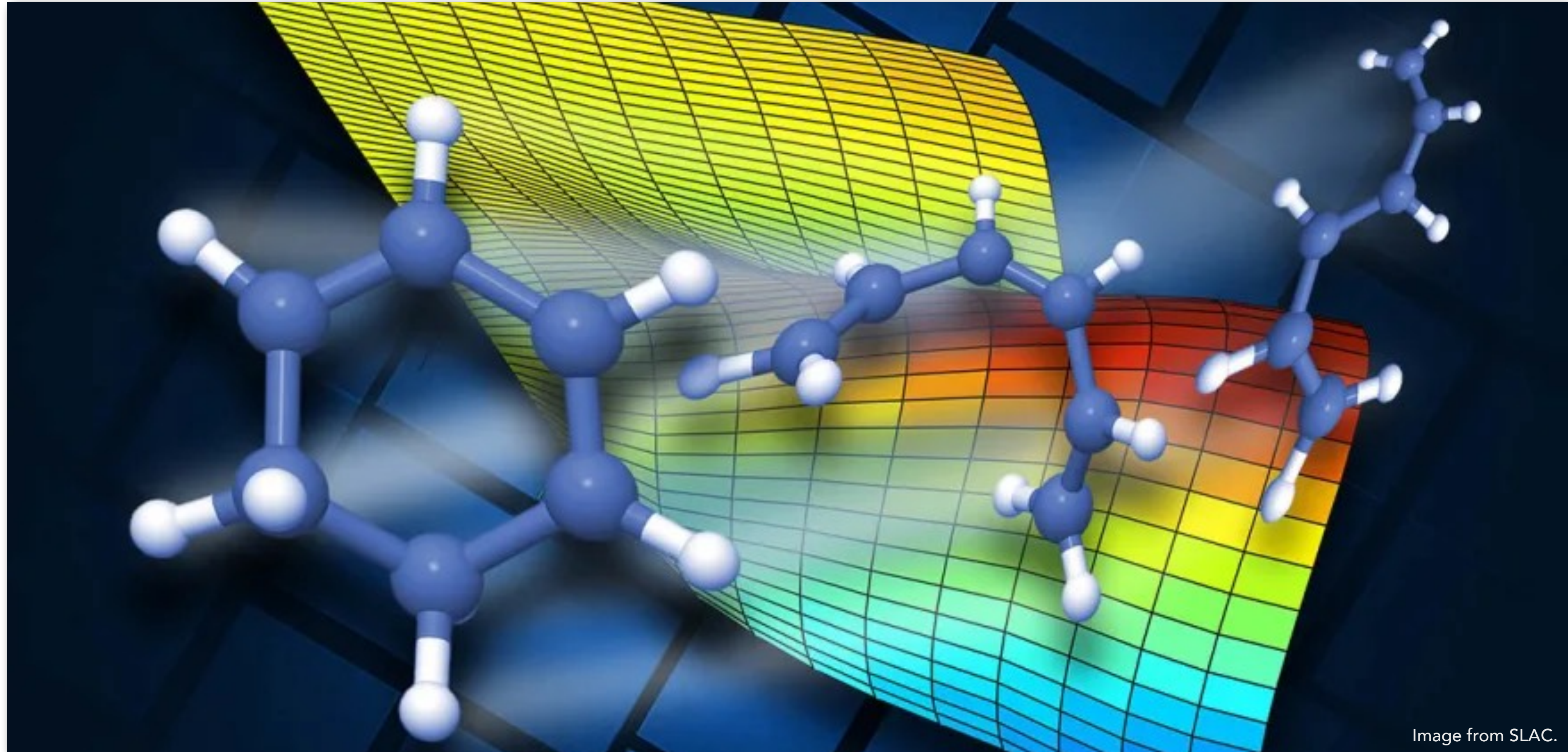
**Micheline B. Soley**, B. Allen, V. S. Batista, in preparation.

**Micheline B. Soley**, D. D. Yavuz, in preparation.

J. Dai,\* E. Palmer,\* J. M. Hawthorne,\* A. Vidwans, **Micheline B. Soley**, in preparation.

# OUTLOOK

Tensor networks and quantum computers' efficiency and ability to enable exact quantum dynamics makes the method well-suited to a wide range of processes in chemistry beyond reach with standard grid-based methods.



T. H. Kyaw\*, **Micheline B. Soley**\*, B. Allen, P. Bergold, C. Sun, V. S. Batista, A. Aspuru-Guzik, (2022) arXiv:2208.10470v1.

**Micheline B. Soley**, P. Bergold, V. S. Batista, JCTC 17 (2021) 3280.

# ACKNOWLEDGEMENTS



University of Wisconsin-Madison Office of the  
Vice Chancellor for Research and Graduate  
Education

## BLUE WATERS

Blue Waters Graduate Research Fellowship,  
Supported by NSF (OCI-0725070, ACI-1238993)  
and the State of Illinois, Joint Effort of UIUC  
and NCSA



## WARF

Wisconsin Alumni Research Foundation

Wisconsin Alumni Research Fund



NSF GRFP (DGE-1144152),  
NSF Grant No. CHE-1900160,  
NSF CCI Center for  
Quantum Dynamics on Modular  
Quantum Devices (2124511)



NERSC

## YQI

YQI Postdoctoral Fellowship

## Y | CRC

Yale High Performance  
Computing Center



Harvard GSAS  
Merit/Graduate Society  
Fellowship