



A EUPEX-ChEESA cooperation: the SPECFEM3D(++) example; context, preliminary results and next steps

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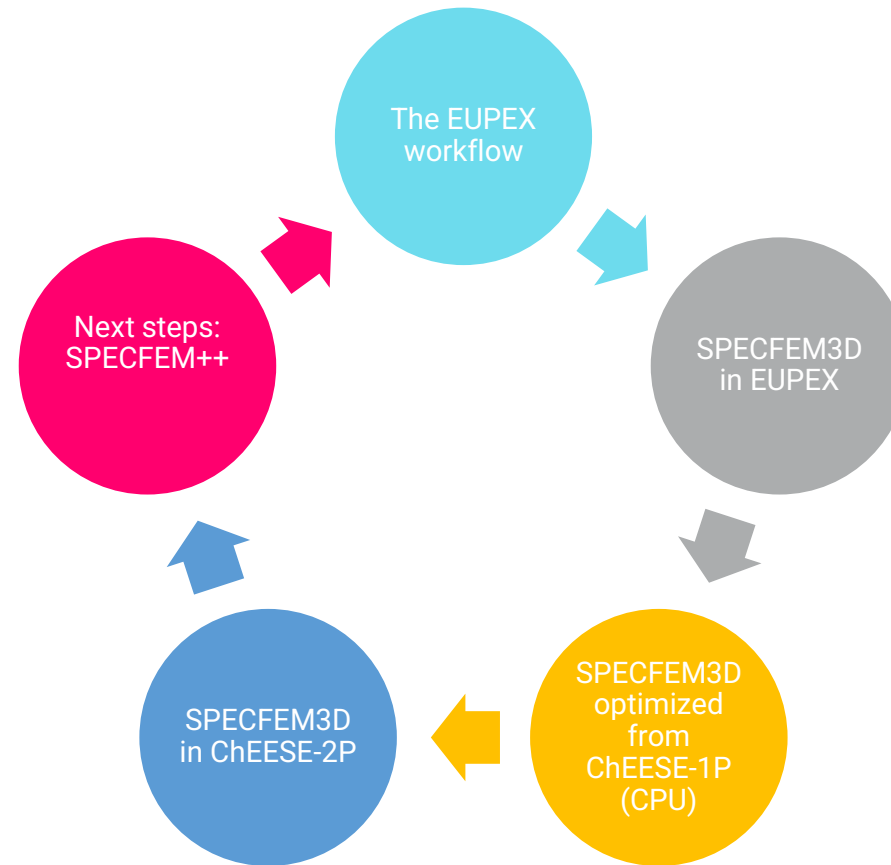
EUPEX Forum 2024



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Agenda



From faults to buildings

Integrated seismological/engineering **probabilistic** workflow

regional scale

ground
motion
acceleration
scenarios

local scale

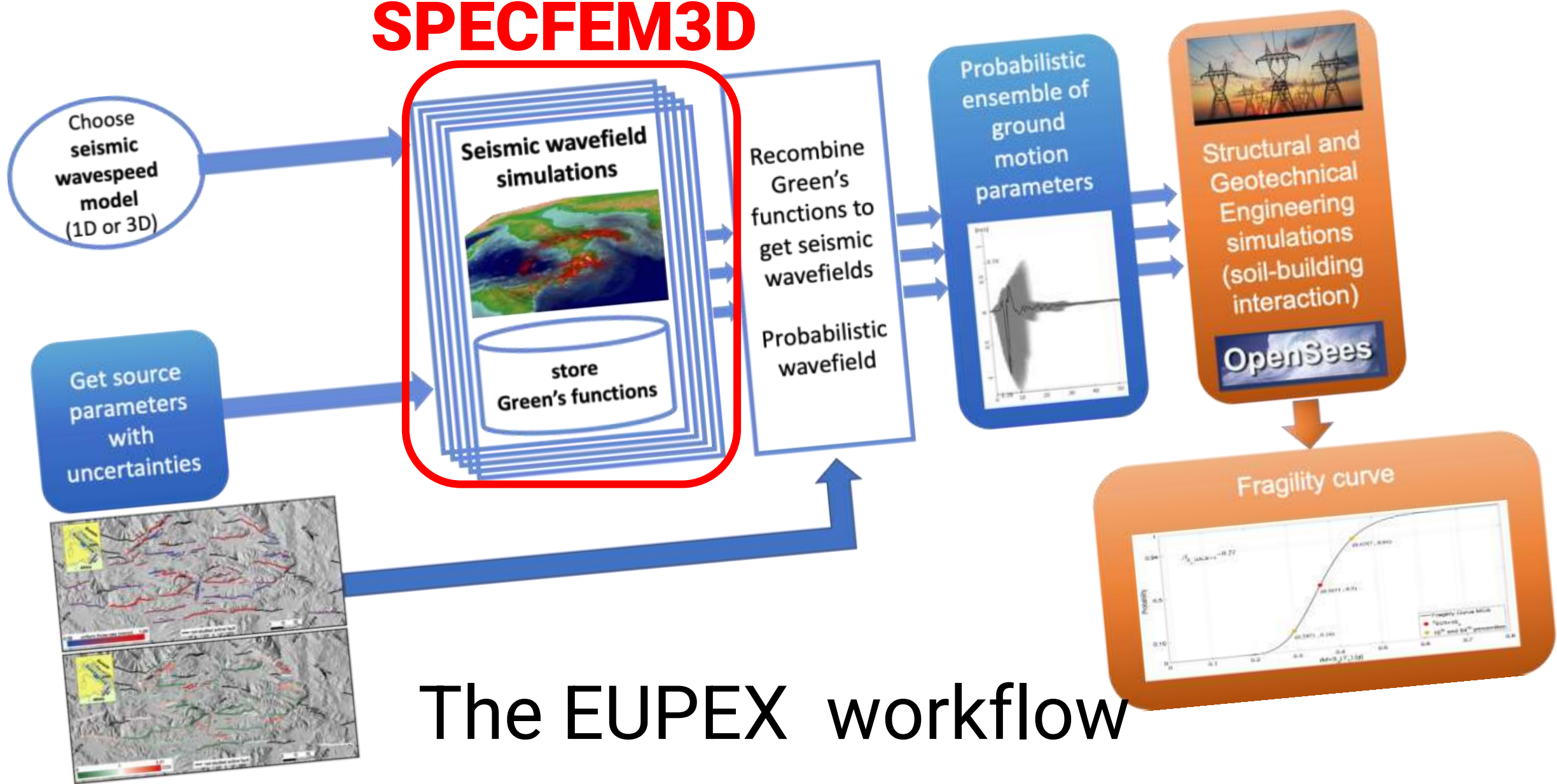
impact on
strategic
buildings

Coupling applications with high-performance parallel streams.

- HPC for natural disaster resilience

HPC4NDR

SPECFEM3D



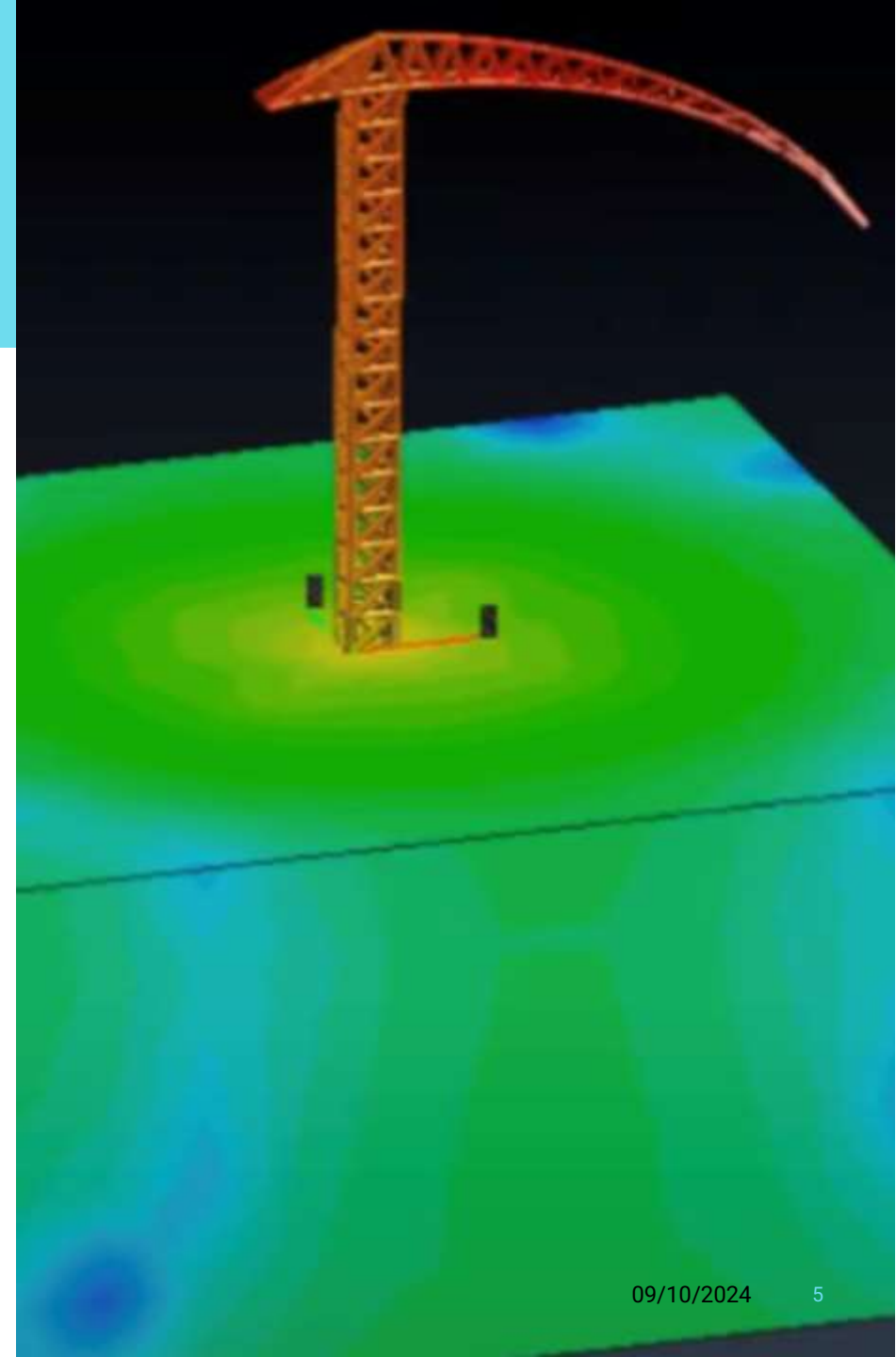
The EUPEX workflow

- From geology to probabilistic seismic engineering analysis

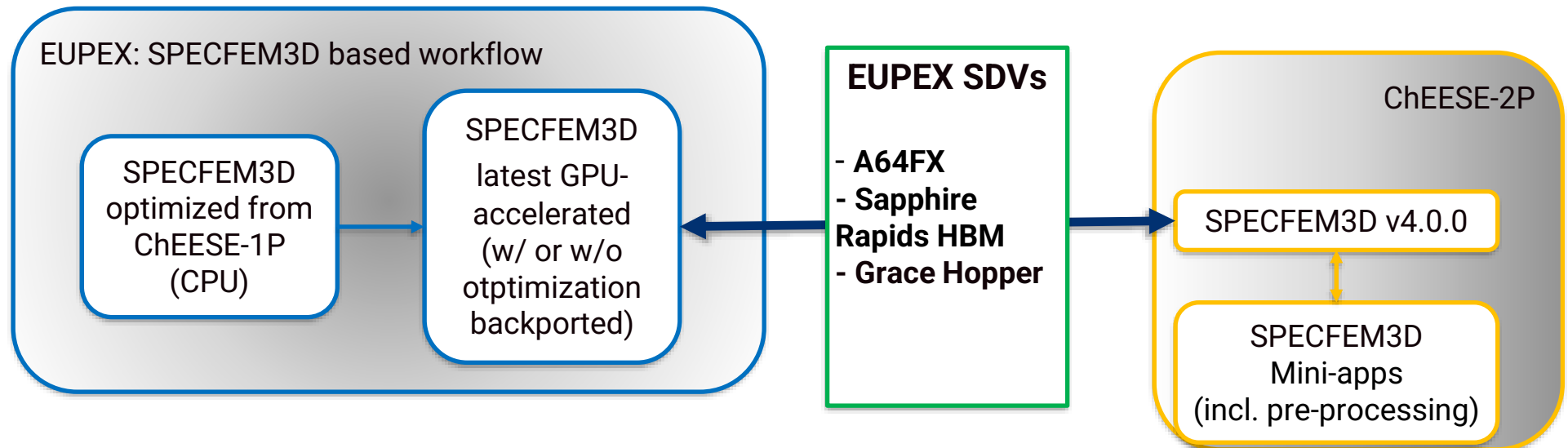
Structural and Geotechnical Engineering simulations

➤ The Engineering applications will be performed using the OPEN-SOURCE framework **OpenSeeS**

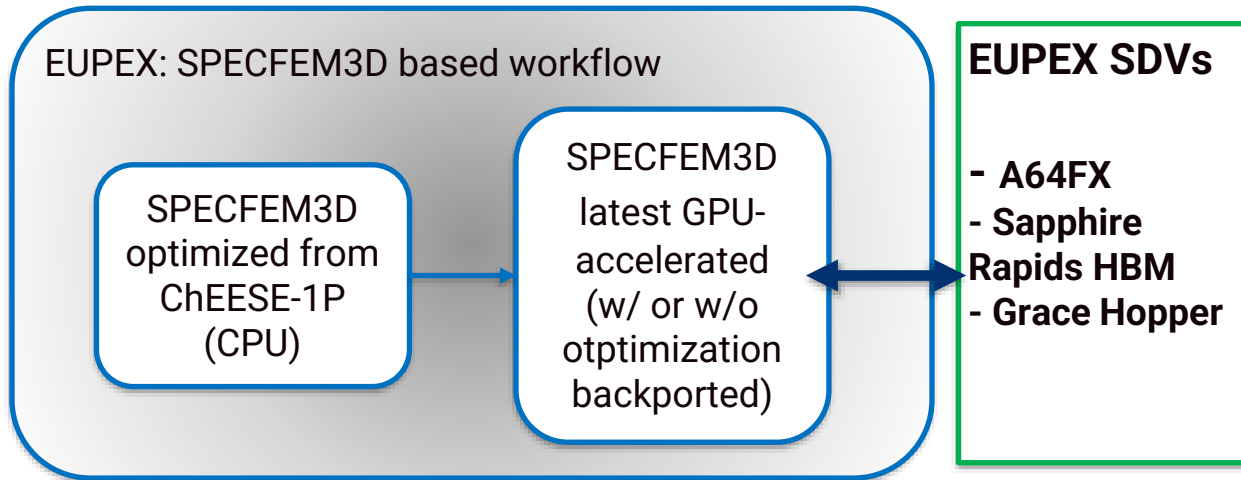
- Soil-Structure Interaction (SSI), of large domains
- HPC capabilities in developing detailed large scale seismic probabilistic frameworks



ChEESE-EUPEX Collaboration Diagram for SPECFEM3D



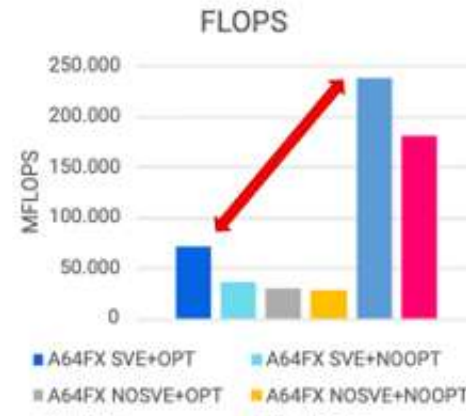
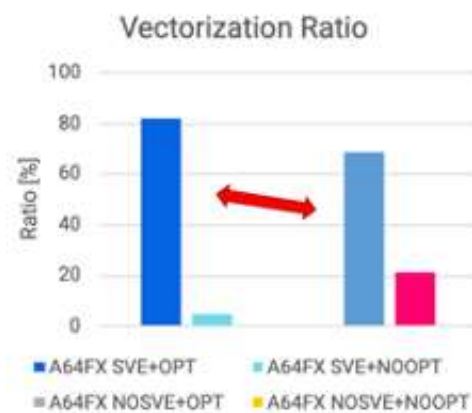
SPECFEM3D IN EUPEX



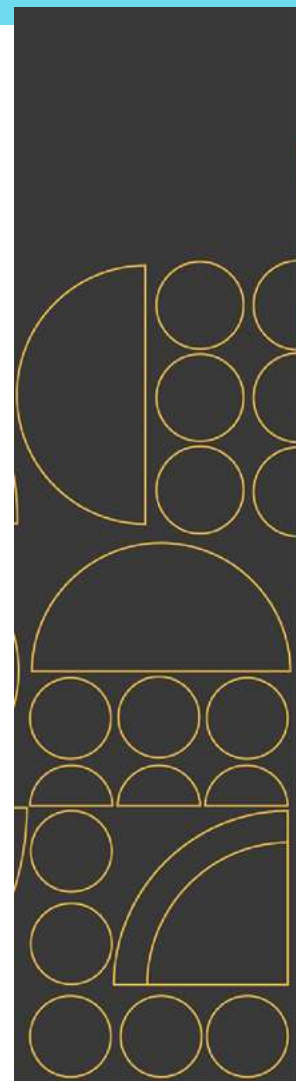
- Target a complete workflow including SPECfem3D
- Focus on optimized version from ChEese-1P (CNRS/Atos/Eviden version)
- Two main branches:
 - Elastic simulation (vectorization improved for CPU architectures)
 - Full SPECfem3D (including seismic tomography features): still missing some optimization/tuning (anisotropic materials). Available in [zenodo](#)
- But target latest accelerated version for production level test case and associated workflow
- A64FX performance tests done, will continue on Grace Hopper (soon)

SPECFEM3D optimized from ChEESE-1P (CPU)

- Performance assessment of the ChEESE-1P optimized version on Irene (A64FX)
- Comparison with G100 (x86 machine in CINECA)
- Preliminary work done evaluating HBM improvement (on-going)



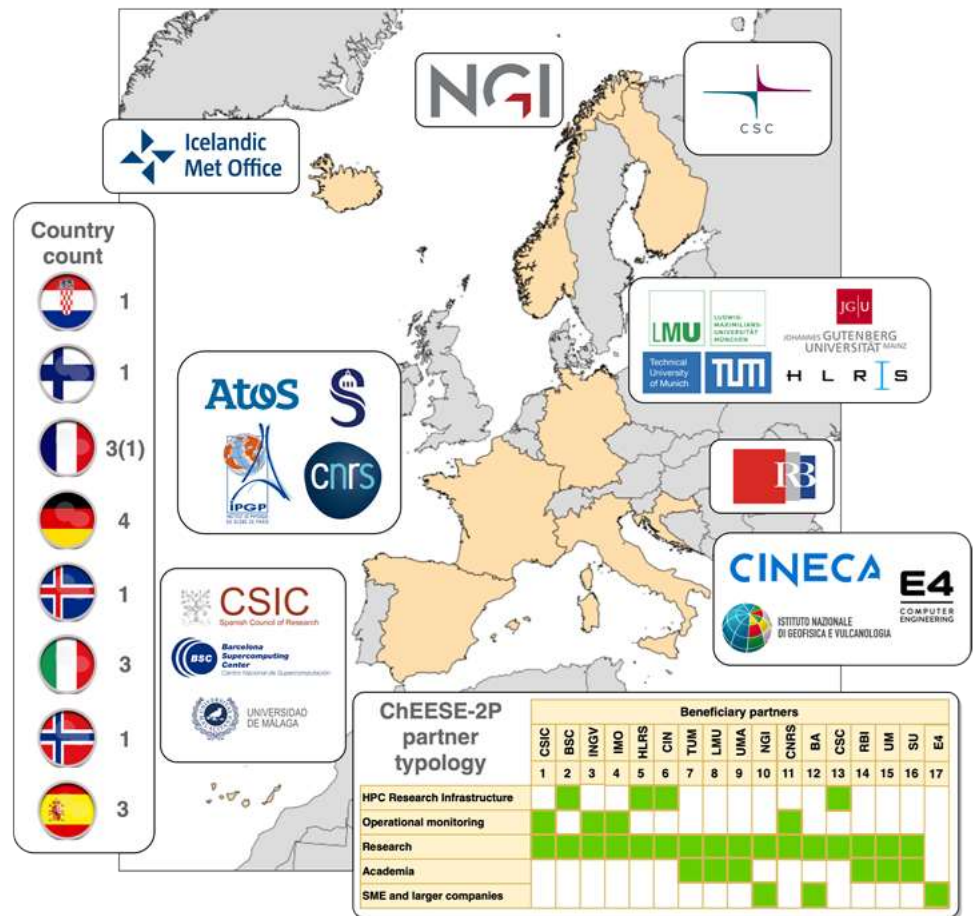
ChEESE-2P CoE



- 17
Beneficiary Organisations
 From 8 different countries
- 1
Affiliated Entity
 IPGP (affiliated to CNRS)

- 4
HPC tier-0 Centers
 BSC, CIN, HLRS, CSC
- 3
Private Companies
 BA, NGI, E4
- 4
Operational Monitoring
 CSIC, INGV, IMO, CNRS
- 6
Academia
 TUM, LMU, UMA, RBI, UM, SU

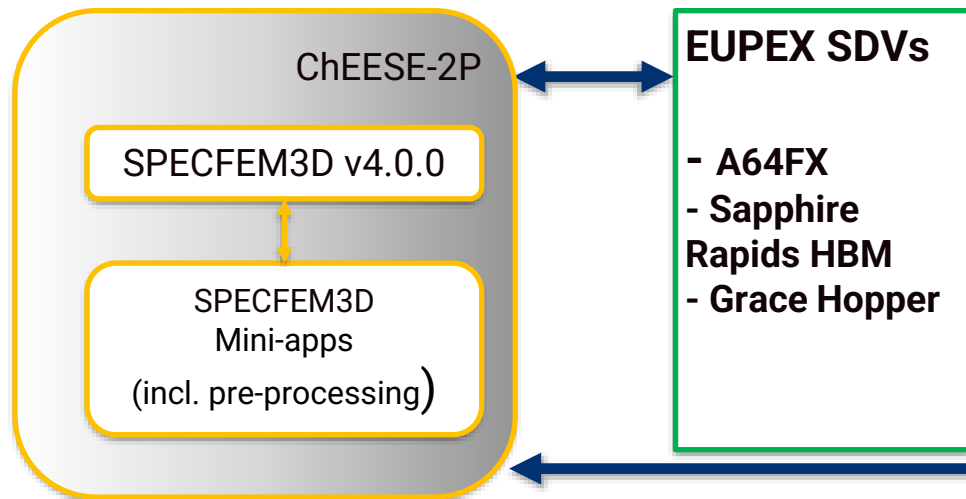
ChEESE-2P Consortium Composition



SPECFEM3D IN ChEESE-2P



ChEESE



- WP2 “Exascale technical challenges in flagship codes”
 - Leonardo booster code audit (scaling etc.) and suggestions using the full app (published in D2.1)
 - Ongoing work for D2.2 (Dec. 2024) on performance portability campaign on GPU (MN5, LUMI, Leonardo) on the mini-apps.
 - Optimization in WP2: Reduce memory access, try different compression strategies (i.e., compressed arrays), need to check precision and do some tests, now on CPU then on GPU.
- WP3 “Co-design with European HPC vendors and technologies”
 - Focus on mini-apps, target ARM SVE and HBM before EUPEX availability.

Area	No	Code	Lead	ChEESE -1P
CS	1	SeisSol	LMU/TUM	yes
	2	SPECFEM3D	CNRS	yes
	3	ExaHyPE	TUM	yes
	4	Tandem	LMU	no
MHD	5	xSHELLS	CNRS	yes
T	6	HySEA	UMA	yes
V	7	FALL3D	CSIC	yes
GD	8	OpenPDAC	INGV	no
	9	LaMEM	UM	no
GL	10	pTatin3D	SU	no
	11	Elmer/ICE	CSC	no

Tier-0 (pre-exascale)			TOP500
LUMI	Finland	~500 PFlop/s peak	5
Leonardo	Italy	~300 PFlop/s peak	6
MN5 ACC	Spain	~250 PFlop/s peak	8



More than 1000 PFlop/s peak aggregated computing power...

...but with a variety of different (accelerated) architectures

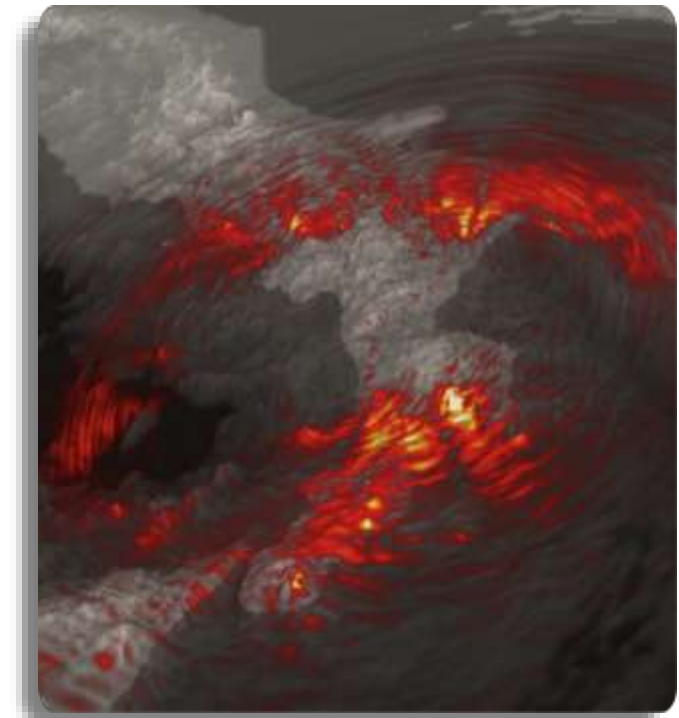
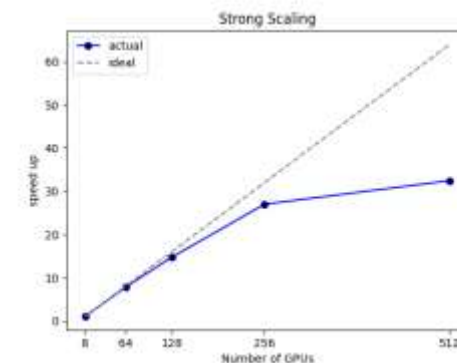
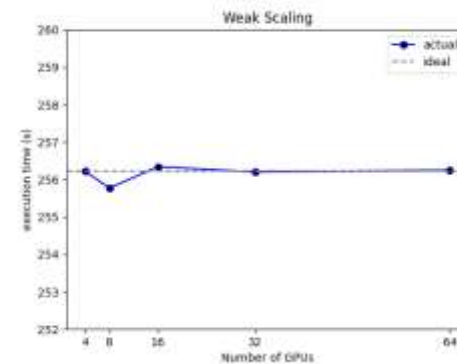
SPECFEM3D IN ChEESE-2P



▶ <https://specfem.org/>

- Spectral-element method:
 - CPU, GPU, MPI (domain decomposition)
- Solves linear seismic wave propagation in 3D models :
 - Elastic, viscoelastic, piroelastic, and fluid solid interactions
 - Implements imaging and Full Waveform Inversion for complex models
- Scalability evaluated on Leonardo booster:
 - Weak scaling : from 4 to 64 GPUs, 360448 mesh elements per GPU
 - excellent weak scaling up to 64 GPUs (and likely beyond)
 - Strong scaling :
 - good strong scaling up to 256 GPUs
 - overlapping MPI communication with GPU computation

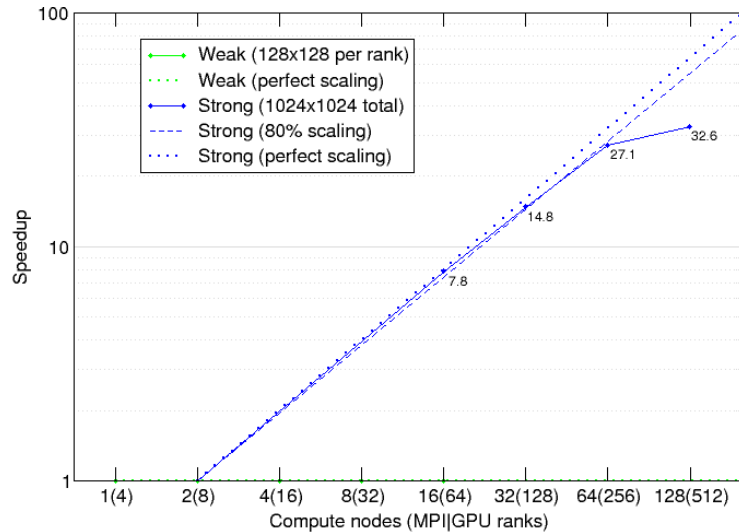
SPECFEM3D has been optimized for various GPU architectures (using primarily CUDA and HIP as GPU programming model).



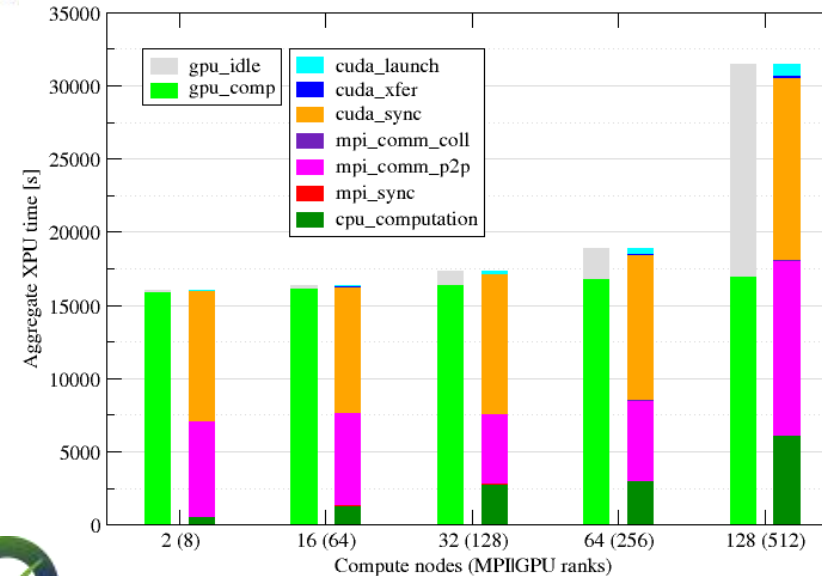
ChEESE-2P GPU assesment on SPECFEM3D



ChEESE



Problem size	1024x1024	1024x1024	1024x1024	1024x1024	1024x1024
MPI GPU ranks	8	64	128	256	512
Wall time [s]	2001.806	255.948	135.478	73.846	61.400
Global scaling efficiency	0.995	0.973	0.919	0.843	0.507
- Computation time scaling	1.000	0.987	0.972	0.950	0.937
- Parallel efficiency	0.995	0.986	0.945	0.887	0.541
-- Load balance efficiency	1.000	0.998	0.996	0.994	0.989
-- Orchestration efficiency	0.995	0.988	0.948	0.892	0.547

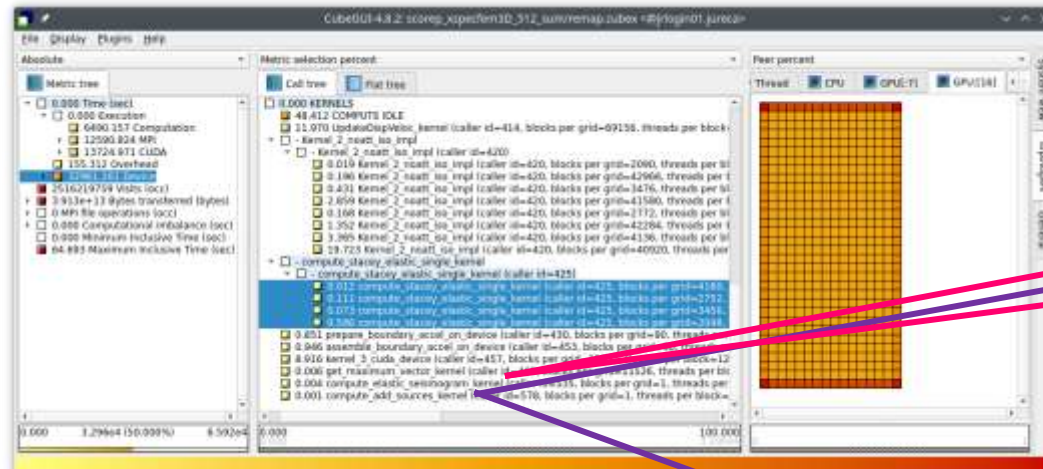


POP3_AR_002

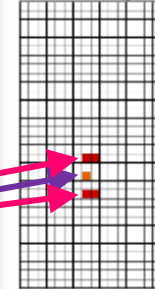
- *iterate_time* (solver) chosen as Focus of Analysis
- Good strong scaling up to 256 GPUs
- With 512 GPUs no longer able to sufficiently overlap MPI communication with CUDA kernels



ChEESE-2P GPU assesment on SPECFEM3D



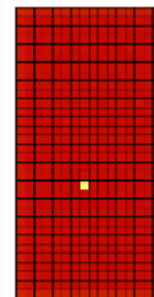
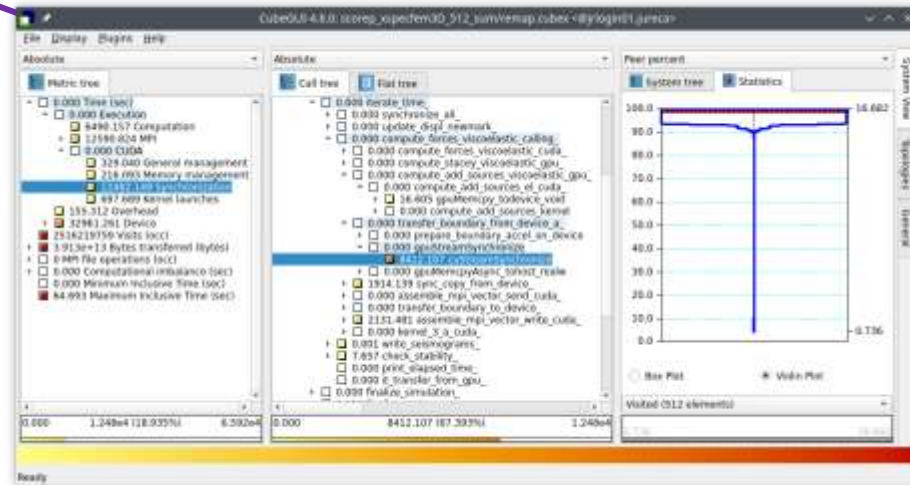
16x32 grid



- 2D domain decomposition on GPUs
- many kernels are well balanced
 - some kernels execute much faster for interior compared to edges
 - only four GPUs handle seismogram receivers
 - only one GPU (#243) executes *compute_add_sources_kernel*

compute_add_sources_kernel executed on single GPU (#243) is rather short, however, results in all other GPUs having very long synchronization times in following *transfer_boundary_from_device_a*

- over two-thirds of CUDA synch time and over 30% of total CPU time



ChEESE-2P co-design on SPECfEM3D



> Mini-apps developed by Vadim Monteiller (CNRS)

- CPU, GPU (CUDA) and MPI
- https://gitlab.com/specfem_cheese_2p/mini_specfem/specfem_mini_app

> Goals

- Ported and optimized for EUPEX and EUIPLOT
- Performance portability by testing different programming models/languages

> Status

- Performance characterization on x86 ✓
- Ported to ARM (Ampere Altra and Grace) ✓
- Vectorization optimization ✓
- HBM study ⌚
- More optimization dedicated to HBM & ARM SVE ↻



ChEESE-2P co-design on SPECFEM3D



> First optimizations to improve vectorization on main kernel

“compute_forces”:

- First attempts by hand: loop reordering, temporary arrays, etc.
- Finally: addition of !\$OMP SIMD directives leads to best performance

> Results on AMD Milan 7763 (1 socket with 64c)

> As mini-apps are all memory bound, the HBM should be effective in improving performance

> Next step: study the performance on HBM and optimize memory layout and access

Mini app	Execution time	Compute bound	Memory bound	Vectorized
elastic_aniso_mpi	259 s	77%	23%	No
elastic_iso_mpi	140s	49%	51%	No
elastic_iso_att_mpi	275 s	49%	51%	No



Add !\$OMP SIMD directives at the right spots

Mini app	Execution time	Speedup	Compute bound	Memory bound	Vectorized
elastic_aniso_mpi	166 s	1,6	61%	39%	Yes
elastic_iso_mpi	116 s	1,2	40%	60%	Yes
elastic_iso_att_mpi	240 s	1,1	37%	63%	Yes

Next steps: SPECFEM++

- > On production: use of the accelerated version of SPECFEM3D (on one node: 30x speedup CPU vs GPU on Jean Zay machine).
- > CUDA, HIP and CPU version in the same code with 2 different memory structures (one for CPU et one for GPU), difficult to maintain.
- > Maintain the CPU version anyway.

Why use Kokkos?

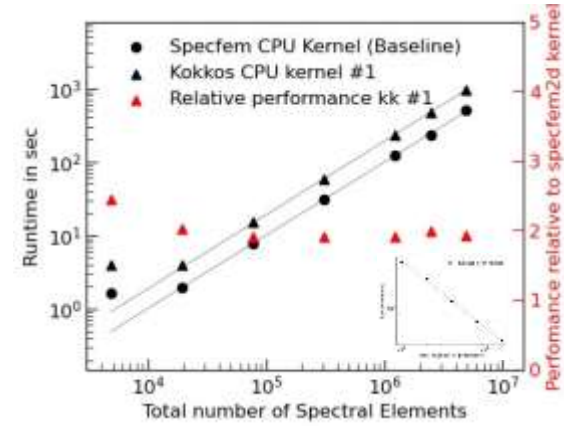
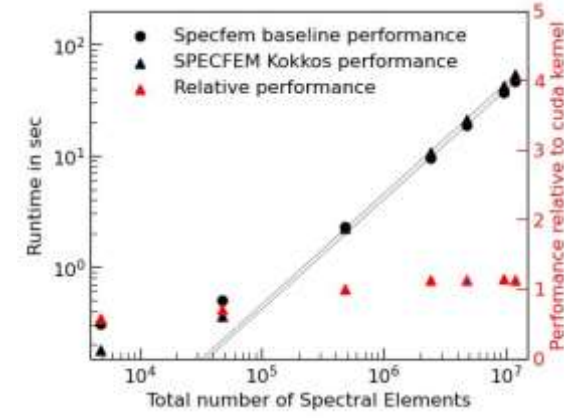
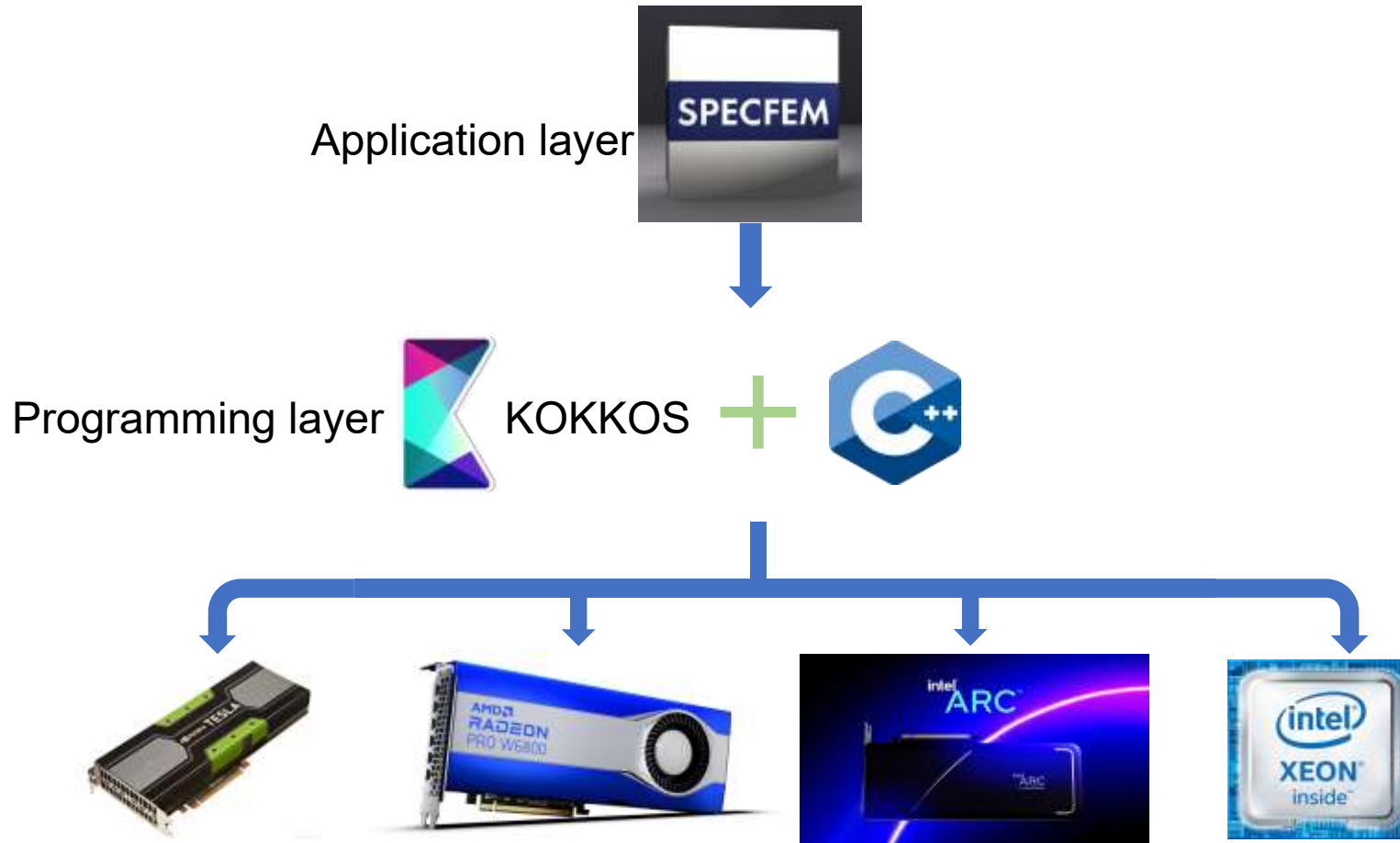
	CUDA/HIP/DPC++	OpenMP 5/ OpenACC	Kokkos
Portability across architectures	No. Need to write separate kernels for every architecture	Yes. Single source code with pragma-based approach	Yes. Single source code implemented using Kokkos functions
Performance	Optimized performance	Tough to optimize for performance	Very good performance
Cost of portability	Very high	Medium	High
Cost of maintenance	Very high. Newer architectures might require tuning of kernels	Low. Assuming compilers do a good job of implementing the standard	Low. Assuming Kokkos backend is always optimized
Compiler dependence	N/A	These are standards, vendors have a flexibility on implementation	N/A
Fortran Support	No. Could use bindings	Yes	No. Could use bindings

Why use Kokkos?

PRINCETON UNIVERSITY

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Next steps: SPECFEM++



Acknowledgement

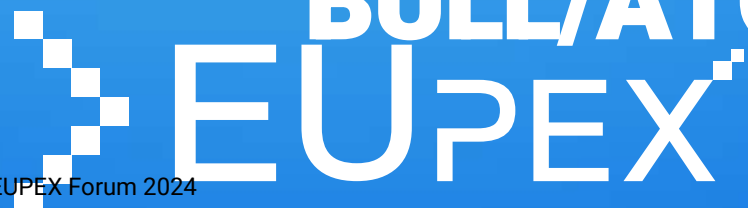
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