

# The new SuperMUC petascale system and applications

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# Thank you to HCMUT Team



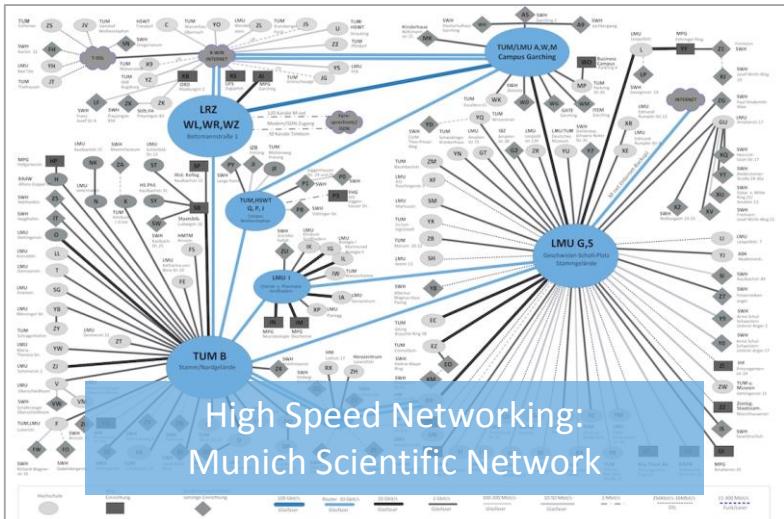
With approx. 250 employees  
for more than 100.000 students and  
for more than 30.000 employees  
including 8.500 scientists



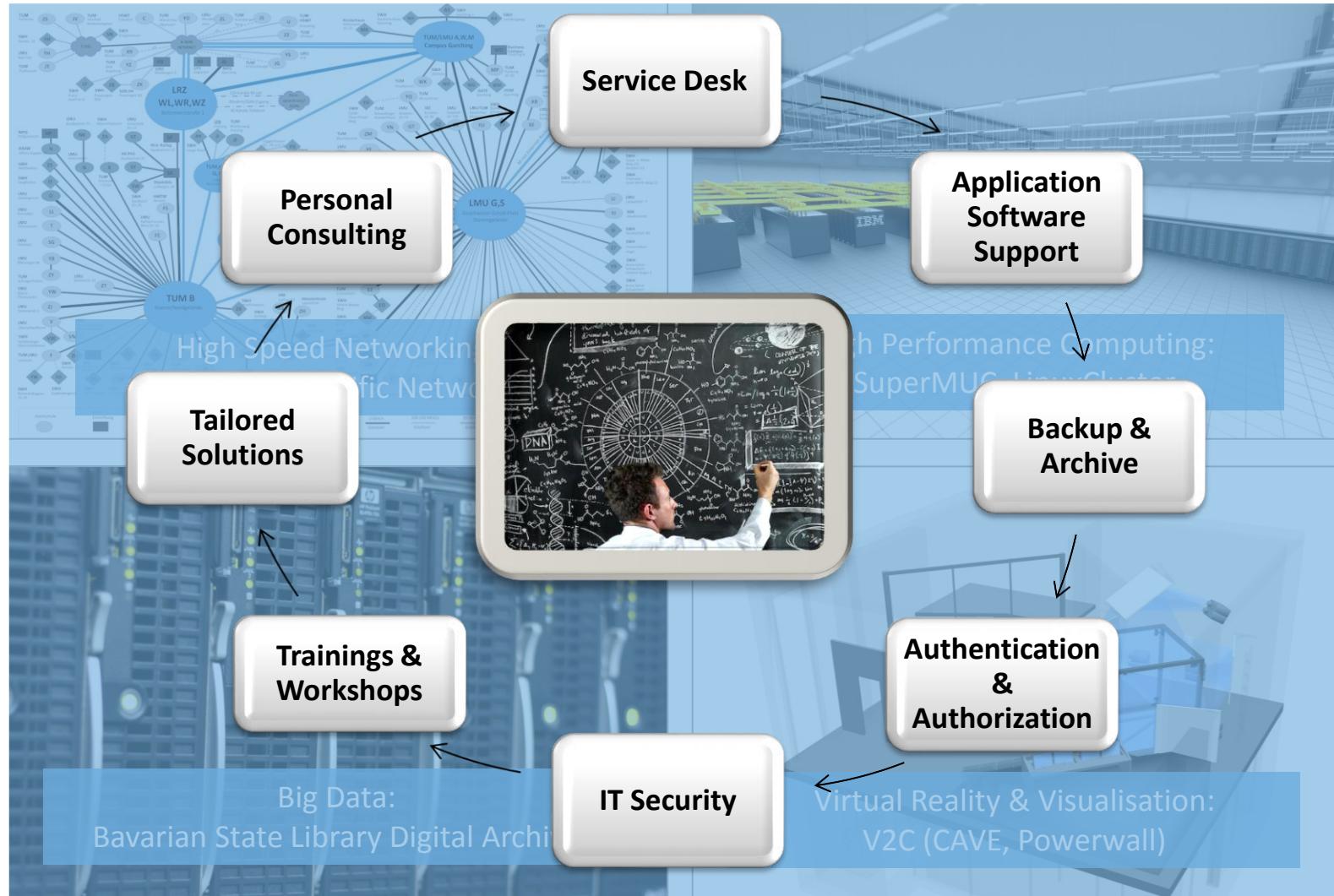
- European Supercomputing Centre
- National Supercomputing Centre
- Regional Computer Centre for all Bavarian Universities
- Computer Centre for all Munich Universities

Photo: Ernst Graf

# LRZ as IT Competence Centre: Operating Cutting-edge IT Infrastructure



# LRZ as IT Competence Centre: Providing Comprehensive IT Services for Science



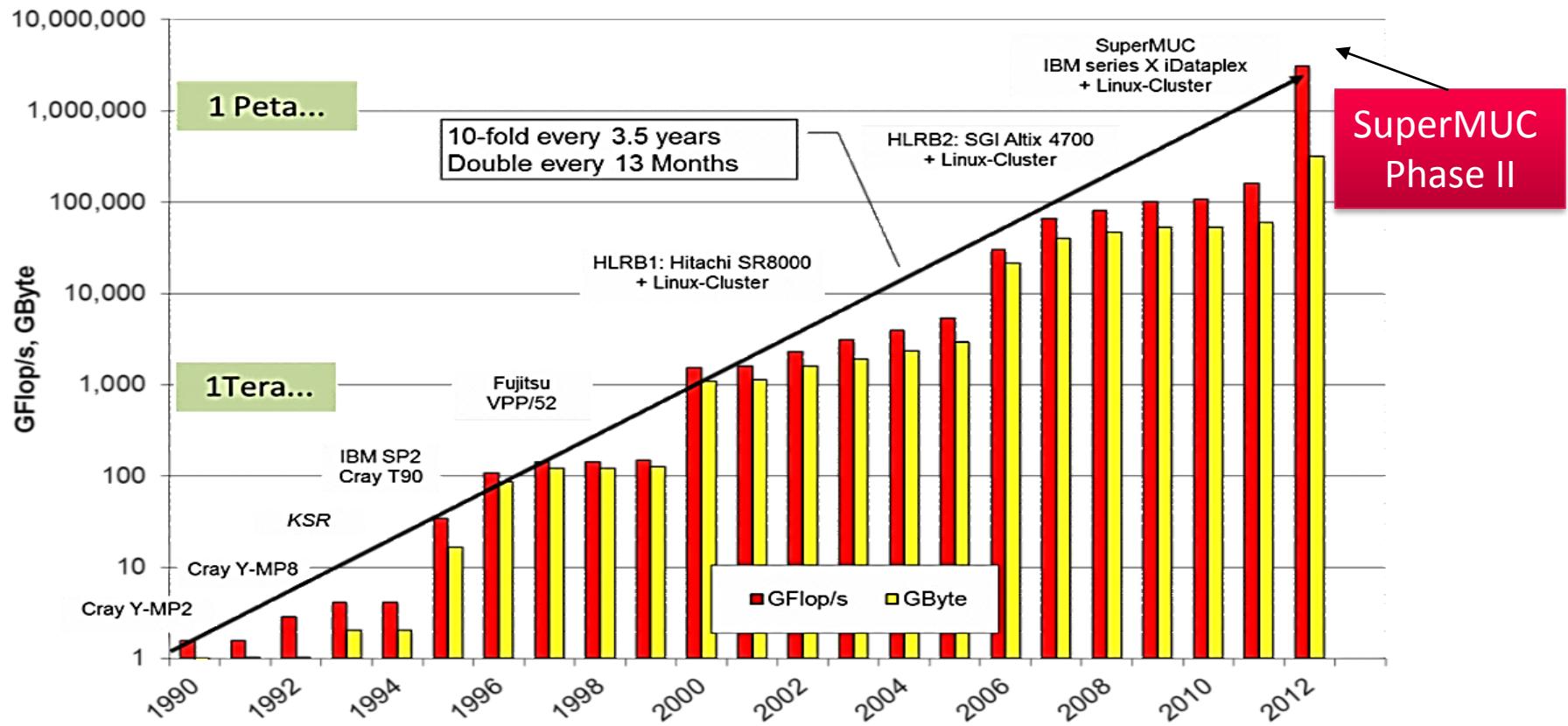


Video: **SuperMUC rendered on SuperMUC by LRZ**

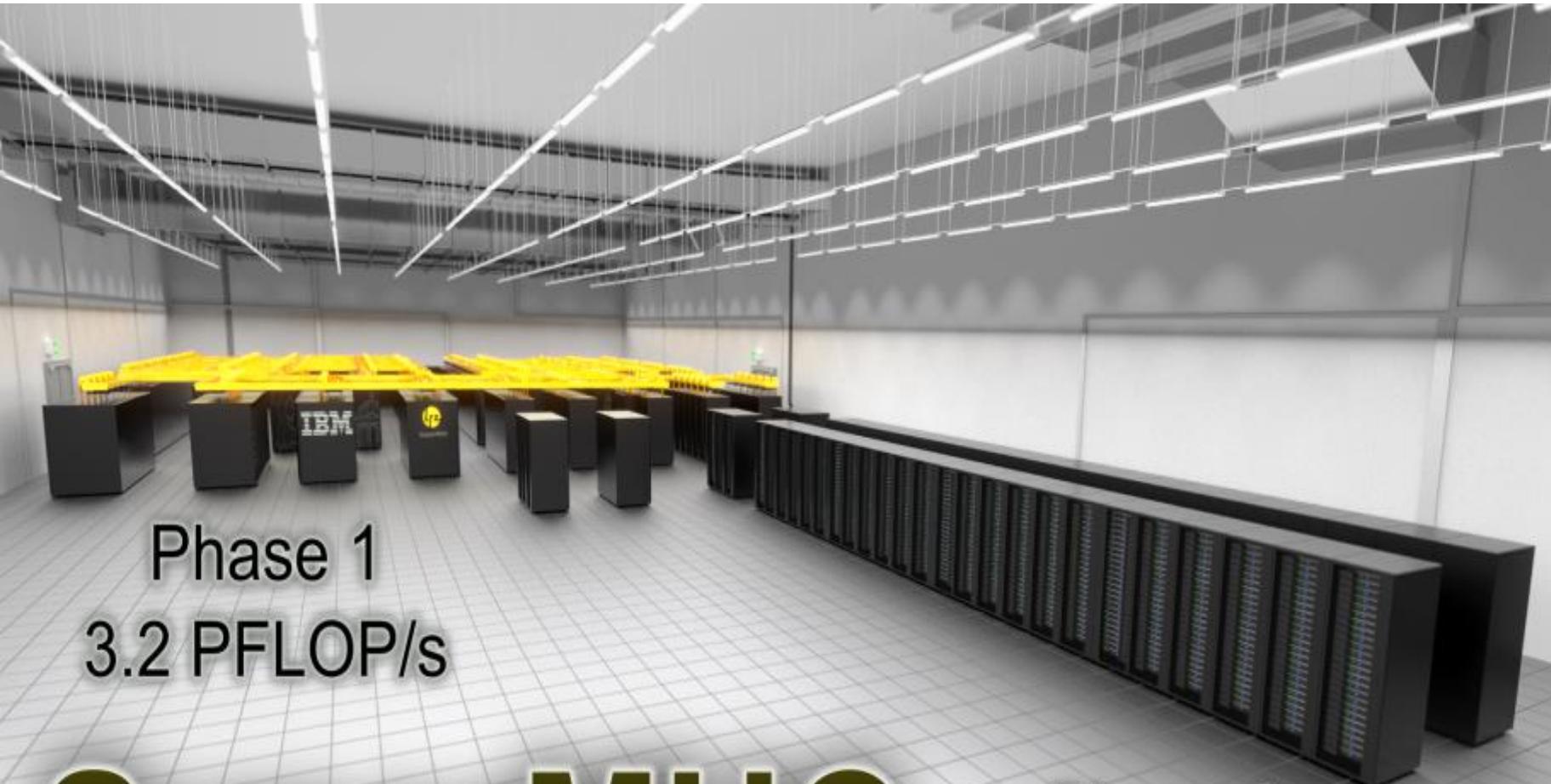
<http://youtu.be/OIAS6iiqWrQ>

# Top 500 Supercomputer List (June 2012)

Rank	Site	Computer/Year Vendor	Cores	R <sub>max</sub>	R <sub>peak</sub>	Power
1	DOE/NNSA/LLNL United States	Sequoia - BlueGene/Q, Power BQC 16C 1.60 GHz, Custom / 2011 IBM	1572864	16324.75	20132.66	7890.0
2	RIKEN Advanced Institute for Computational Science (AICS) Japan	K computer, SPARC64 VIIIfx 2.0GHz, Tofu interconnect / 2011 Fujitsu	705024	10510.00	11280.38	12659.9
3	DOE/SC/Argonne National Laboratory United States	Mira - BlueGene/Q, Power BQC 16C 1.60GHz, Custom / 2012 IBM	786432	8162.38	10066.33	3945.0
4	Leibniz Rechenzentrum Germany	SuperMUC - iDataPlex DX360M4, Xeon E5-2680 8C 2.70GHz, Infiniband FDR / 2012 IBM	147456	2897.00	3185.05	3422.7
5	National Supercomputing Center in Tianjin China	Tianhe-1A - NUDT YH MPP, Xeon X5670 6C 2.93 GHz, NVIDIA 2050 / 2010 NUDT	186368	2566.00	4701.00	4040.0
6	DOE/SC/Oak Ridge National Laboratory United States	Jaguar - Cray XK6, Opteron 6274 16C 2.200GHz, Cray Gemini interconnect, NVIDIA 2090 / 2009 Cray Inc.	298592	1941.00	2627.61	5142.0
7	CINECA Italy	Fermi - BlueGene/Q, Power BQC 16C 1.60GHz, Custom / 2012 IBM	163840	1725.49	2097.15	821.9
8	Forschungszentrum Juelich (FZJ) Germany	JuQUEEN - BlueGene/Q, Power BQC 16C 1.60GHz, Custom / 2012 IBM	131072	1380.39	1677.72	657.5
9	CEA/TGCC-GENCI France	Curie thin nodes - Bullx B510, Xeon E5-2680 8C 2.700GHz, Infiniband QDR / 2012 Bull	77184	1359.00	1667.17	2251.0
10	National Supercomputing Centre in Shenzhen (NSCS) China	Nebulae - Dawning TC3600 Blade System, Xeon X5650 6C 2.66GHz, Infiniband QDR, NVIDIA 2050 / 2010 Dawning	120640	1271.00	2984.30	2580.0



# SuperMUC Phase 1 + 2



# SuperMUC

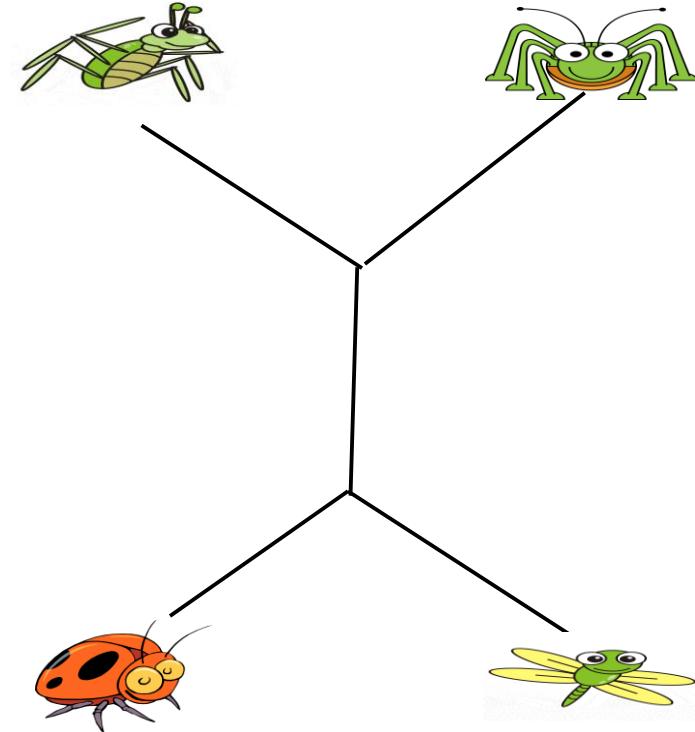
Phase 2  
3.2 PFLOP/s

- Computational Fluid Dynamics: Optimisation of turbines/wings, noise reduction
- Fusion: Plasma in a future fusion reactor (ITER)
- Astrophysics: Origin and evolution of stars and galaxies
- Solid State Physics: Superconductivity, surface properties
- Geophysics: Earth quake scenarios
- Material Science: Semiconductors
- Chemistry: Catalytic reactions
- Medicine and Medical Engineering: Blood flow, aneurysms, air conditioning
- Biophysics: Properties of viruses, genome analysis
- Climate research: Currents in oceans
- ...

# Phylogenetic Tree Computation

 ACGT  
 ACC  
 ACGG  
 AAGC

ACGT
ACC-
ACGG
AAGC



Sequencing



Alignment



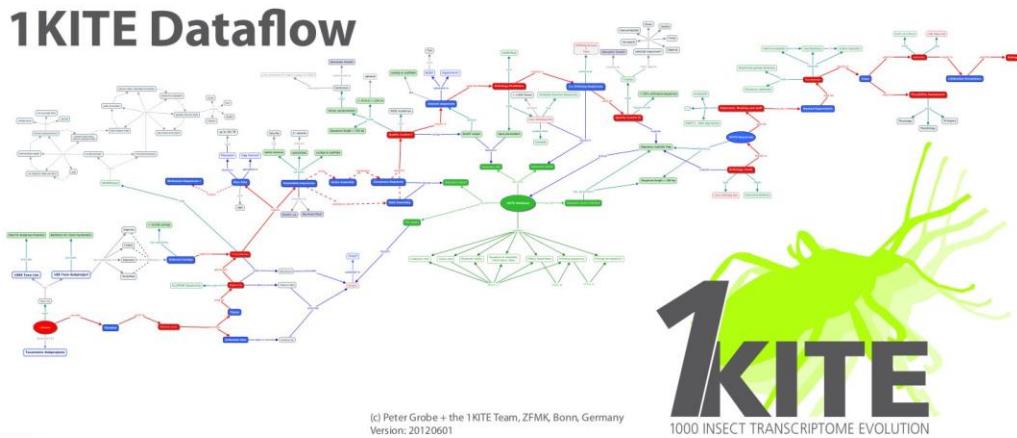
Phylogenetic Tree

Alexandros Stamatakis, H-ITS

- 4226517247809112252219618802377042809718932383449  
8822942857479880831434032178759024536798491951168  
3076494692867414802738570221298292428457687814873  
4552121861861600804474608426626044448936698500560  
2468116186441264227425440726676614927906540649360  
2976397461917469326750931190889241406694054603576  
66015625
- $\approx 4.22 \times 10^{301}$

- Alexandros Stamatakis  
Scientific Computing Group,  
Heidelberg Institute for Theoretical Studies (HITS) /  
Exelixis Lab
  
- „Big Data“ and High Performance Computing
- Novel software and applications needed
- Reading the data: only 1 minute (instead of 15 minutes)
- 1000 Processors: 17 hours (instead of 10 days)
- Load balancing

## 1KITE Dataflow



# Scientific Results - Publications



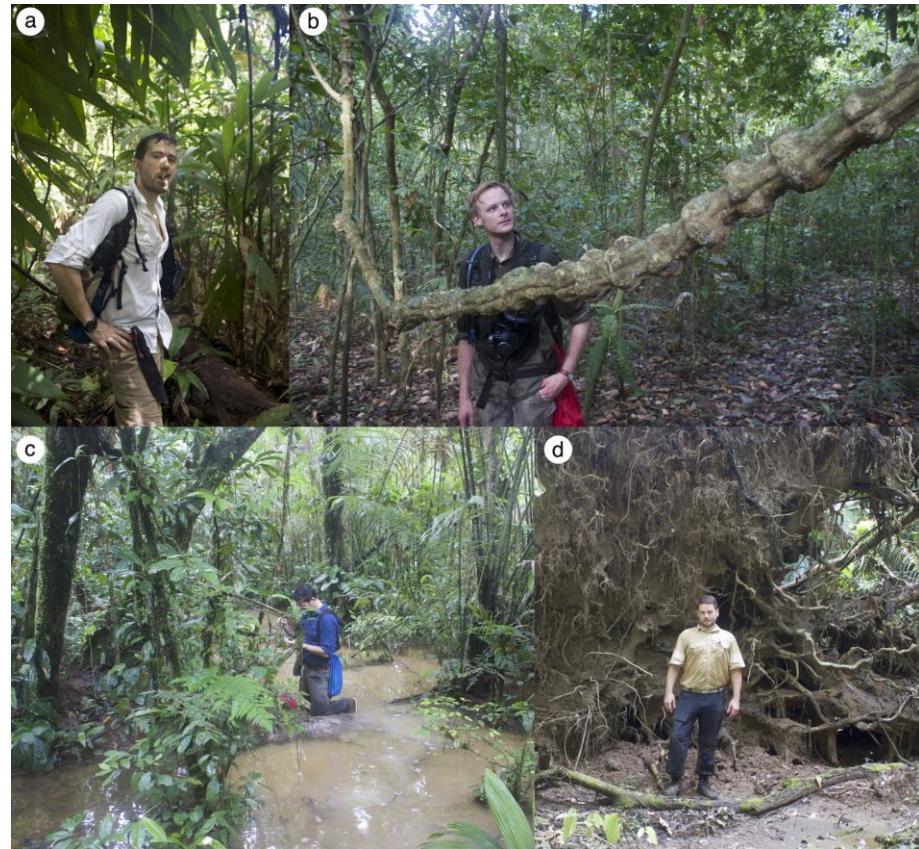
All aTwitter over an Internet study p. XXXX  
The extragalactic background's uneven glow pp. XXXX & XXXX  
A cellular target for human norovirus pp. XXXX & XXXX



Alexandros Stamatakis, H-ITS

- Neotropical Rainforests are hyperdiverse ecosystems
- Since Humboldt and Bonpland, we know about the high animal and plant richness
- New study now finds that unicellular eukaryotes are even more diverse
- Particularly the parasitic Apicomplexa dominate these forests
- Their presence might drive the diversity of macro-organisms

Micah Dunthorn/TU Kaiserslautern

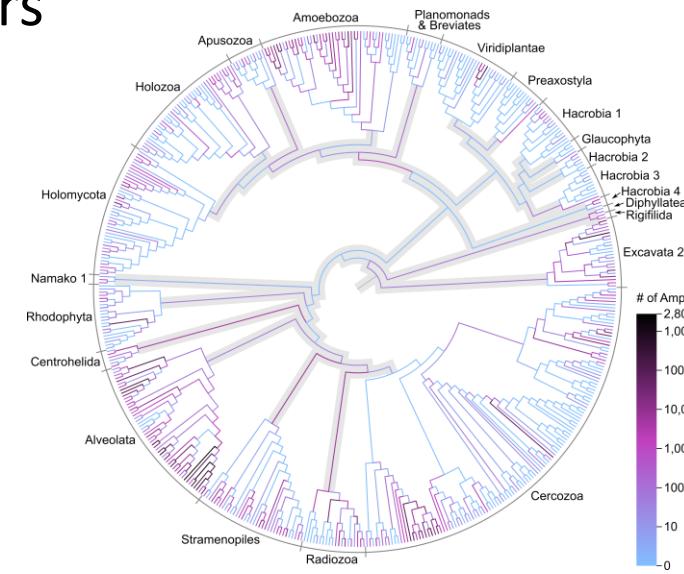


<https://natureecoevocommunity.nature.com/channels/521-behind-the-paper/posts/15402-a-larger-microbial-perspective-of-tropical-rainforests>

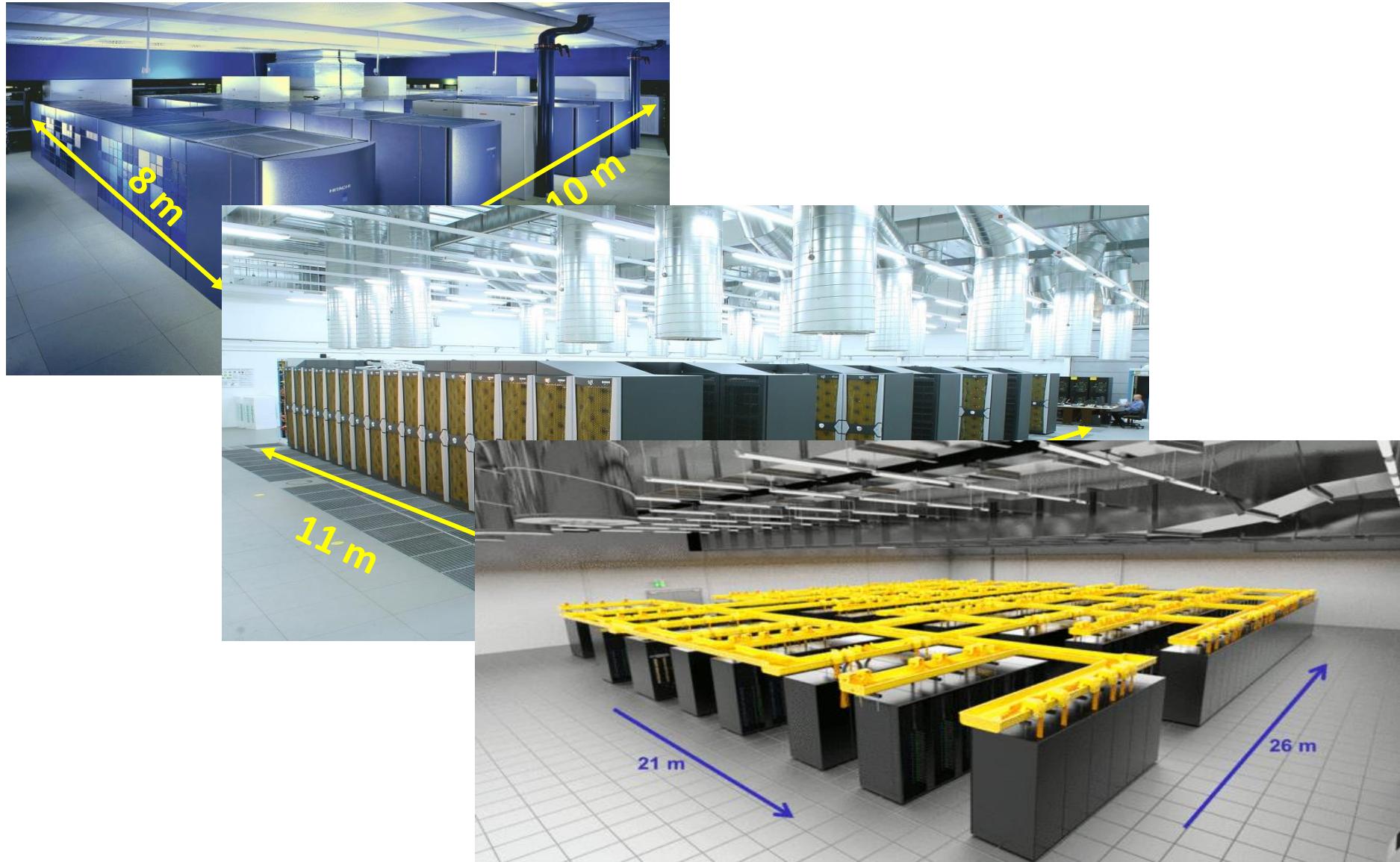
- More than 130 million DNA sequences were analysed
  - Most of them belong to yet unknown microbial species
  - Thus, a thorough method was necessary for classifying those sequences
  - The method takes the evolutionary history of known species into account
  - But this comes at the cost of increased computational needs
  - Approximately 1 million computation hours on **SuperMUC** were necessary



Mahé et al. (2017). Parasites dominate hyperdiverse soil protist communities in Neotropical rainforests.  
Nature Ecology and Evolution 1:09.  
DOI: 10.1038/s41559-017-0091



# SuperMUC and its predecessors



# LRZ Building Extension

Picture: Horst-Dieter Steinhöfer

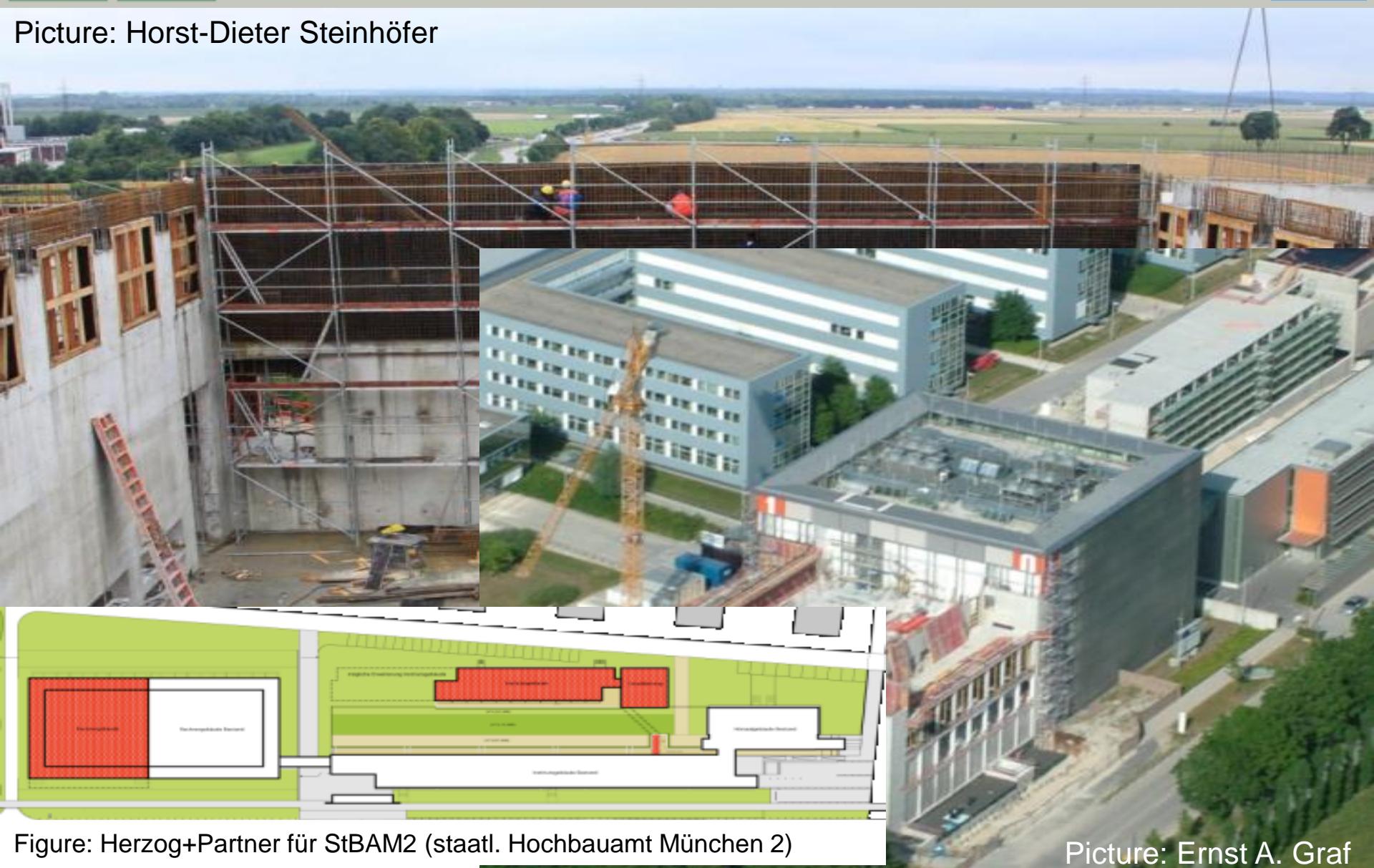
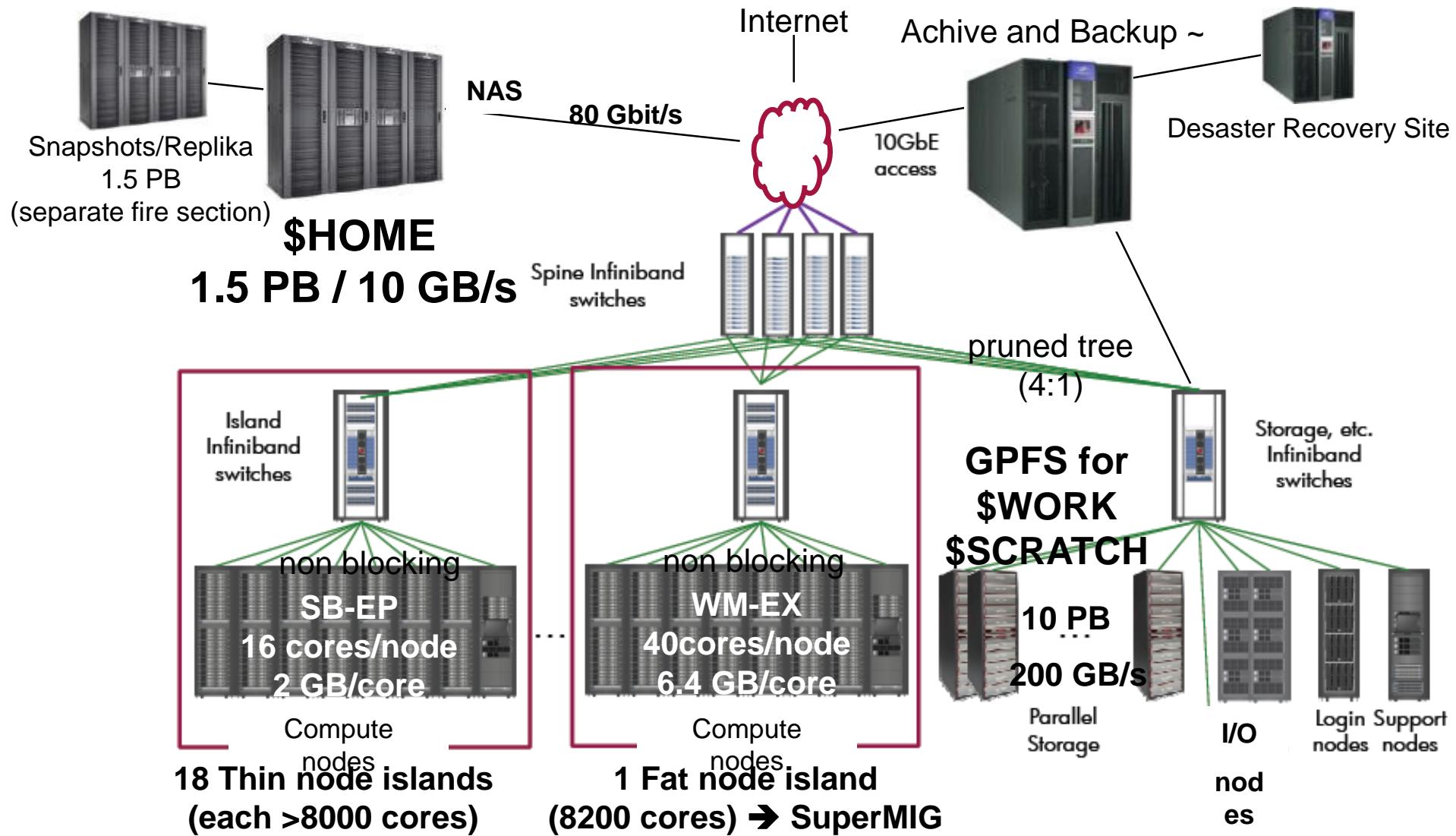


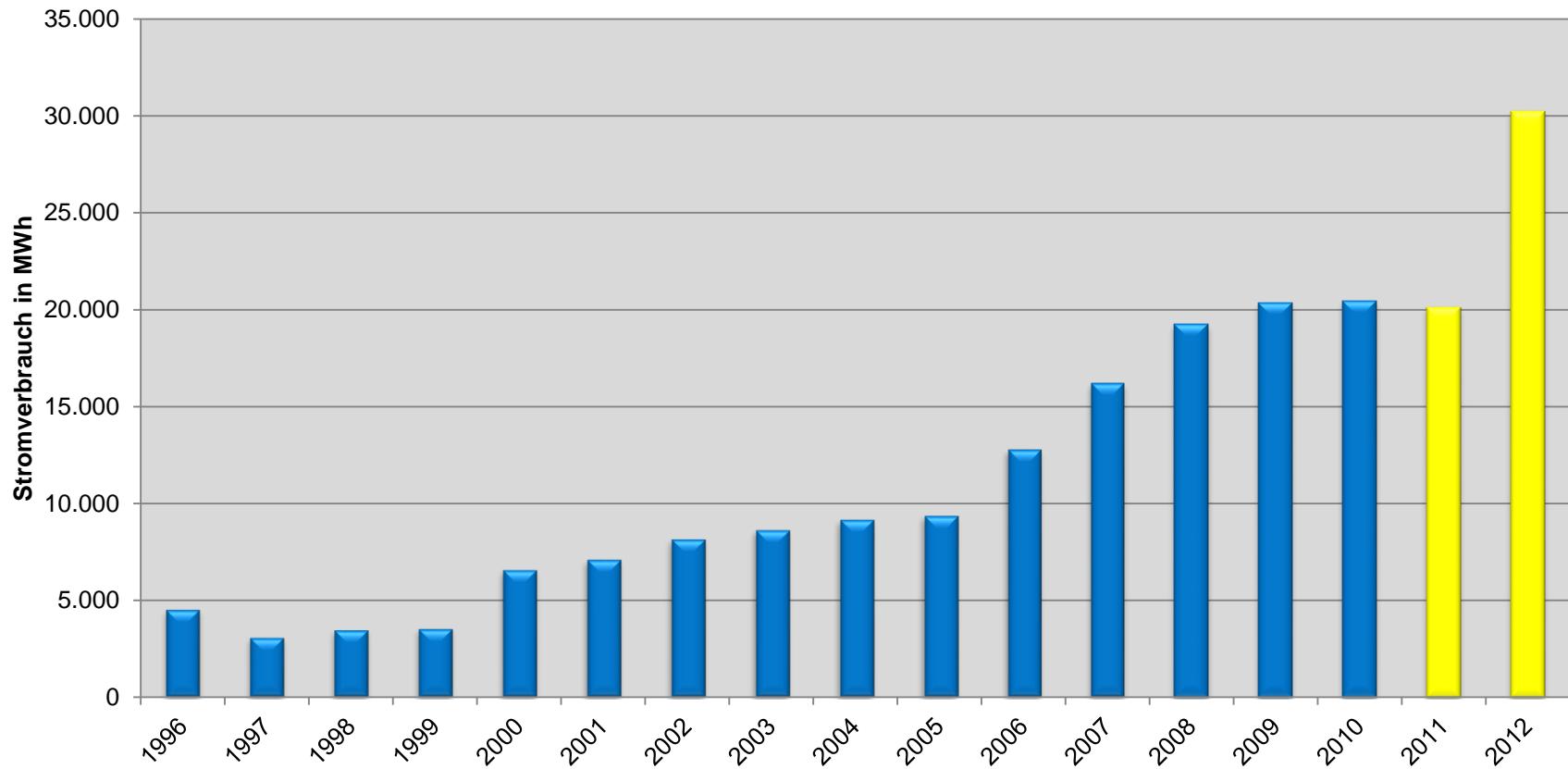
Figure: Herzog+Partner für StBAM2 (staatl. Hochbauamt München 2)

Picture: Ernst A. Graf

# SuperMUC Architecture



# Power Consumption at LRZ



# Cooling SuperMUC



# Energy Efficiency on SuperMUC @ LRZ

Photos: Torsten Blöth, Lenovo



- ✓ Usage of Intel Xeon E5 2697v3 processors
- ✓ Direct liquid cooling
  - 10% power advantage over air cooled system
  - 25% power advantage due to chiller-less cooling

- ✓ Energy-aware scheduling
  - 6% power advantage
  - ~40% power advantage
  - Annual savings: ~2 Mio. € for SuperMUC Phase 1 and 2

# Increasing Numbers of Cores

Date	System	Flop/s	Cores
2000	HLRB-I	2 Tflop/s	1512
2006	HLRB-II	62 Tflop/s	9728
2012	SuperMUC	3200 Tflop/s	155656
2015	SuperMUC Phase II	3.2 + 3.2 Pflop/s	229960

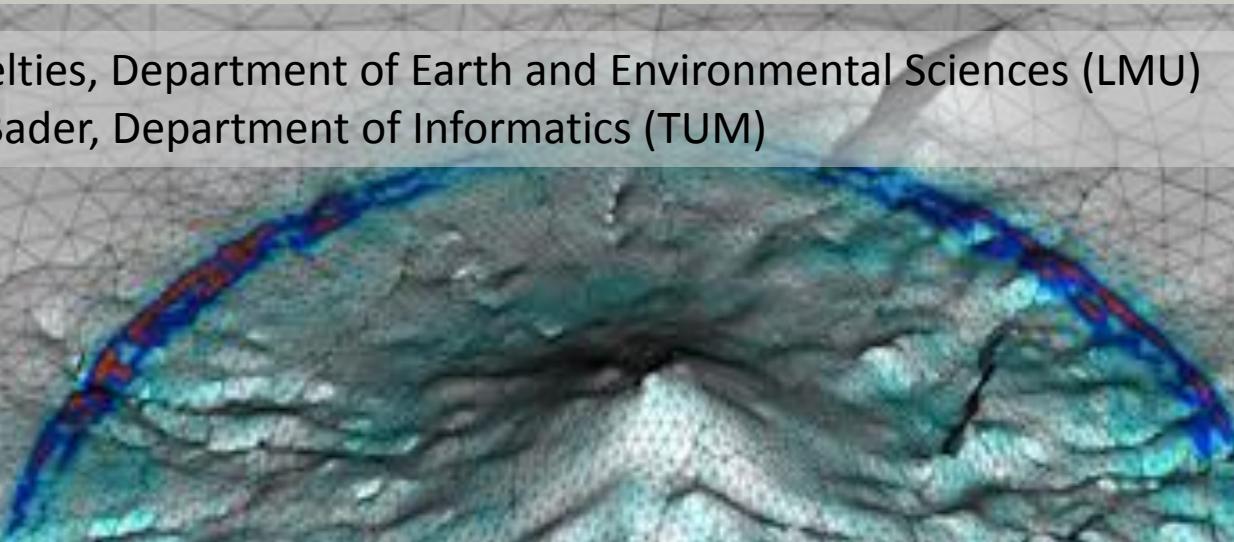
## ■ Results:

Name	MPI	# cores	Description	TFlop/s/island	TFlop/s max
Linpack	IBM	★ 128000	TOP500	161	2560
Vertex	IBM	★ 128000	Plasma Physics	15	245
GROMACS	IBM, Intel	☆ 64000	Molecular Modelling	40	110
Seissol	IBM	☆ 64000	Geophysics	31	95
walBerla	IBM	★ 128000	Lattice Boltzmann	5.6	90
LAMMPS	IBM	★ 128000	Molecular Modelling	5.6	90
APES	IBM	☆ 64000	CFD	6	47
BQCD	Intel	★ 128000	Quantum Physics	10	27

## ■ Sustained TFlop/s on 64000/128000 cores

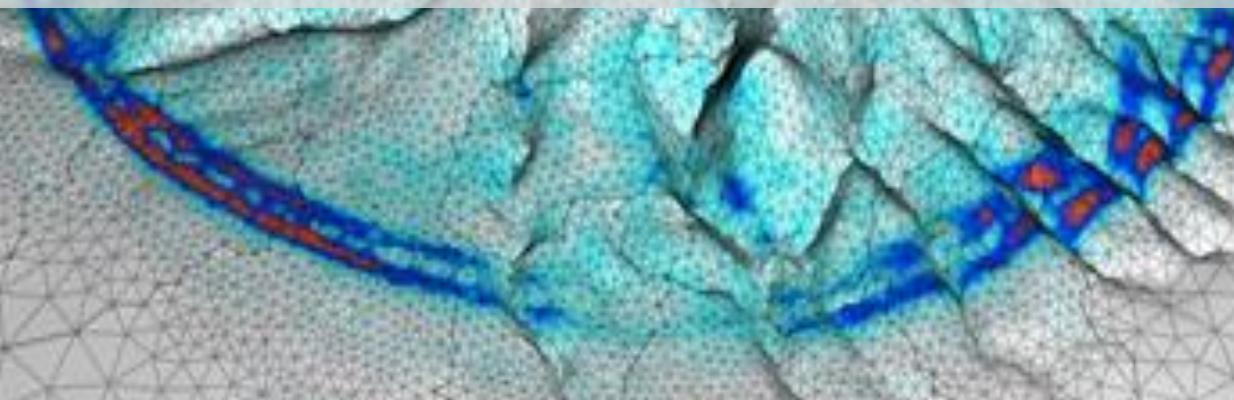
Dr. Christian Pelties, Department of Earth and Environmental Sciences (LMU)

Prof. Michael Bader, Department of Informatics (TUM)



1,42 Petaflop/s on 147.456 Cores of SuperMUC  
(44,5 % of Peak Performance)

[http://www.uni-muenchen.de/informationen\\_fuer/presse/presseinformationen/2014/pelties\\_seisol.html](http://www.uni-muenchen.de/informationen_fuer/presse/presseinformationen/2014/pelties_seisol.html)



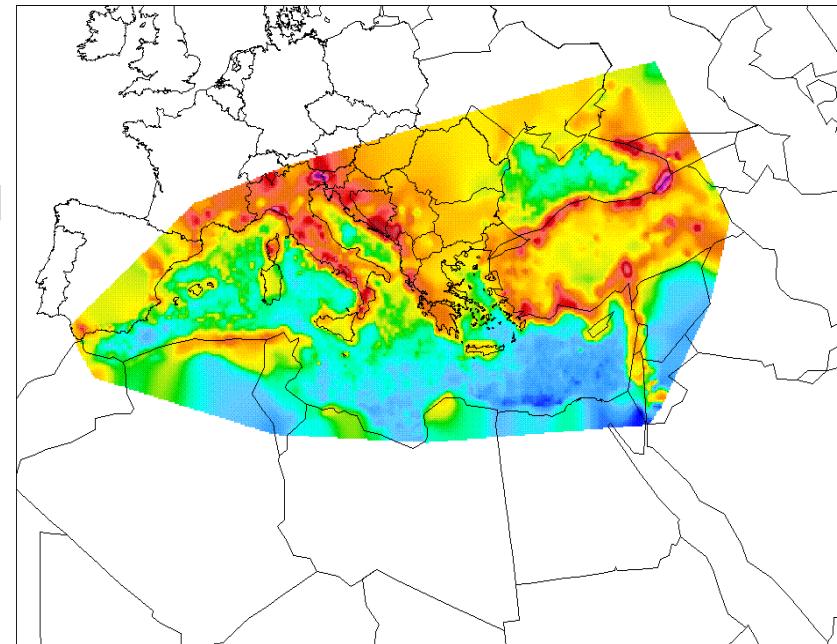
Picture: Alex Breuer (TUM) / Christian Pelties (LMU)

## ■ LRZ benefits

- Understanding the (current and future) needs and requirements of the respective scientific domain
- Developing future services for all user groups
- Thematic focusing: **Environmental Computing**

## ■ EU Project Series DRIHM\*

- Flash Project estimates for 1990-2006
- > 29 billion euros in damages produced by floods
- > 4,500 total number of casualties



SSMI and raingauge observations (1978-1994)

# Professor Peter V. Coveney



- holds a chair in Physical Chemistry
- is an Honorary Professor in Computer Science at University College London (UCL)
- is Professor Adjunct at Yale University School of Medicine (USA).
  
- is Director of the Centre for Computational Science (CCS) and of the Computational Life and Medical Sciences Network (CLMS) at UCL.
- <https://www.ucl.ac.uk/chemistry/people/peter-coveney>
  
- leads CompBioMed, A Centre of Excellence in Computational Biomedicine
- <http://www.compbioemed.eu>



- Goal: *advance the role of computationally based modelling and simulation within biomedicine.*
- Three related user communities:
  - academic,
  - industrial and
  - clinical researchers
- All wish to build, develop and extend such capabilities in line with the increasing power of high performance computers.
- Three distinct exemplar research areas:
  - cardiovascular,
  - molecularly-based and
  - neuro-musculoskeletal medicine.



1 of 9 European Centres of Excellence in HPC  
Official start on 1 October 2016; 3 years

■ Target question:

*Can we use the genomic data from an individual candidate and predict whether a standard drug for the treatment of breast cancer will help or not?*

■ Goal:

**A demonstration of feasibility  
with the power of high performance computing**

■ Key questions:

- Provide an answer to the question above
- Determine how to use IT-Infrastructures for this question
- Detect insufficiencies of using IT-Infrastructures for this question
- Derive a workflow for utilizing HPC in daily operation

- Running on all cores of SuperMUC Phase1+2



- Docking simulation of potentials drugs for breast cancer
- 37 hours total run time
- 241,672 cores
- 8.900.000 CPU hours
- 5 Terabytes of data produced

EU CoE CompBioMed

<http://www.compbio-med.eu>

EU Projects COMPAT and MAPPER

<http://www.compat-project.eu>

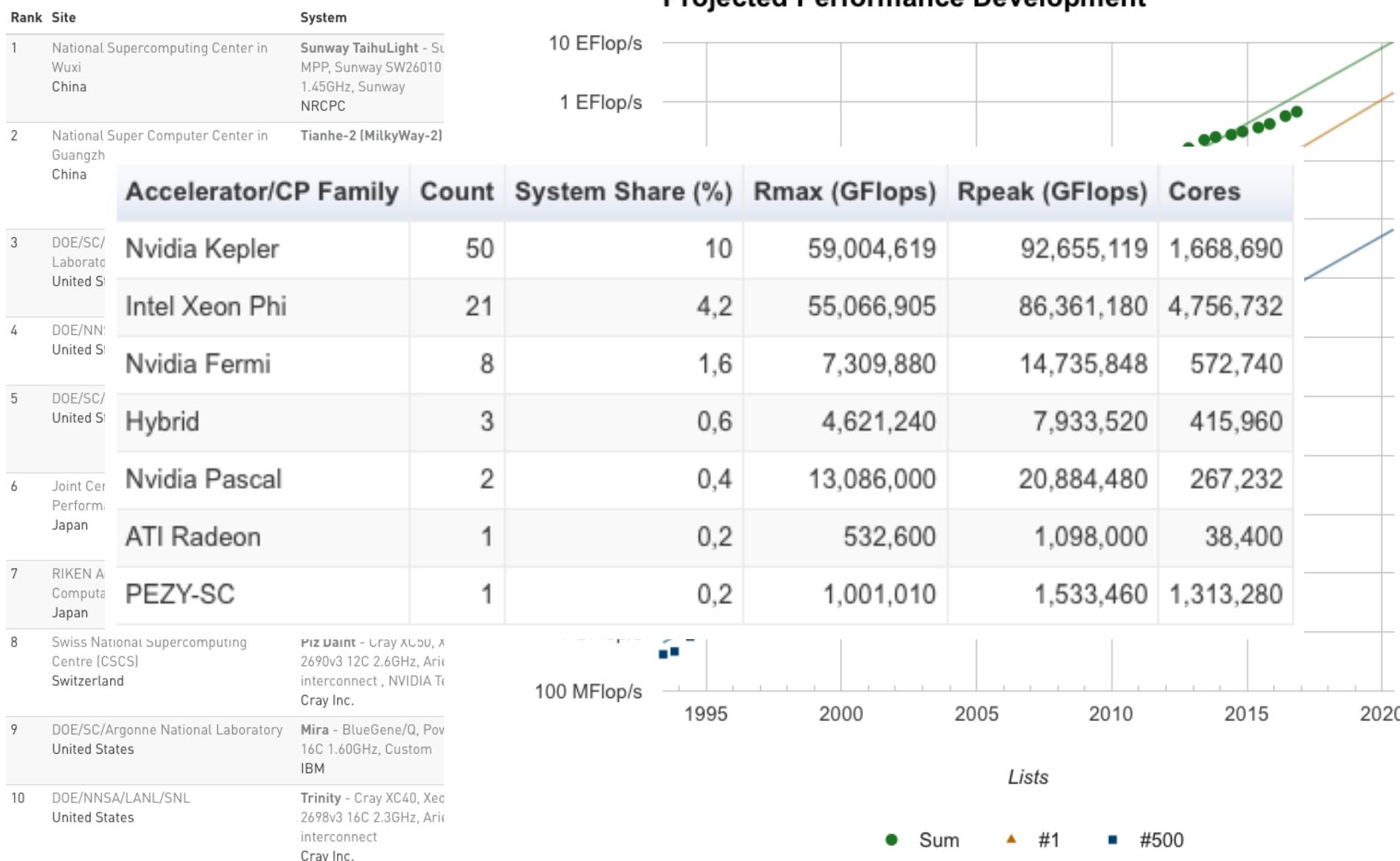
Until today:

- HLRB-II (pre-SuperMUC): Top 500 06/2007: 56,5 Tflop/s
- SuperMUC Phase 1: Top 500 06/2012: 2897 Tflop/s

Coming up:

- **SuperMUC NG (Next Generation)** – Procurement on-going

# Consulting the Top 500 List - www.top500.org

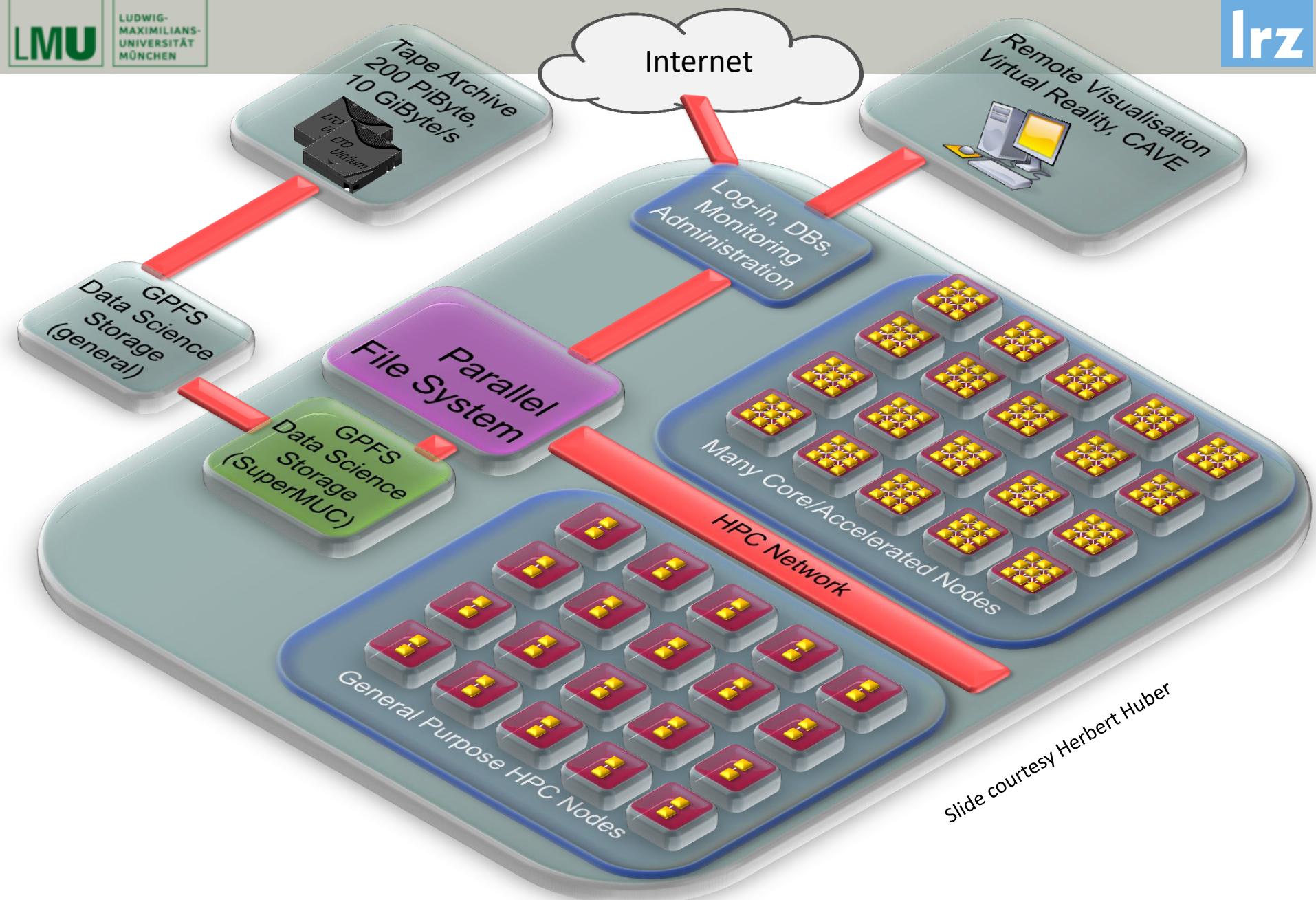


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Coming up:

- **SuperMUC NG (Next Generation)** – Procurement on-going
- **Selection criteria:**
  - LRZ application mix (compute, memory, bandwidth characteristics)
    - Number of cores
    - Memory per core
    - Interconnect
  - Accelerators (Manycore, GPGPU, ...)
  - Virtualization (Docker, Cloud, ...)
  - Workflow engines, HTC applications, ...
  - Power consumption (in total, over time, ...)



## Conclusions

- Excellent research needs excellent tools
- Supercomputers provide the highest possible computational performance, interconnectivity and memory capacity
- The complexity of (super-)computers (such as SuperMUC NG) is steadily increasing (not only on the extreme scale)
- Demand of domain science drives computer science research to new frontiers
- Users need the possibility to execute (and optimize) their codes on the full size machines
- The **LRZ Partnership Initiative Computational Science (piCS)** tries to improve user support

<http://www.sciencedirect.com/science/article/pii/S1877050914003433>

1. Choose focus topics to serve as lighthouse
  - National agreement within GCS: LRZ focuses on Environment (& Energy)
2. Choose user communities
  - Already active at LRZ?
  - Not active at LRZ?
3. Invite them for introductory piCS Workshops
  - Show faces & tour
  - Discussion on joint topics, requirements, interests, ...
4. Establish links between communities and specific points-of-contact
  - Whom to talk to, if there are questions?
  - When to talk to them? In general, as early as possible
  - Maybe, place people into the research groups (weekly, for a certain period)
5. Run joint lectures (e.g. hydrometeorology and computer science)
6. Apply for joint projects
7. Use **HPC Machines** efficiently to do science

# The new SuperMUC petascale system and applications

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Contributions from: A. Bode, A. Stamatakis, L. Czech, A. Frank, M. Brehm, H. Huber, M. Bader, F. Jamitzky, A. Parodi, ...