

# Maxeler University program: The idea

## What can we do for the academic groups?

- ✓ Provide dataflow HPC hardware and software free or at strongly reduced cost
- ✓ Share our expertise in dataflow implementation of computational algorithms
- ✓ Help in securing research funding for projects involving dataflow computing

## How can we profit from the interaction with academic groups?

- ✓ We would like to see major, non-incremental scientific results being achieved with dataflow technology – we consider this as the best possible advertising
- ✓ We would like to expand the use of dataflow systems to entirely new areas where their commercial potential is not at all clear
- ✓ We would like to create an expertise base in dataflow system programming, especially in the key scientific centres in the world

*“The exciting thing about this program is that it covers a wide spectrum of application areas in high performance computing and brings together centres of excellence in HPC using FPGA in USA, Europe, Japan and China into a powerful community”*

(Ram Subramanian, Worldwide Manager of the Xilinx University Program)

# Maxeler University Program Members



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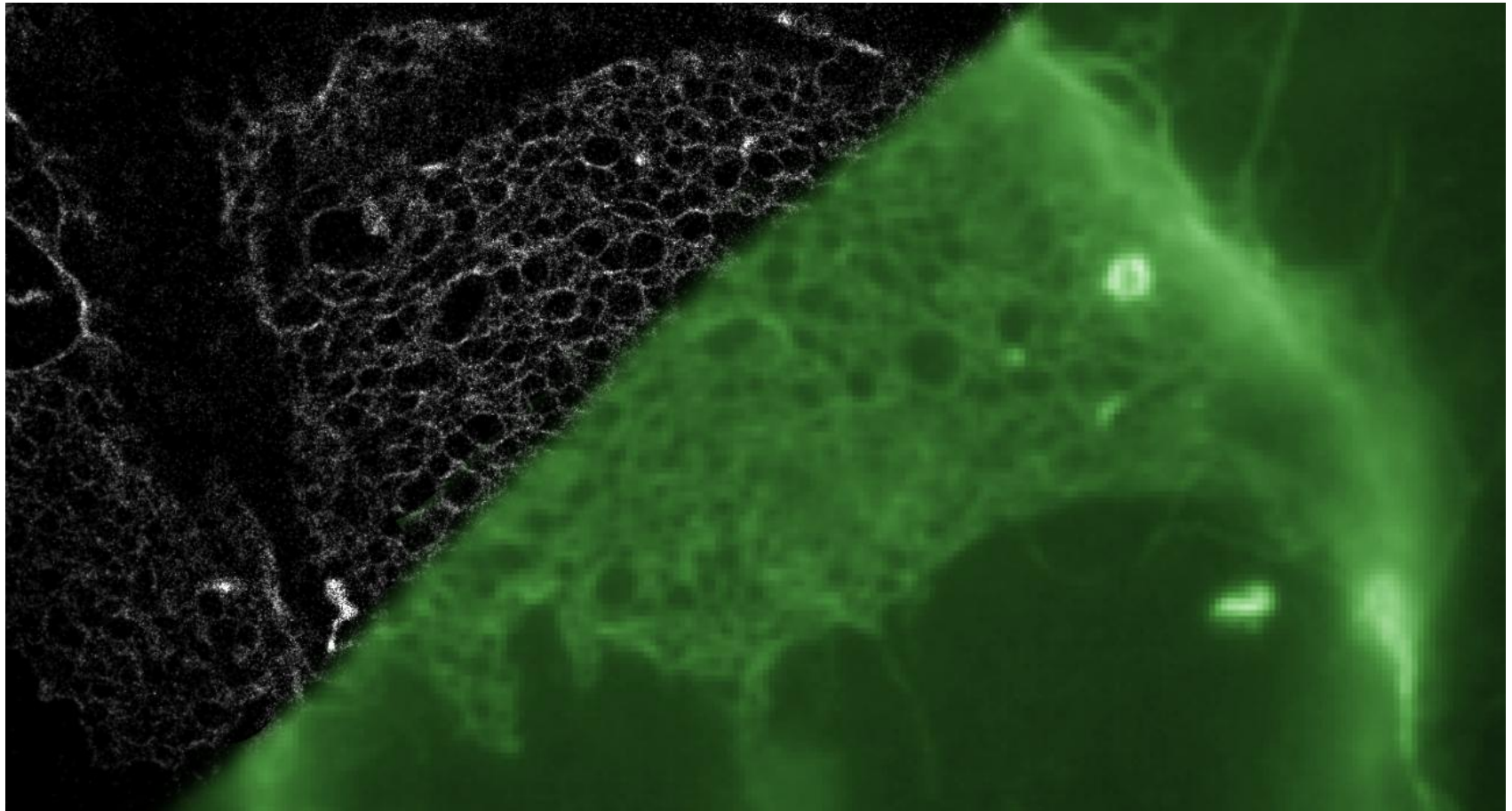


# Maxeler University Program Members

- Tsinghua University
- University of Tokyo
- Technion-Israel Institute of Technology
- University of Oslo
- University of Hamburg
- Nanyang Technological University
- National University of Singapore
- Korea University
- Japan Advanced Institute of Science and Technology
- University of Kaiserslautern
- Universität Erlangen-Nürnberg
- Manchester Metropolitan University
- Politecnico di Milano
- Hunan University
- Stanford University
- Universität Paderborn
- Universität Hamburg
- Goethe-Universität Frankfurt am Main
- Tohoku University
- University of Lisbon
- Chalmers University of Technology
- Northeastern University
- University of Edinburgh
- Heriot-Watt University
- Delft University of Technology
- University of Oxford
- Hong Kong University
- University of Illinois at Chicago
- The Chinese University of Hong Kong
- Max-Planck-Institut für molekulare Genetik
- Karlsruhe Institute of Technology
- Nagasaki University
- University of California Berkeley
- University of British Columbia
- Imperial College London
- University of Brno
- University of Novi Sad
- Kragujevac University
- AGH University of Science and Technology
- University of Windsor, Ontario
- Manchester Business School
- Fudan University
- Hong Kong Polytechnic University
- Univ. Sts Cyril and Methodius
- TU Crete
- UCL
- UCLA
- UBC-EOS
- Chosun University
- Ludwig-Maximilians Universität München

# MAX-UP: The scope of subjects

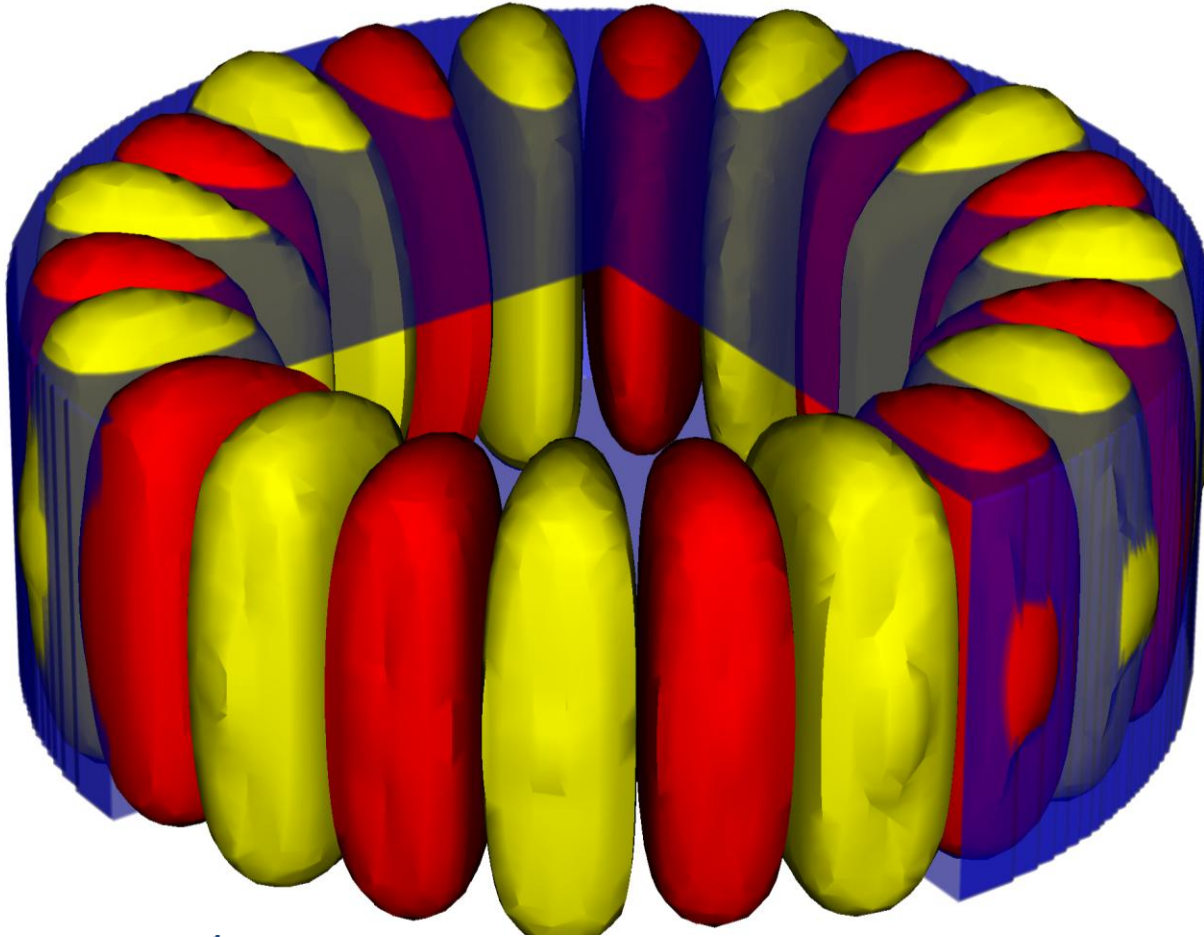
- **Financial modelling**
- **Quantum chemistry**
- **Bio-chemical engineering**
- **Geophysics**
- **Atmospheric physics**
- **Strong-field laser physics**
- **Virtual reality and computer gaming**



Localization microscopy enhances the resolution of fluorescence light microscopy (shown in green) by about an order of magnitude. Single fluorescent molecules act as switchable markers. Their detected signals can be fitted with a two-dimensional Gaussian distribution and thus located with sub-pixel resolution. Using MaxCompiler we achieved a hardware acceleration of 225 in signal detection and fitting.



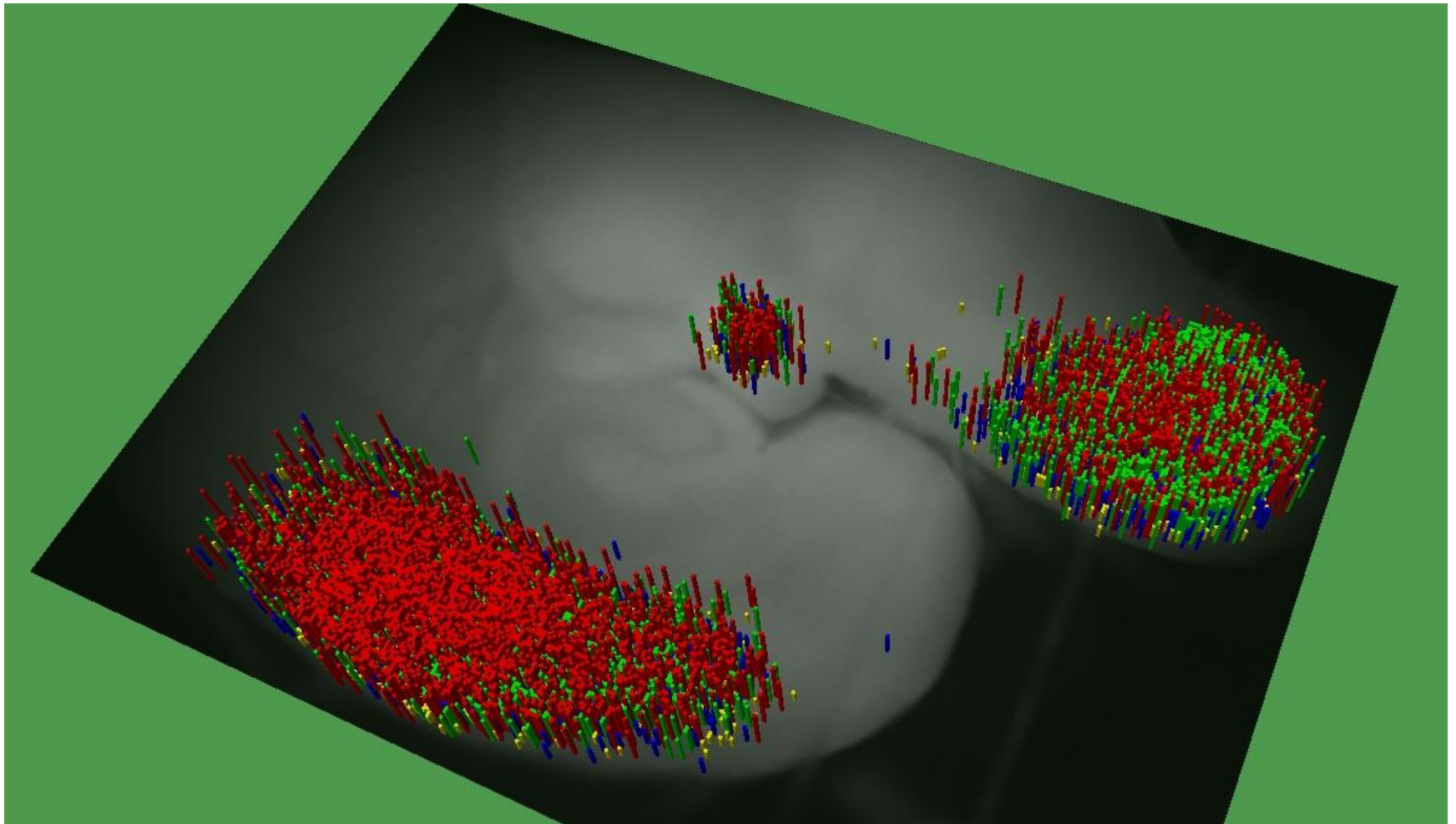
KIRCHHOFF-  
INSTITUTE  
FOR PHYSICS



Acceleration of  
simulations of  
computational  
nanophotonics using  
Maxeler MaxGenFD  
Finite Difference  
Compiler for dataflow



**UNIVERSITÄT PADERBORN<sup>0</sup>**  
*Die Universität der Informationsgesellschaft*

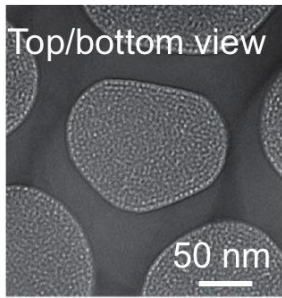


**UIC** University of Illinois  
at Chicago

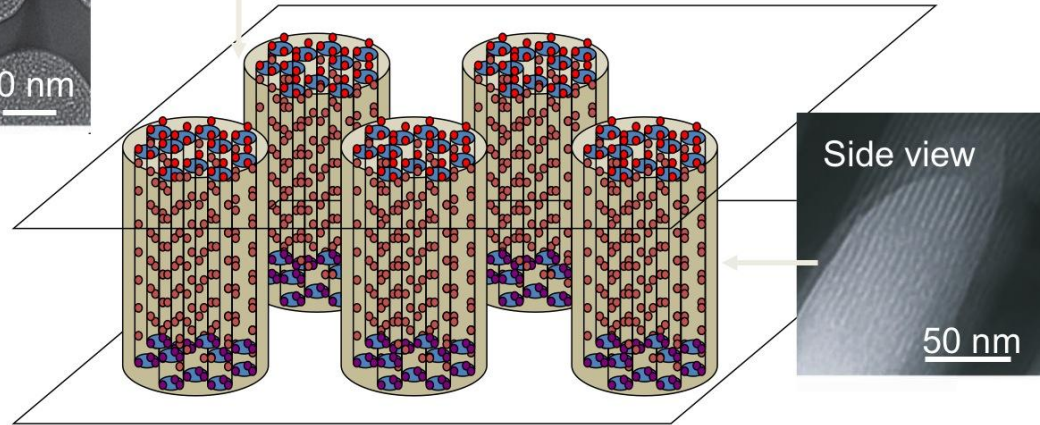
### Brain network analysis

Dynamic community analysis on corpus callosum to identify functional regions. By accelerating the linear correlation calculation, Maxeler dataflow engines can build brain networks on-the-fly while the lab experiment is running.

# Bio-inspired membrane for water desalination and purification

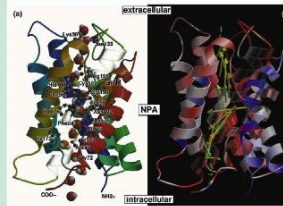
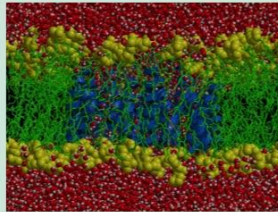


Nanoporous silica with controlled surface structure and charges ( ● ● ● )



Mechanism revealed via multi-scale and molecular simulations

Aquaporins  
in cell  
membrane



Simulation-Aided Design and  
Synthesis of Bio inspired  
Membranes for Water Desalination  
and Purification





**CAVE  
automatic  
virtual  
environment**

This project will investigate the use of dataflow architectures in creation of ***high-end immersive virtual reality***. The goal is to build a proof of concept simulator with ***extremely low latency***.



# A MAX-UP project example

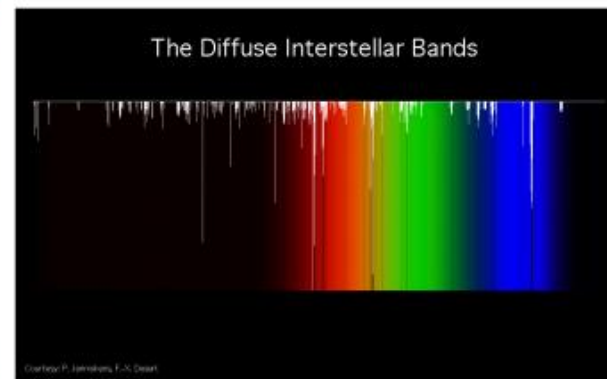
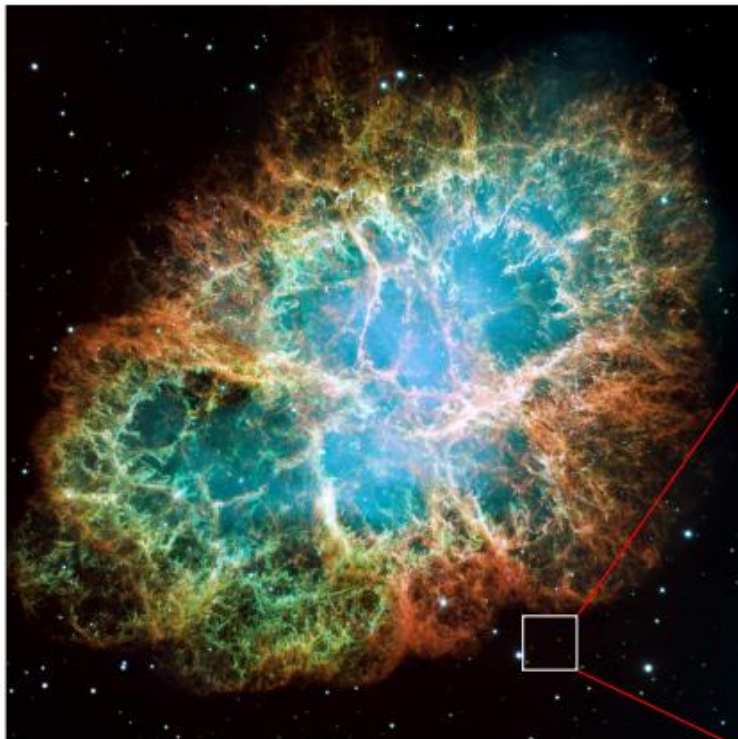
W Vigor, A Thom\*, M Bearpark, Chemistry, Imperial College; R Bruce, Maxeler  
\*now at Cambridge

Imperial College  
London

Astrochemistry

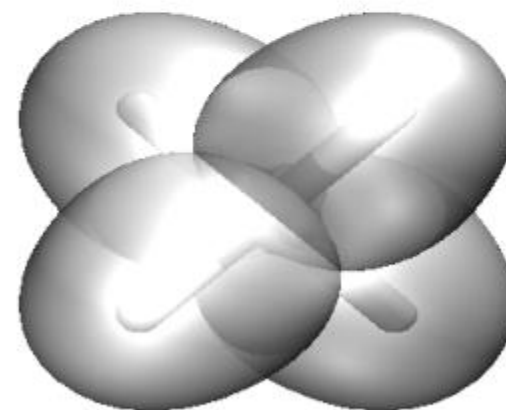
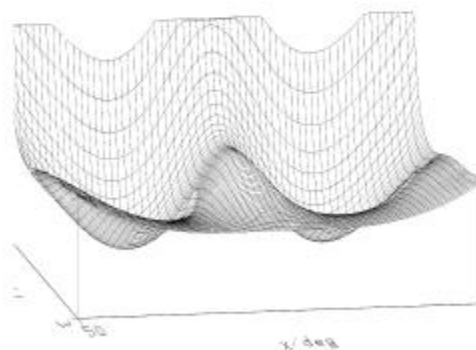
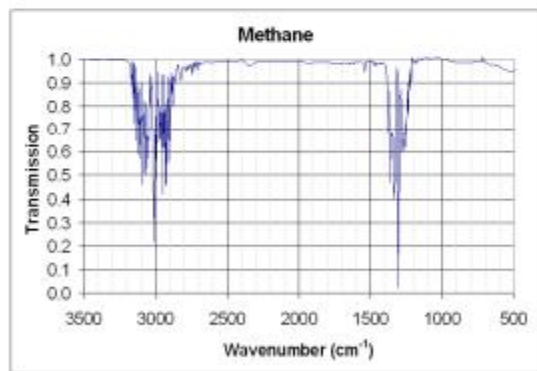
MAXELER  
Technologies  
MAXIMUM PERFORMANCE COMPUTING

coronene:  $C_{24}H_{12}$



<http://www.nasa.gov/>

P. Jenniskens and F.-X. Desert, *Astronomy and Astrophysics Supp. Ser.* **106**, 39–78 (1994)



Vibrational spectrum

⇐ Energy surface

⇐ Electronic structure

R. Marquardt and M. Quack, *J. Phys. Chem. A* **108**, 3166–3181 (2004)



$N$

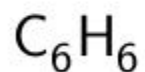
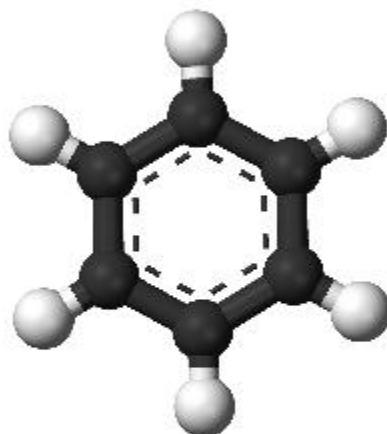
10

$M$

172

$N_d$

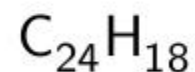
$10^9$



42

510

$10^{62}$



156

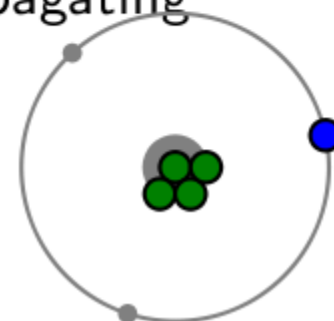
2500

$10^{253}$

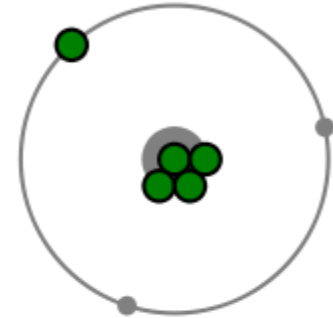
- ▶ Solutions to  $\mathbf{H}\mathbf{c} = E\mathbf{c}$  also solve  $e^{-\tau\mathbf{H}}\mathbf{c} = e^{-\tau E}\mathbf{c}$ .
- ▶ Propagate  $\frac{\partial\mathbf{c}}{\partial\tau} = -\mathbf{H}\mathbf{c}$ .
- ▶ Lowest energy eigenvector  $\mathbf{c}$  becomes dominant.

$$\frac{\partial c_i}{\partial\tau} = -\sum_j H_{ij}c_j.$$

- ▶ Represent  $c_i$  as populations of signed **psips** propagating through space.



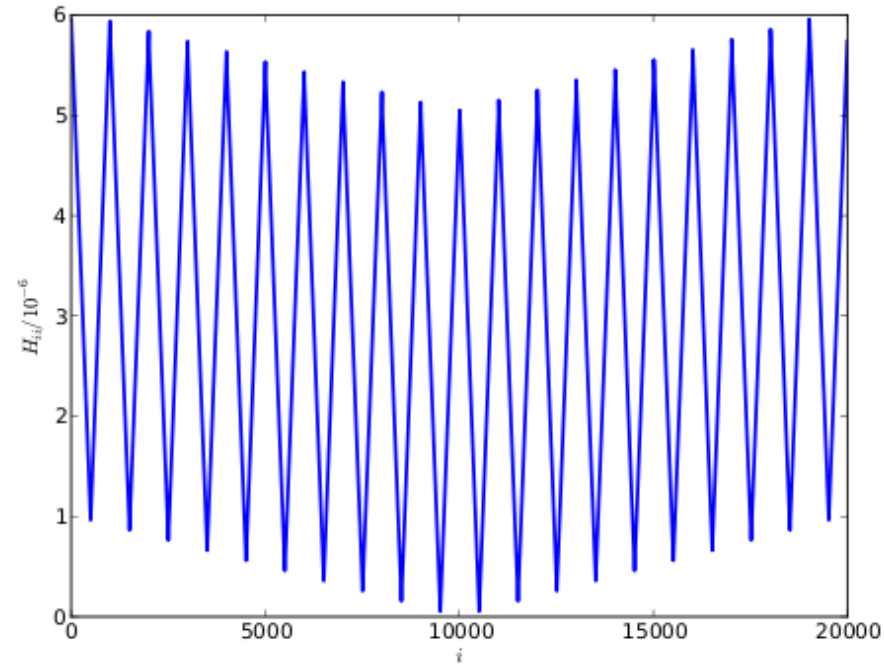
$$\frac{\partial c_i}{\partial \tau} = -[H_{ii} - S]c_i - \sum_{j \rightarrow i} H_{ij}c_j.$$



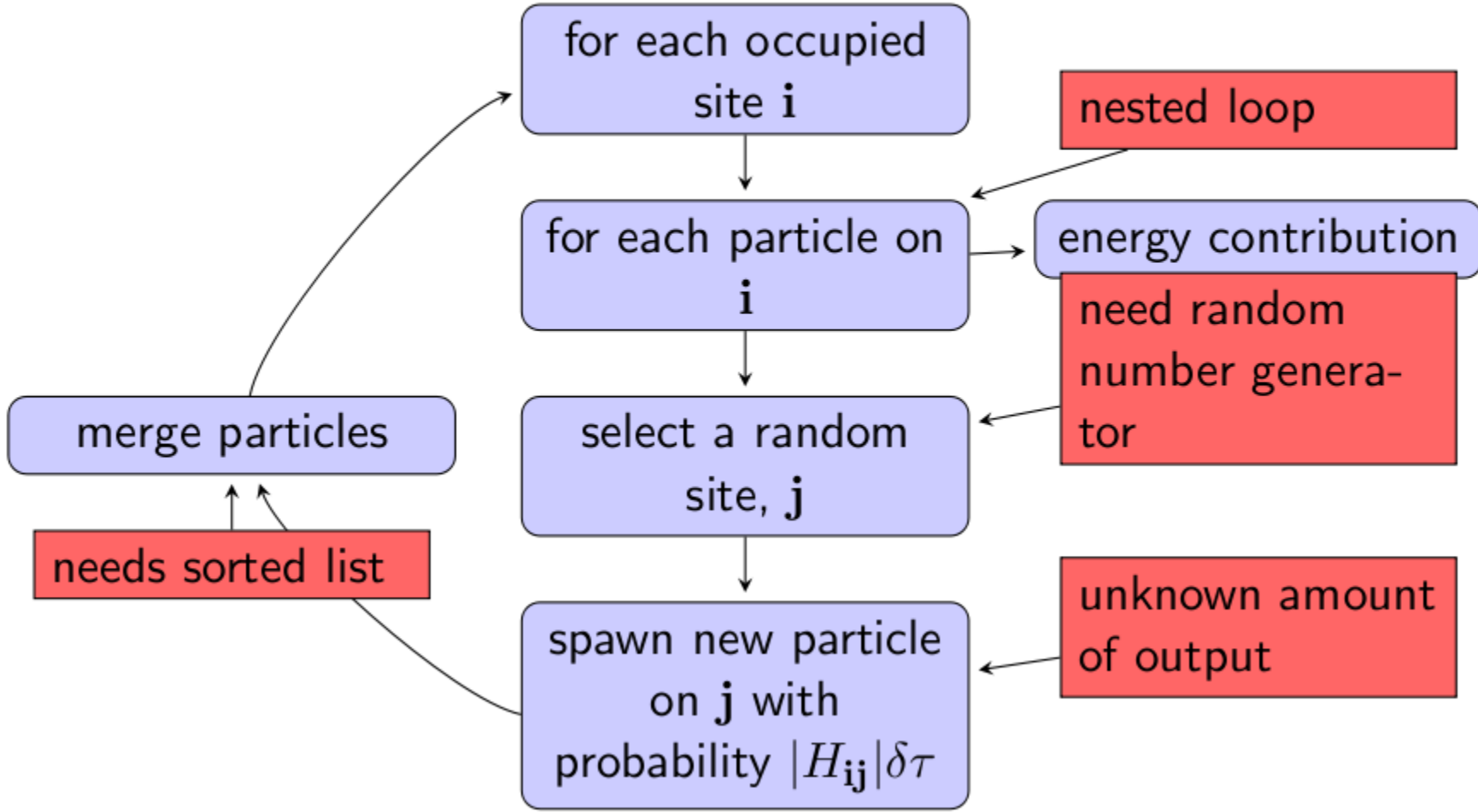
- ▶ Given a psip at  $\mathbf{j}$ , at time  $\tau + \delta\tau$  **spawn** a new one at randomly chosen  $\mathbf{i}$  based on  $-\delta\tau H_{ij}$ . This samples  $\sum_j$ .
- ▶ Psips at  $\mathbf{i}$  **die** based on  $\delta\tau(H_{ii} - S)$ .
- ▶ Start  $S = \min H_{ii}$ . Once enough psips created change  $S$  in response to population. e.g. for growth, make  $S$  more negative, so more death.
- ▶ Opposite-signed psips at the same determinant **annihilate**.

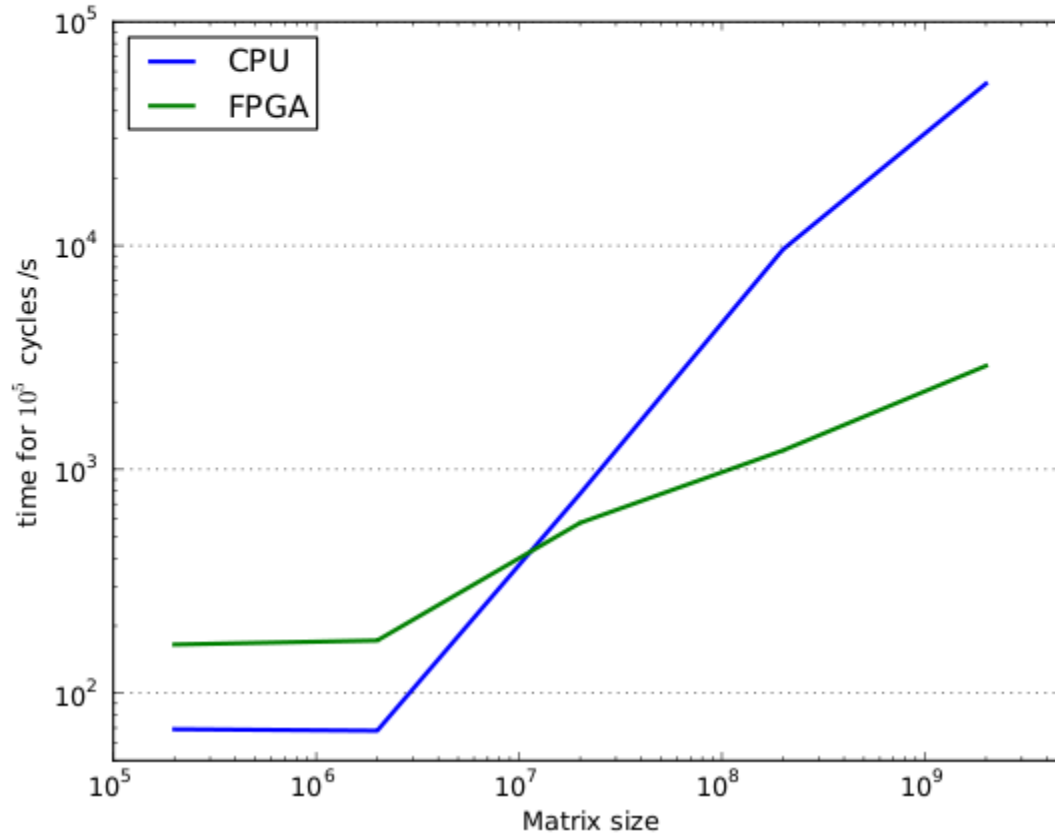
- ▶ Real elements of  $\mathbf{H}$  are very complicated.
- ▶ Sparse Toy matrix used for testing.

$$\mathbf{H} = \begin{pmatrix} H_{11} & a & 0 & 0 & \dots & 0 \\ a & H_{22} & a & 0 & \dots & 0 \\ 0 & a & H_{33} & a & & 0 \\ \vdots & & \ddots & \ddots & \ddots & 0 \\ 0 & \dots & 0 & a & H_{p-1p-1} & a \\ 0 & 0 & \dots & 0 & a & H_{pp} \end{pmatrix}$$



# Problems





## What next?

- 2D Hubbard model
- Real life FCI for molecules, such as polycyclic aromatic hydrocarbons (PAHs)

# MAX-UP: The future

- Strengthening existing subjects: bio-informatics, quantum chemistry, ... with new collaborations
- Bringing new subjects: chemical engineering, bio-engineering, ...
- Strengthening the existing UK, German, US bases of MAX-UP
- Expansion to Eastern Europe, South America, Australia, ...
- Research funding applications with MAX-UP members: iCASE PhD studentships (UK), ERC synergy grants, ...

Acceleration of an algorithm based on the Gross Pitaevskii equation.

Execution time for two iterations:

CPU execution time: 50s

Dataflow execution time: 4.9s

Achieved speedup: 10.2x

Programmer:

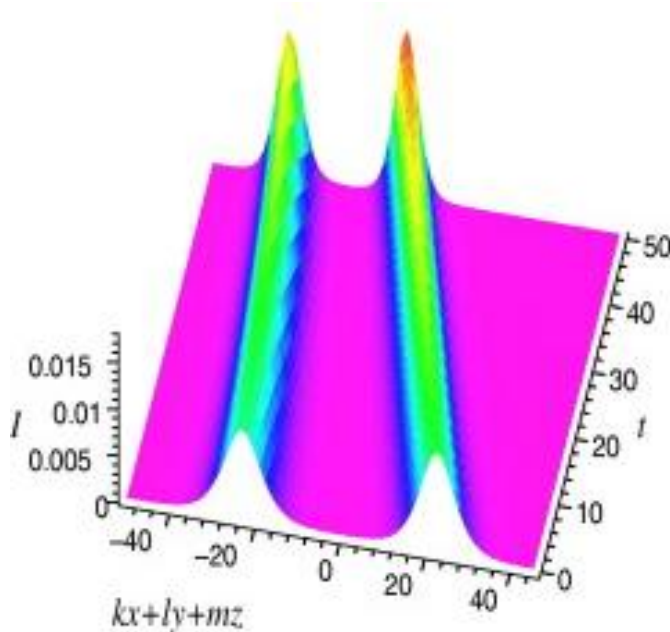
Sasha Stojanovic

[stojsasa@etf.bg.ac.rs](mailto:stojsasa@etf.bg.ac.rs)

Advisor:

Veljko Milutinovic

[vm@etf.bg.ac.rs](mailto:vm@etf.bg.ac.rs)



University of  
Belgrade

Note: Comparison is made between a single core CPU without using any of extensions available in today's modern processors and a single dataflow implementation on the oldest and smallest MAX 2 Maxeler card. Using modern CPUs equipped with several cores, each one with extensions for parallel processing of data, and modern Maxeler card with higher capacity, bandwidth to memory and host, and working frequency, will result in decreased execution times in both cases, but it is expected that the speedup will not change significantly.