

MODULAR SUPERCOMPUTING: a system-wide orchestration of heterogeneous resources

ADAC, 24.01.2022 | Estela Suarez (JSC)



Mitglied der Helmholtz-Gemeinschaft

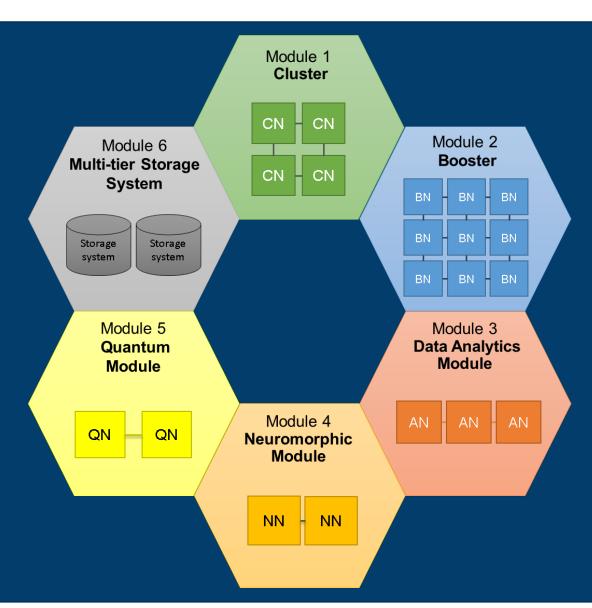
OUTLINE

• System architecture

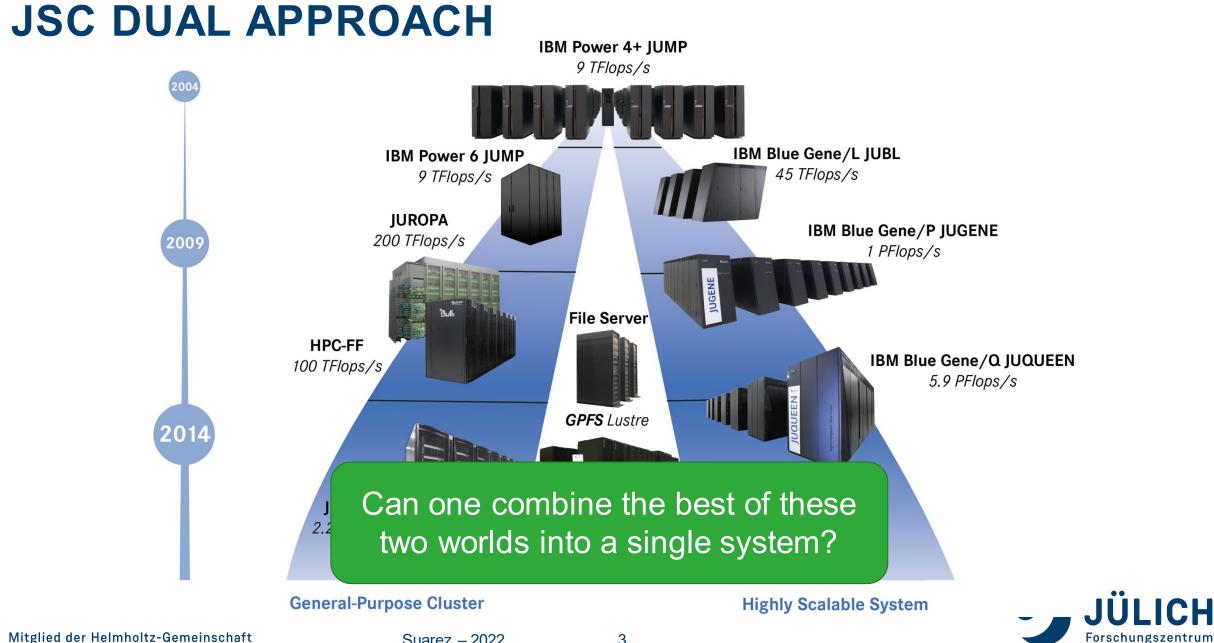
- From dual architecture to the Modular Supercomputing Architecture (MSA)
- Hardware implementations of MSA

• Software

- Software stack
- ParaStation Modulo
- Scheduler
- Application experience
- Conclusions and next steps







THE DEEP PROJECTS

2011-2021: The DEEP projects

- **DEEP** (2011 2015)
 - Introduced Cluster-Booster architecture
- **DEEP-ER** (2013 2017)
 - Added I/O and resiliency functionalities
- **DEEP-EST** (2017 2021)
 - Modular Supercomputer Architecture

2021-2024 The SEA projects

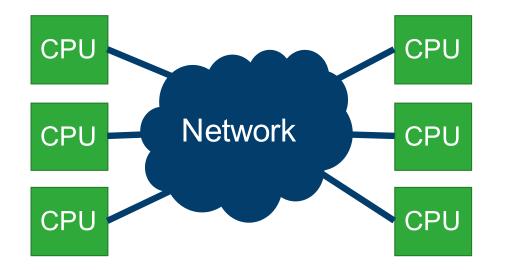
• DEEP-SEA, IO-SEA, RED-SEA





HOMOGENEOUS

General Purpose Cluster



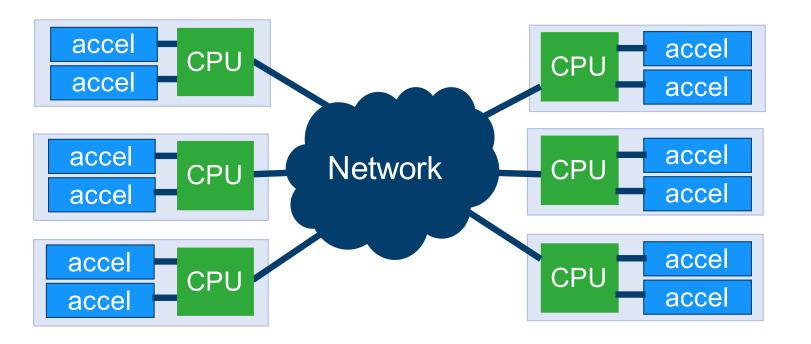


Nodes contain only CPUs



HETEROGENOUS MONOLITHIC

Every node contains accelerators (e.g. GPUs)



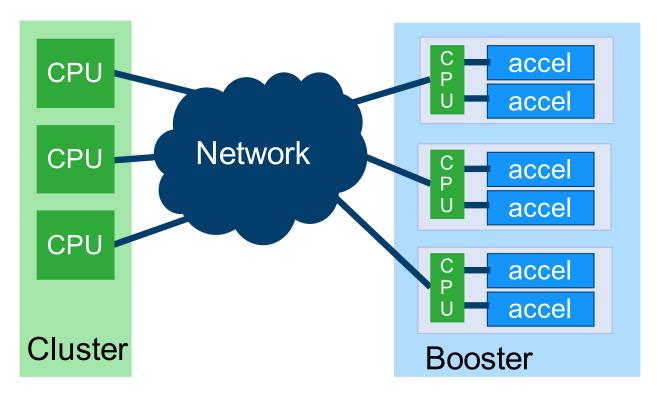
+: Energy efficient

- +: Easy management
- -: Static assignment
 - of accelerators to CPUs
- -: Difficult to efficiently share resources
- Every node contains CPU(s) and some accelerator
- All nodes are equal → "monolithic"



HETEROGENOUS MODULAR

Different nodes are grouped in "modules"



t: Energy efficient

- +: Better scalability
- +: High flexibility
- **+:** Dynamic resource assignment
- -: Complexity

- All nodes within one module are equal
- Different modules have different configurations \rightarrow "modular"

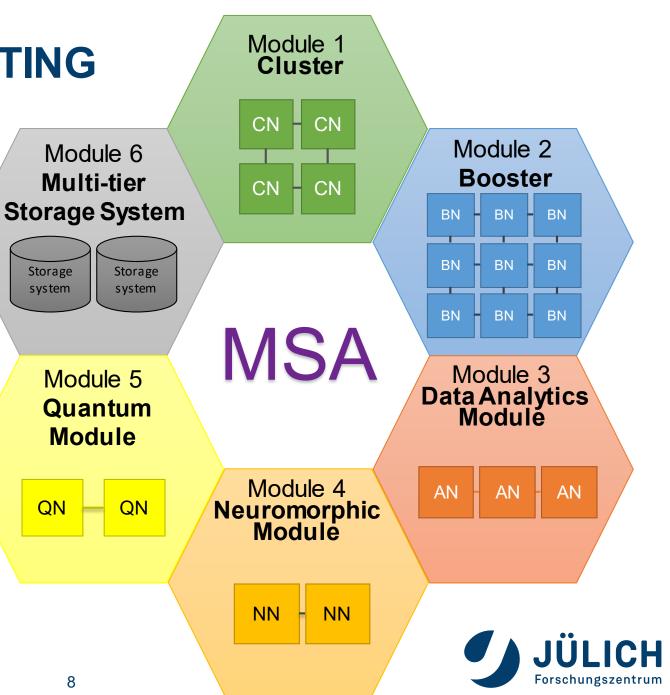


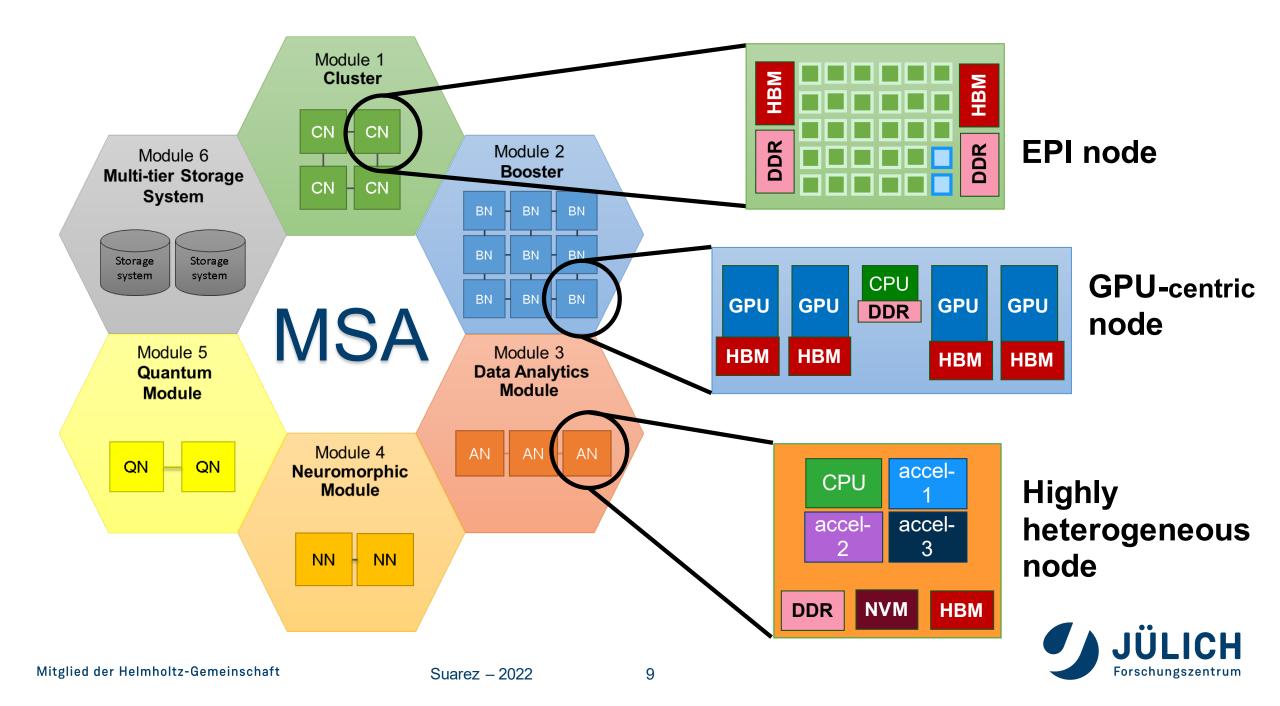


MODULAR SUPERCOMPUTING ARCHITECTURE

- Composability of heterogeneous resources
- Cost-effective scaling

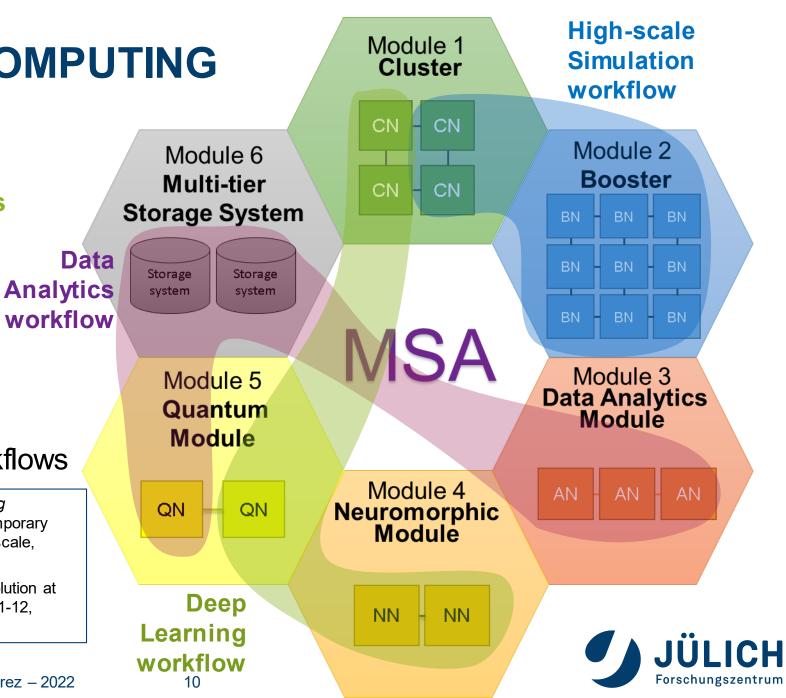
- E. Suarez, N. Eicker, Th. Lippert, "*Modular Supercomputing Architecture: from idea to production*", Chapter 9 in Contemporary High Performance Computing: from Petascale toward Exascale, Volume 3, p 223-251, CRC Press. (2019)
- E. Suarez, N. Eicker, and Th. Lippert, "Supercomputer Evolution at JSC", Proceedings of the 2018 NIC Symposium, Vol.49, p.1-12, (2018)





MODULAR SUPERCOMPUTING ARCHITECTURE

- **Composability of heterogeneous** resources
- Cost-effective scaling
- Effective resource-sharing
- Match application diversity
 - Large-scale, complex workflows -
- E. Suarez, N. Eicker, Th. Lippert, "Modular Supercomputing Architecture: from idea to production", Chapter 9 in Contemporary High Performance Computing: from Petascale toward Exascale, Volume 3, p 223-251, CRC Press. (2019)
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THE HARDWARE PROTOTYPES



2015



DEEP Prototype 128 Xeon + 284 KNC nodes InfiniBand + 1.5Gbit Extoll 550 TFlop/s

2016



DEEP-ER Prototype 16 Xeon + 8 KNL nodes 100Gbit Extoll 40 TFlop/s

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2020

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DEEP-EST Prototype 55 Cluster + 75 Booster + 16 Data Analytics 100 Gbit Extoll + InfiniBand + Eth 800 TFlop/s



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MODULAR SUPERCOMPUTER JUWELS



#7

JUWELS Cluster #44

Green500:

Intel Xeon (Skylake) processor InfiniBand **TOP500:** 2,500 compute nodes **10 PFLOP/s** peak (CPU-based) JUWEL S Booster
Entry in Nov'20
me 7402 processorRank 7 World 3IA A100 GPUsRank 1 EuropeDR DragonFly+70 PFLOP/seak (GPU-based)



TOP5 AI: Rank 4 (3 in 2021)

JUWELS is designed for simulation and

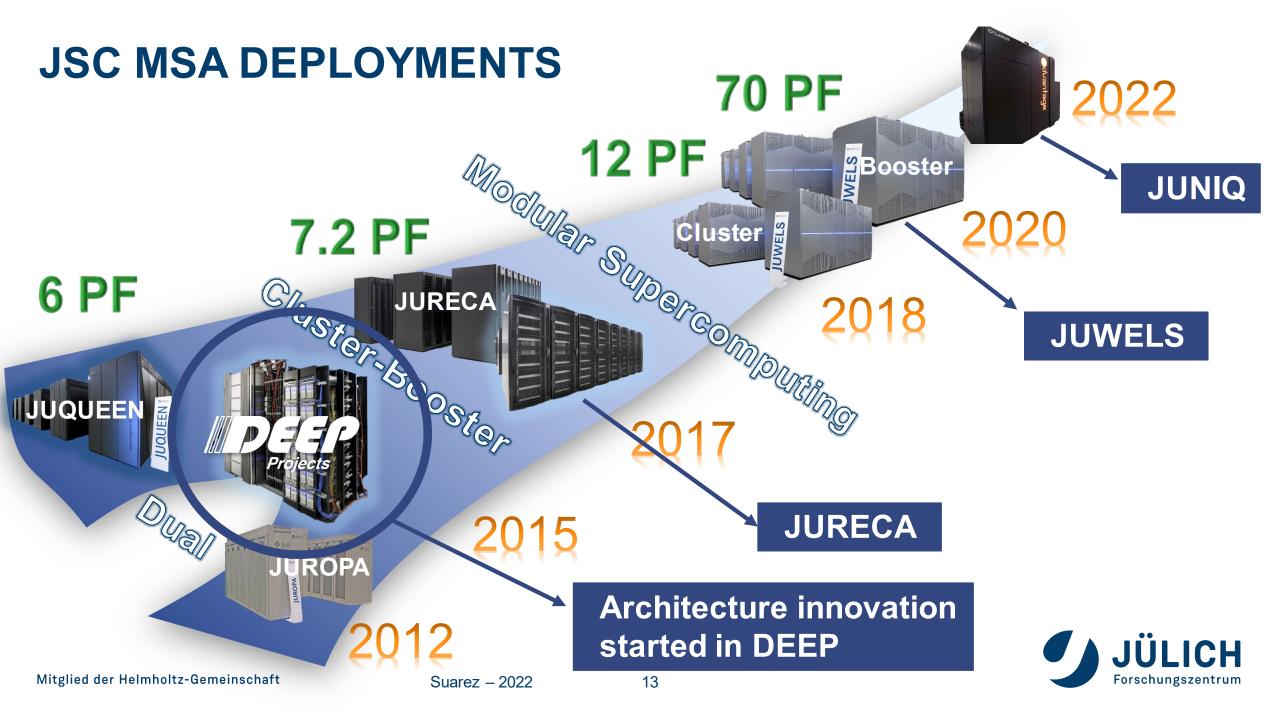
Rank 1 in TOP250



Funded through SiVeGCS (BMBF, MWK-NRW)



large-scale machine learning



MELUXINA

Cluster: 570 CPU nodes

• AMD EPYC 7H12, 2× 64C @2.6GHz, 512 GB (~ 4 GB / core)

Booster: 200 GPU nodes

- AMD EPYC 7452, 2× 32C @ 2.35GHz, 512 GB (~ 8 GB / core)
- 4× NVIDIA A100 Ampere, 40GB HBM2

Smaller partitions

20 Large Memory nodes: CPU node with 4096 GB, 1.92 TB SSD
20 FPGA nodes: CPU node with, 2× Stratix FGPA10MX (16GB HBM)
20 Cloud nodes: CPU node with 4096 GB, 1.92 TB SSD

System-wide

- NVIDIA/Mellanox InfiniBand HDR 200 Gb/s
- Atos BullSequana XH2000
- ParTec ParaStation Modulo Software

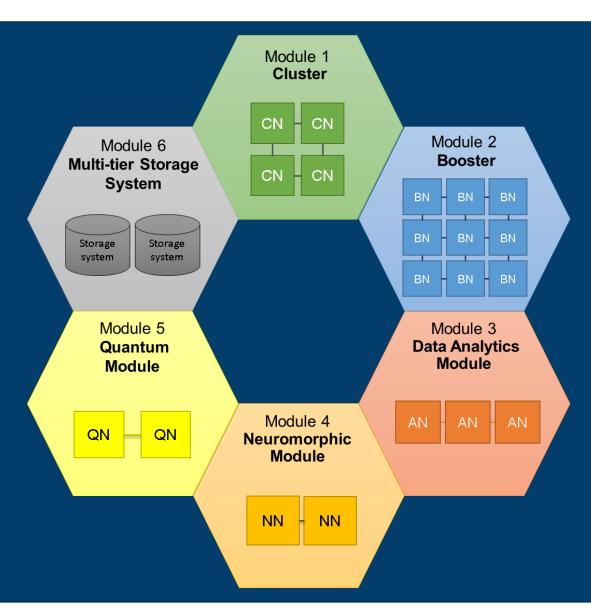
Source: Valentin Plugaru

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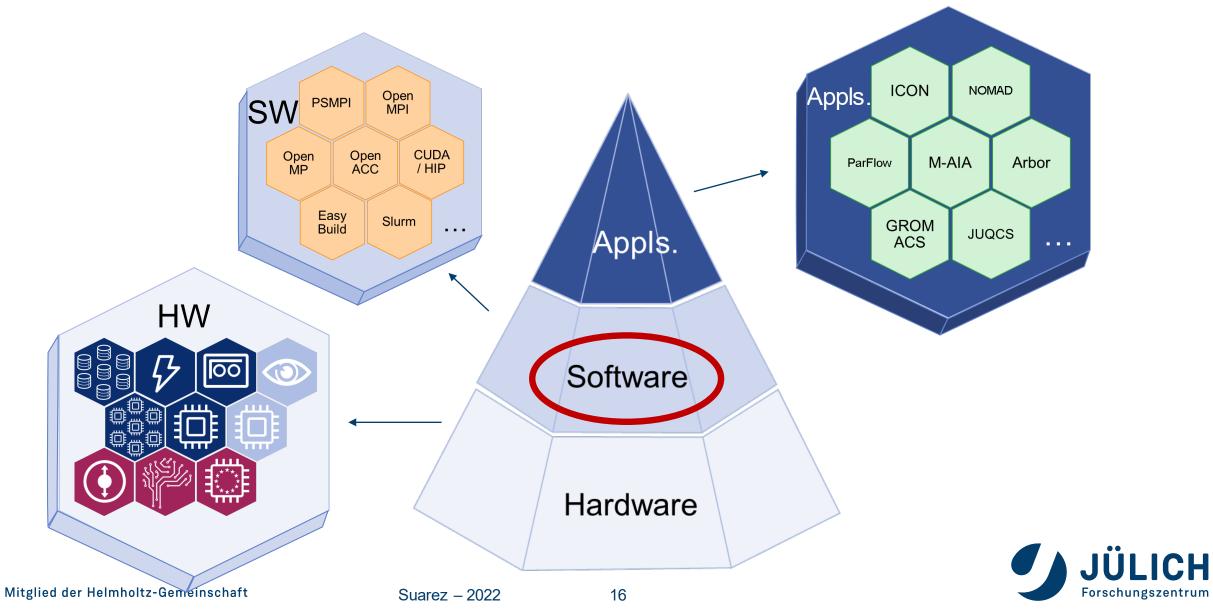
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MATCHING APPLICATIONS AND HARDWARE



SOFTWARE ENVIRONMENT





ParaStation MODULO

scalasca 🗖







- Scheduler: Slurm, psslurm (ParaStation Modulo)
- Filesystem: BeeGFS, GPFS
- **Compilers**: Intel, GCC, NVIDIA HPC SDK
- **Debuggers**: Intel Inspector, TotalView
- Programming: ParaStation MPI, OpenMP, OmpSs, CUDA
- Performance analysis tools: Scalasca, Score-P Extrae/Paraver, Vampir, Intel Advisor, VTune...
- Benchmarking tools: JUBE

SIONIib

• I/O Libraries: SIONlib, SCR, HDF5,...

Eicker et al., Bridging the DEEP Gap - Implementation of an Efficient Forwarding Protocol, Intel European Exascale Labs - Report 2013 34-41
Clauss et al., Dynamic Process Management with Allocation-internal Co-Scheduling towards Interactive Supercomputing, COSH@HiPEAC, (2016)

ParaStation Modulo



- ParaStation ClusterTools
 - Tools for system provisioning and system management
- ParaStation HealthChecker & TicketSuite
 - Automated error detection & error handling
 - Ensuring integrity of the computing environment

Source: Thomas Moschnv

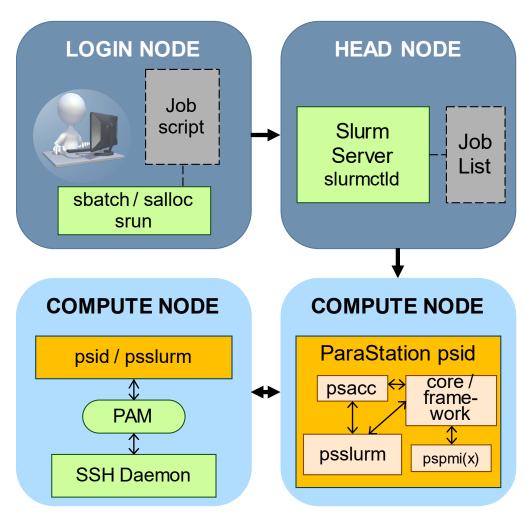
- Keeping track of issues
- Powerful analysis tools
- ParaStation Process Management & ParaStation MPI
 - Runtime environment tuned for the largest distributed memory supercomputers
 - Optimally support the **Modular Supercomputing Architecture**





ParaStation Process Manager

- Scalable network of process management daemons
 - Process startup and control, I/O forwarding, ...
 - Precise resource monitoring
 - Proper cleanup after jobs
 - Daemons run on the compute nodes
- psslurm: full integration with Slurm
 - Plugin to ParaStation Management
 - Reduce number of daemons on compute nodes
 - Replace node-local Slurm daemon
 - Integration with ParaStation HealthChecker
 - Possible to fix problems and add unique features







Source: Thomas Moschny

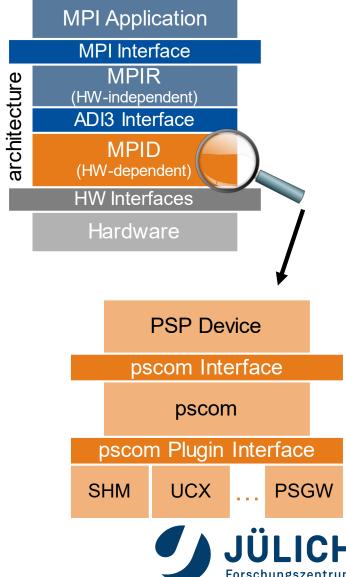
ParaStation

ParaStation MPI Library

- Based on MPICH 3.4.2 (MPI-3.1 compliant)
 - Supports MPICH tools (tracing, debugging, ...)
 - MPICH layers beneath ADI3 replaced by ParaStation PSP Device
 - Powered by pscom low-level communication library
 - Maintains MPICH ABI compatibility
- Support for various transports and protocols via pscom plugins
 - Support for InfiniBand, Omni-Path, Extoll, (soon BXI)
 - Multiple transports / plugins can be used concurrently
 - Gateway capability via PSGW plugin
 - CUDA awareness via GPUDirect
- Proven scaling up to ~3,500 nodes and ~140,000 procs. / job







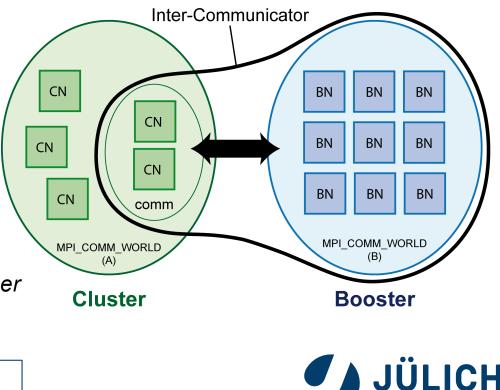
MPICH

ParaStation Global MPI for MSA

- An MPI application can run:
 - Using only Cluster nodes
 - Using only Booster nodes
 - Distributed over Cluster and Booster
 - In this case two executables are created
 - <u>Collective offload</u> process
 - Transparent data exchange via MPI
- ParaStation Global MPI
 - Uses MPI_Comm_spawn()
 - Collective spawn groups of processes from Cluster to Booster (or vice-versa)
 - Inter-communicator
 - Connects the 2 MPI_COMM_WORLD
 - Contains all parents on one side and all children on the other
 - Returned by MPI_Comm_spawn for the parents
 - Returned by MPI_Get_parent by the children

 Clauss et al., Dynamic Process Management with Allocation-internal Co-Scheduling towards Interactive Supercomputing, COSH@HiPEAC, (2016)

- One can also start two parts of a code and connect them via MPI_Connect()
- Or have one single common MPI_COMM_WORLD and split it into subcommunicators via MPI_Comm_Split()



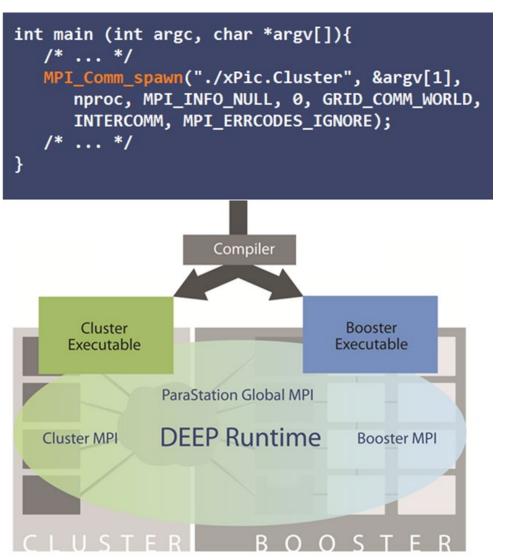
ParaStation

COMPILE AND RUN

Compilation

- Creates two executables (if different CPU architecture)
 - $\circ \quad \text{One for CLUSTER} \ \text{code}$
 - One for BOOSTER code
- Batch system
 - Reserves required resources
- Execution
 - Script starts Booster code
 - This code calls MPI_Comm_spawn() with Cluster executable
 - Optional: **xenv** to load suitable environment modules
- Runtime + Scheduler + FS
 - Detect ParaStation MPI calls
 - Distribute child binaries

salloc --partition=cluster -N 4
 : --partition=booster -N 12
srun --het-group=1 -N 4 -n 8
 ./app_booster



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Heterogeneity from user's PoV



- Slurm supports the ability to submit heterogeneous jobs (since v 17.11)
 - form job pack (het-job) allocation using colon notation for salloc, sbatch, srun
 - even allowing different executables

```
$ srun -N 1 -p part1 ./first \
    : -N 2 -p part 2 ./second
```

- Full support for job packs in ParaStation psslurm, with unique features for modular jobs:
 - Support for heterogeneous jobs with common MPI_COMM_WORLD
 - For each job in the job pack, resources can be specified individually
 - Support global resources (e.g. gateways): psgw plugin to psmgmt + spank plugin
 - o Compensates for Slurm's inability to handle global resources
 - Extends salloc, srun and sbatch
- Modular here means: Jobs across heterogeneous hardware

Source: Thomas Moschny

Either with a common MPI_COMM_WORLD, or with separated / interconnected MPI_COMM_WORLDS





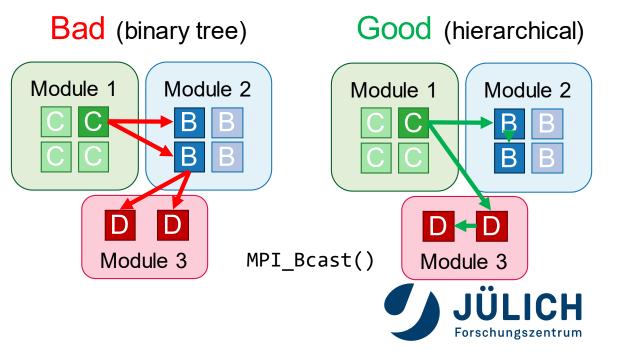
Explicit MSA-extensions to ParaStation MPI **ParaStation**

- API additions to retrieve topology information
 - Querying the module ID: MPI_Info_get (MPI_INFO_ENV, "msa_module_id", ..., value, ...);
 - Splitting communicators according to the topology:

MPI_Comm_split_type (oldcomm, MPIX_COMM_TYPE_MODULE, ..., &newcomm);

• Modularity-aware MPI collectives

- Optimized patterns for collectives that take topology of the MSA system into account
- Assumption: Inter-module communication is the bottleneck
- Dynamic updates of the communication patterns supported, e.g. for malleable jobs (experimental)



Hiearchical collectives (MSA awareness)

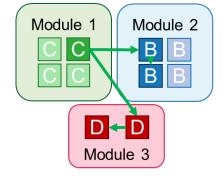
- General rules to optimize collectives \rightarrow execute these steps in order:
 - 1) Do all module-internal gathering and/or reduction operations (if required)
 - 2) Conduct the inter-module operation with a single process per module
 - 3) Perform a strict module-local data distribution
- Multi-level hierarchy awareness
 - Apply this set of rules recursively,
 i.e., node level, module level, system level
- Performance heavily depends on concrete settings, i.e.:
 - Number of processes / gateway nodes
 - Distribution of the ranks in the communicator
 - Message sizes (and hence the collective communication pattern)

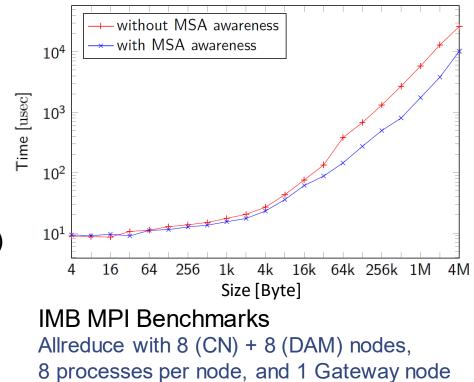


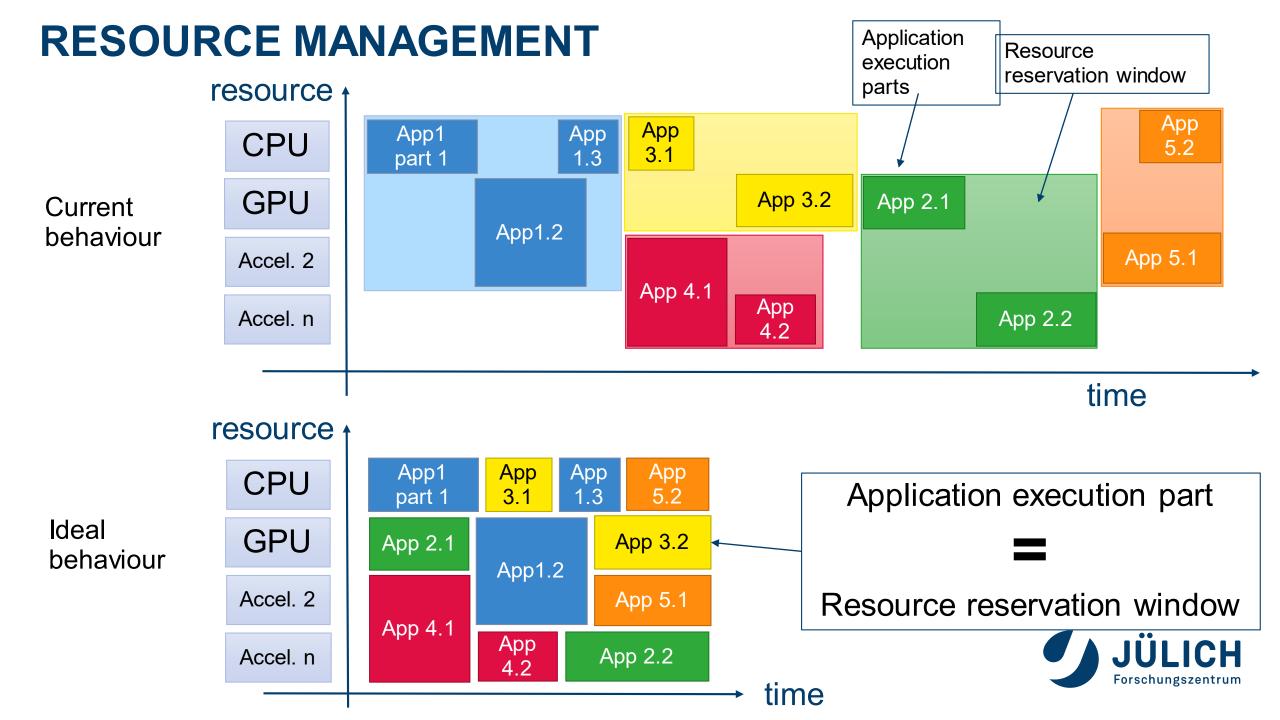
Source: Thomas Moschny

ParaStation

Good (hierarchical)







Improved Workflow Support (experimental)

- New parameter --delay introduced in sbatch command for job packs
 - Amount of time, the next job should wait after start of the first job in a job pack
- **Goal**: Overlapping job execution
 - Currently not supported by Slurm -
 - Whole job pack either accepted or rejected 0
 - All jobs allocated and run in parallel Ο
 - All jobs wait for allocation if any of the jobs can not be 0 allocated at the moment





Time

Typical Workflow supported

by Slurm





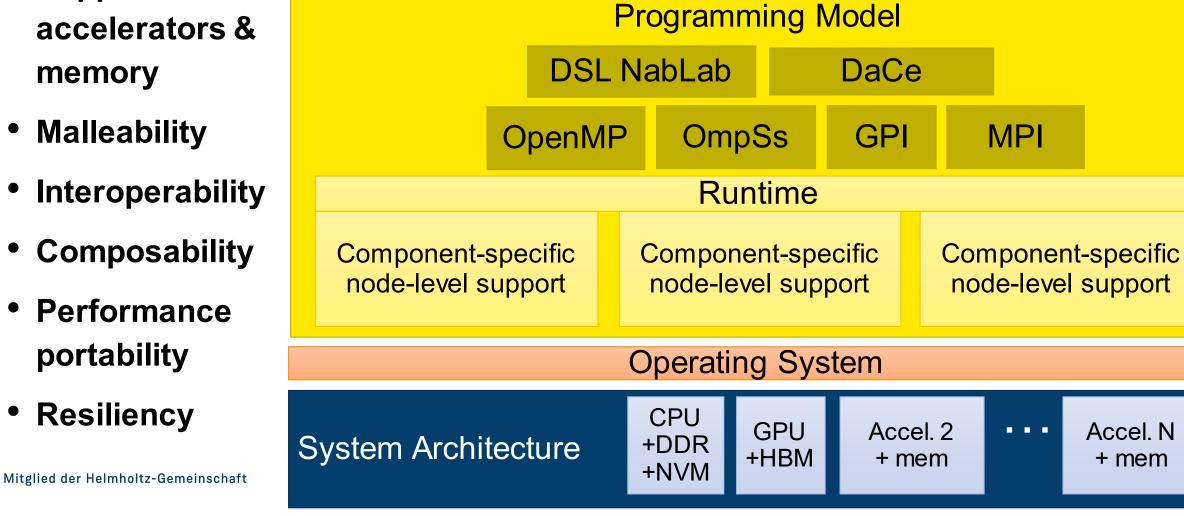
EXTENDING THE SOFTWARE STACK



Accel. N

+ mem

- Support for accelerators & memory
- Malleability
- Interoperability
- Composability
- Performance portability
- Resiliency



Application / Workflow

Quantum integration in MSA



Module 6

Multi-tier Storag System

Module

Quantun

Module

• New usage models

- Tightly coupled simulations: benefit from efficient data exchange
- Hybrid quantum-HPC simulations: combining quantum and classical algorithms
- Workflows comprising stages on the QPU, with pre- and post-processing on HPC modules
- Integrate QPU and its front-end into the managements stack
 - Low-latency connection to other modules via federated, high-speed network
 - Unified environment: Integrated in the user-, SW-, schedule- and resource- mgmt.
 - Provide "direct" access of the QPU via a web-based portal
 - Redirect portal requests through the global scheduler/resource manager
 - Pseudo-shared usage model as prerequisite
- Exact requirements depend on the use case and are subject to research



Module 1 Cluster

Module 4 Neuromorphic

Module

NN - NN

Module 2

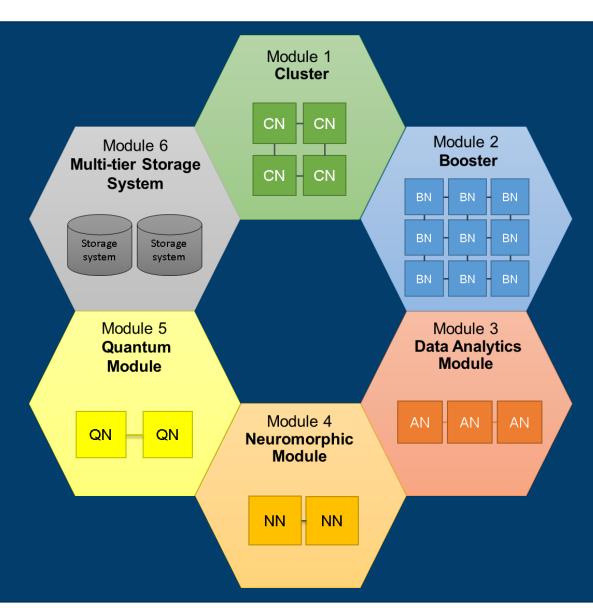
Module 3

Data Analytics

Module

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Architecture Use-Modes





Cluster-Booster use mode

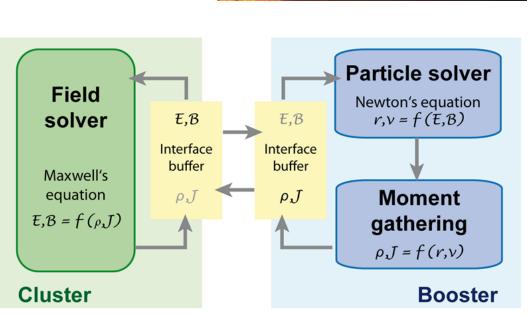
Code partition Workflow I/O forward

- Kreuzer, et al., Application Performance on a Cluster-Booster System. IPDPSW HCW (2018) [10.1109/IPDPSW.2018.00019]
- Kreuzer et al. The DEEP-ER project: I/O and resiliency extensions for the Cluster-Booster architecture. HPCC'18 proceedings (2018) [10.1109/HPCC/SmartCity/DSS.2018.00046]
- Wolf et al., PIC algorithms on DEEP: The iPiC3D case study. PARS-Mitteilungen 32, 38-48 (2015)
- Christou et al., EMAC on DEEP, Geoscientific model devel.(2016) [10.5194/gmd-9-3483-2016]
- Kumbhar et al., Leveraging a Cluster-Booster Architecture for Brain-Scale Simulations, Lecture Notes in Computer Science 9697 (2016) [10.1007/978-3-319-41321-1_19]
- Leger et al., Adapting a Finite-Element Type Solver for Bioelectromagnetics to the DEEP-ER Platform. ParCo 2015, Advances in Parallel
- Computing, 27 (2016) [10.3233/978-1-61499-621-7-349]

Application use case: xPic

- Space Weather simulation
 - Simulates plasma produced in solar eruptions and its interaction with the Earth magnetosphere
 - Particle-in-Cell (PIC) code
 - Authors: KU Leuven
- Two solvers:
 - Field solver: Computes electromagnetic (EM) field evolution
 - Limited code scalability
 - Frequent, global communication
 - **Particle solver**: Calculates motion of charged particles in EM-fields
 - $_{\circ}$ Highly parallel
 - $_{\circ}$ Billions of particles
 - Long-range communication

A. Kreuzer, J. Amaya, N. Eicker, E. Suarez, *"Application performance on a Cluster-Booster system",* 2018 IEEE International Parallel and Distributed Processing Symposium Workshops (IPDPSW), HCW (20th International Heterogeneity in Computing Workshop), Vancouver (2018), p: 69 - 78. [doi: 10.1109/IPDPSW.2018.00019]



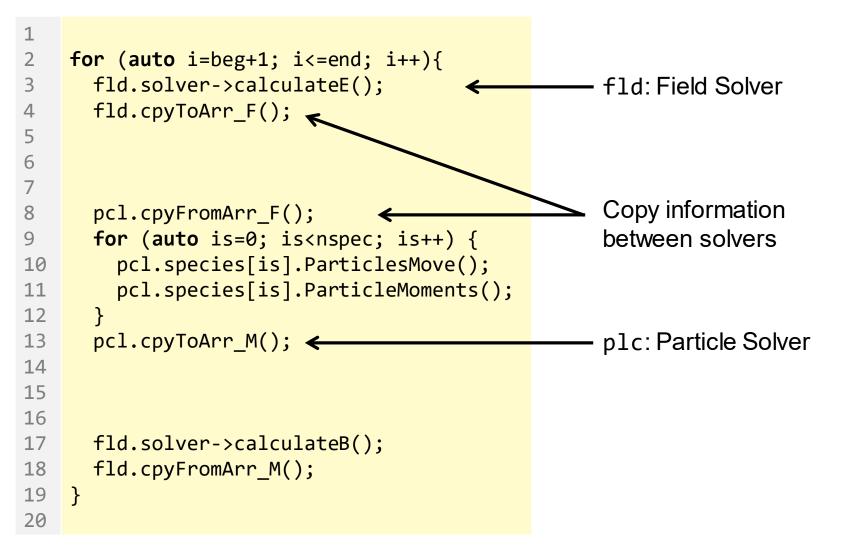




ATHOLIEKE UNIVERSITEIT COC

xPic – ORIGINAL CONFIGURATION







xPic – CODE PARTITION



```
#ifdef CLUSTER
1
   for (auto i=beg+1; i<=end; i++){</pre>
2
      fld.solver->calculateE();
3
4
      fld.cpyToArr F();
5
      ClusterToBooster();
6
      // Auxiliary computations
      ClusterWait();
7
8
9
10
11
12
13
   BoosterToCluster();
14
15
   BoosterWait();
16
      fld.solver->calculateB();
17
     fld.cpyFromArr M();
18
19 }
                                         #endif
20 #endif
```

```
#ifdef __BOOSTER__
for (auto i=beg+1; i<=end; i++){</pre>
```

```
ClusterToBooster();
```

```
ClusterWait();
pcl.cpyFromArr_F();
for (auto is=0; is<nspec; is++) {
   pcl.species[is].ParticlesMove();
   pcl.species[is].ParticleMoments();
}
pcl.cpyToArr_M();
BoosterToCluster();
// I/O and auxiliary computations
BoosterWait();</pre>
```



Field solver: 6× faster on Cluster Particle solver: 1.35 × faster on Booster 35

• Overall performance gain:

1× 28% × gain compared to Cluster alone
node 21% × gain compared to Booster alone

8× 38% × gain compared to Cluster only
nodes 34% × gain compared to Booster only

- 3%-4% overhead per solver for C+B communication (point to point)

A. Kreuzeret al. "*Application Performance on a Cluster-Booster System*", 2018 IEEE IPDPS Workshops (IPDPSW), Vancouver, Canada, p 69 - 78 (2018) [10.1109/IPDPSW.2018.00019]

xPic – (1-NODE) PERFORMANCE RESULTS

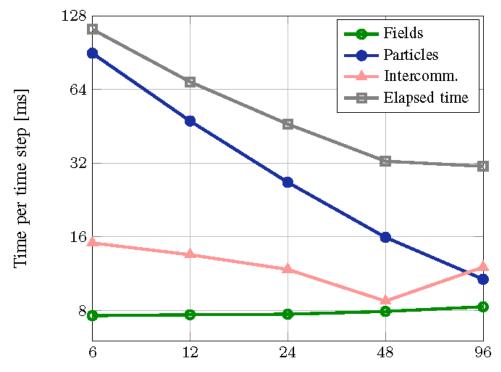
	45			
Runtime [s]	40	■ Cluster	=	
	35	☐ Booster		
	30			
	25			
	20			
Ru	15			
	10	P 774		
	5			
	0			
		Fields	Particles	Total
		Γ	Γ	7
		#cells per node	4096	KATHOLIEKE UNIVERSITE
		#particles per cell	2048	LEUVEN
		Compilation flags	-openmp, -mavx (Cluster) -xMlC-AVX512 (Booster)	
', 2018 IEEE IPDPS Workshops .00019]				



xPic – STRONG SCALING on JURECA



Variable-ratio modular strong scaling



(4 Cluster nodes)

Number of Booster nodes

#cells per node	36864	
#particles per cell	1024	
#blocks per MPI process	12, 32 or 64	
Compilation flags	-mavx (Cluster) -openmp, xMIC-AVX512 (Booster)	

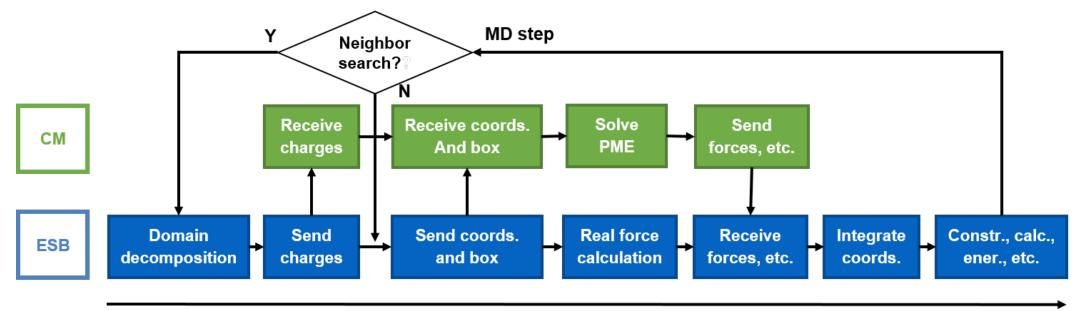
- Code portions can be scaled-up independently
 - Particles scale almost linearly on Booster
 - Fields kept constant on the Cluster (4CNs)
- A configuration is reached where same time is spent on Cluster and Booster
 - Additional 2× time-saving is enabled via overlapping



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GROMACS: multi-module usage in MD simulations

- Best mapping on MSA depends on the problem size and aims at optimizing the computational load
 - <10⁴ particles: only on Cluster (CPU)
 - ~ 10⁵ particles: Booster or DAM (Data Analytics Module)
 - >10⁶ particles (large macromolecules): pair interactions on GPU, run PME on CPUs



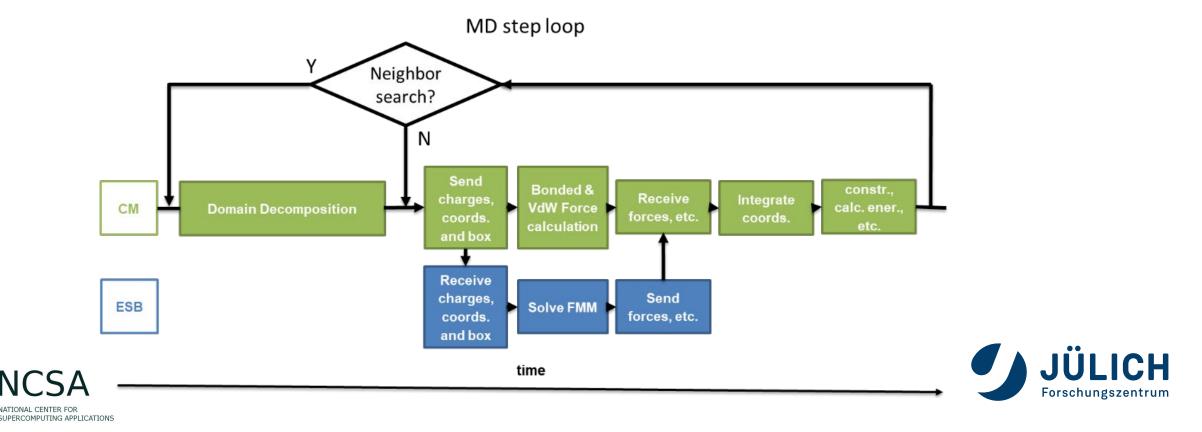




JPERCOMPUTING APPLICATIONS

GROMACS: multi-module usage in MD simulations

- Best mapping on MSA depends on the problem size and aims at optimizing the computational load
 - <10⁴ particles: only on Cluster (CPU)
 - ~ 10^5 particles: Booster or DAM (Data Analytics Module)
 - >10⁶ particles (large macromolecules): pair interactions on GPU, run PME on CPUs
 - Very large volume (>10⁶ nm³): Replace PME with FMM (Fast Multipole Method) running on ESB



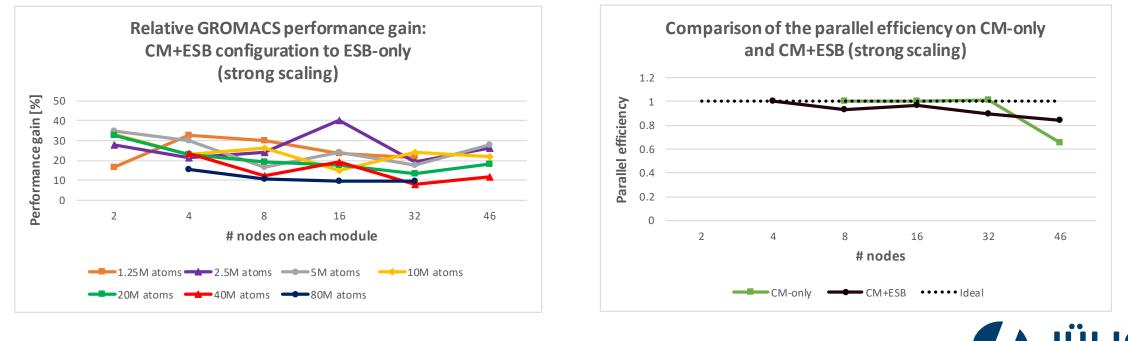
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einschaft

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NextDBSCAN: multi-module usage in ML

- Parallel algorithm for density-based clustering of arbitrary data sets
 - Performance and flexibility gain by running on multiple modules

See presentation in ADAC applications group:

DBSCAN Clustering and Modular Supercomputing: Lessons Learned

Ernir Erlingsson (University of Iceland)

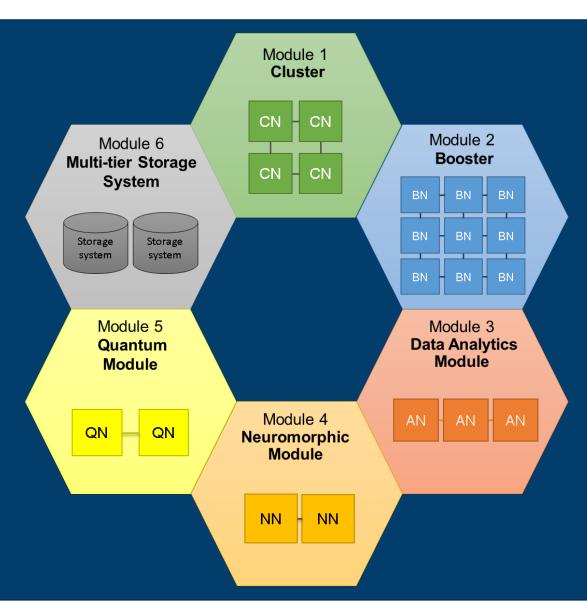
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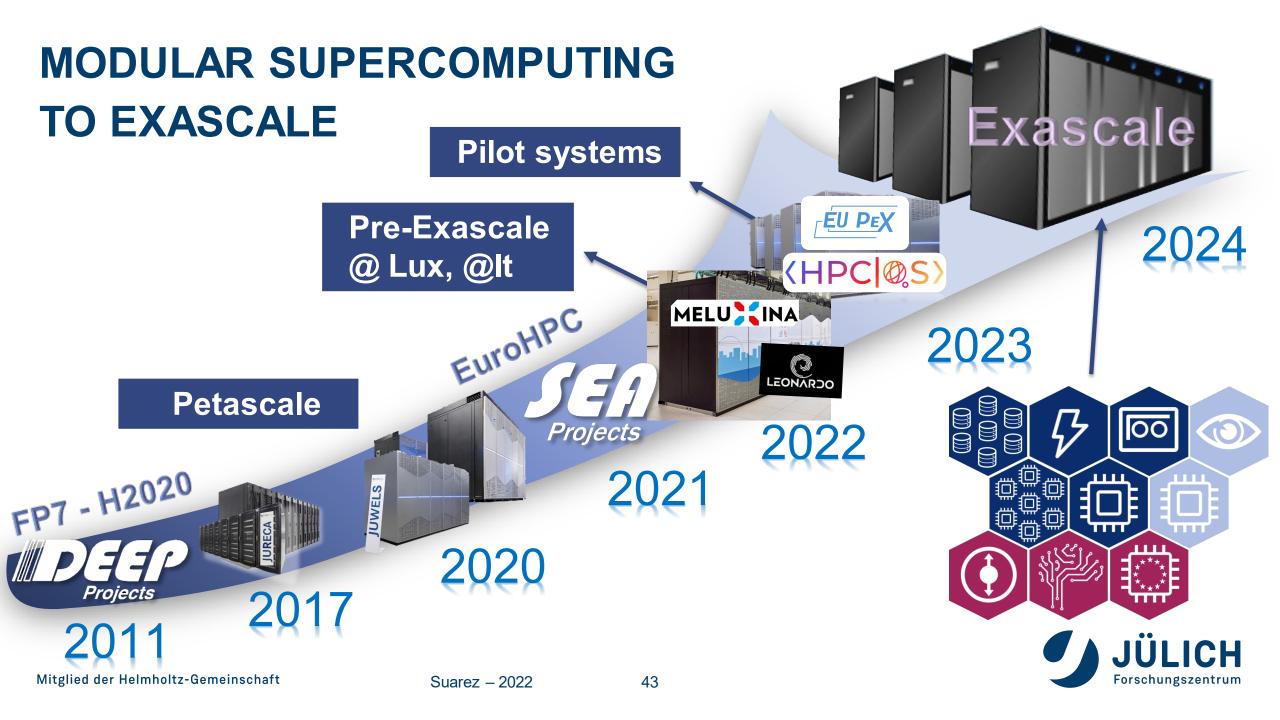
CONCLUSIONS

- The Modular Supercomputing Architecture (MSA)
 - Orchestrates heterogeneity at system level
 - Allows scaling hardware in economical way (Booster \rightarrow Exascale)
 - Serves very diverse application profiles
 - Maximum flexibility for users, without taking anything away (still can use individual modules)

• Distribute applications on the MSA give each code-part a suitable hardware

- Straight-forward implementation for workflows
- Partition at MPI-level interesting for multi-physics / multi-scale codes
- Monolithic codes do not need to be divided
- Current / Upcoming implementations of MSA
 - DEEP system, JURECA, JUWELS
 - MELUXINA (Luxembourg EuroHPC Petascale system)
 - EUPEX and HPCQS pilots
 - ... Exascale !





THANK YOU!



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