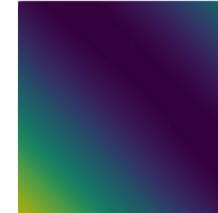
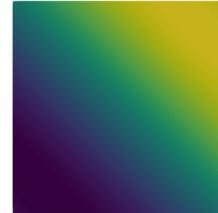


Using Trixi.jl for Adaptively Coupling Multiphysics Problems

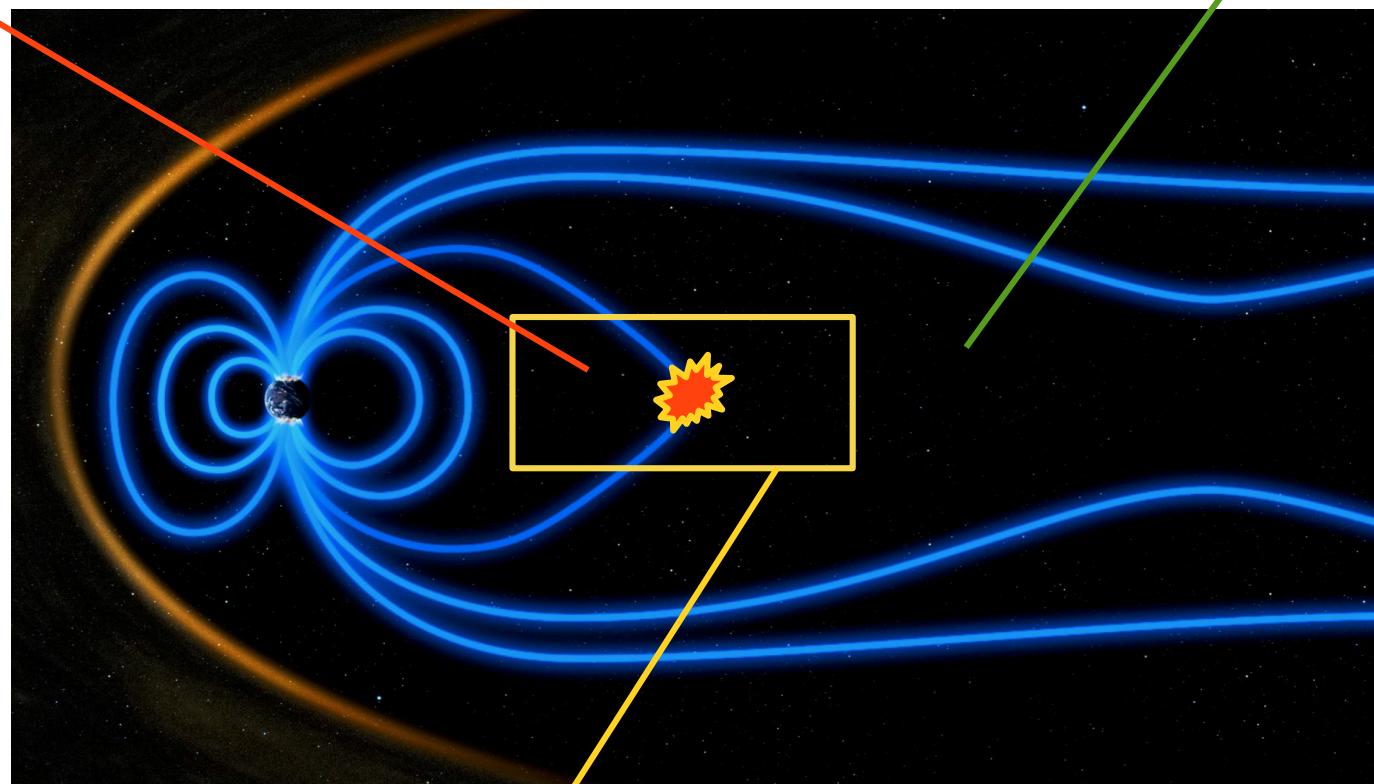
Simon Candelaresi, Michael Schlottke-Lakemper



Modeling

kinetic model
(expensive)

magnetohydrodynamics (MHD)
(cheap)

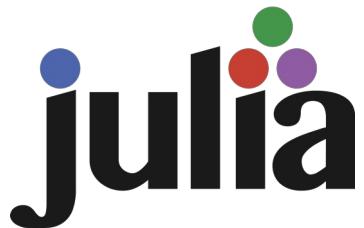


interface coupling

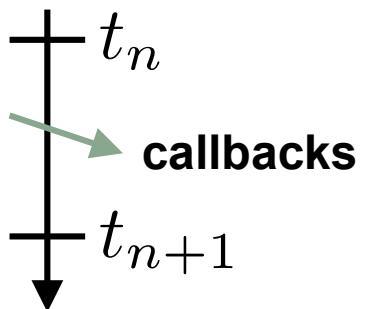
(ESA)



open source:
github.com/trixi-framework/Trixi.jl

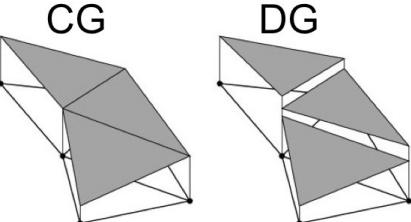


OrdinaryDiffEq (SciML)

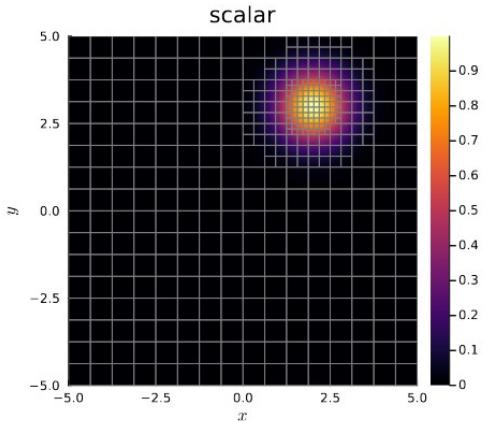


Trixi.jl

Discontinuous Galerkin



(Stack Overflow)



```
using Trixi

equations = LinearScalarAdvectionEquation2D((0.5, -0.3))
solver = DGSEM(polydeg = 3)
cells_per_dimension = (16, 16)

mesh = StructuredMesh(cells_per_dimension, (-1, -1), (1, 1))

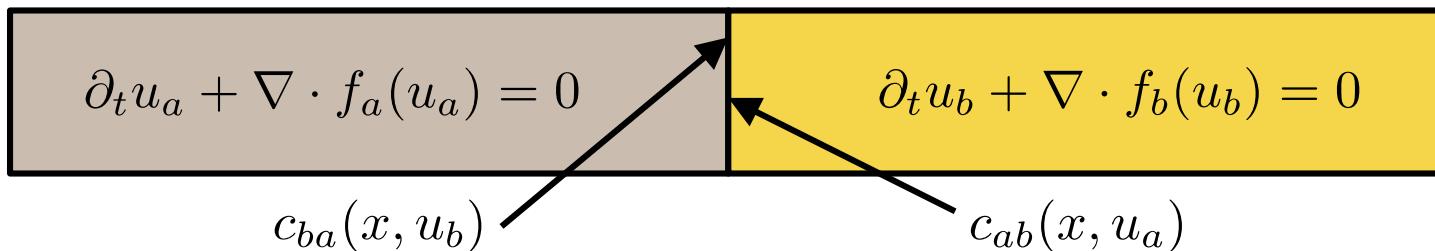
semi = SemidiscretizationHyperbolic(mesh, equations,
                                      initial_condition, solver)
ode = semidiscretize(semi, (0.0, 1.0));

stepsize_callback = StepsizeCallback(cfl = 1.6)
callbacks = CallbackSet(stepsize_callback)

sol = solve(ode, CarpenterKennedy2N54(),
            dt = 1.0, callback = callbacks);
```

Coupling via Converter Functions

Two system with any number of shared variables, including 0:

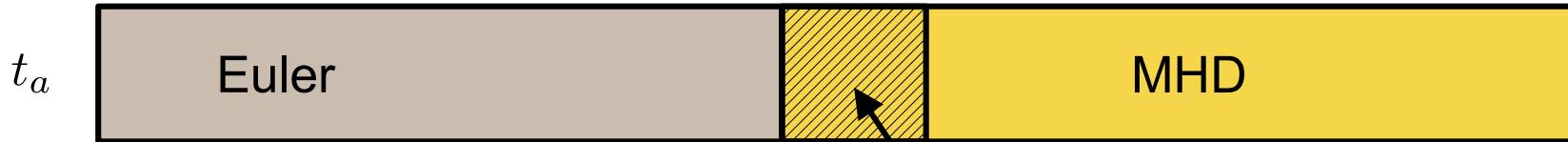


2.5d MHD $\rho, v_1, v_2, v_3, p, B_1, B_2, B_3, \Psi$	ρ, v_1, v_2, p	2d Euler
--	---------------------	----------

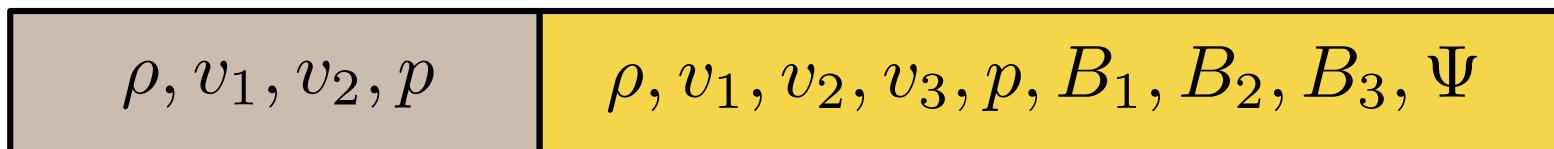
```
coupling_function12 = (x, u, equations_other, equations_own)
                      -> SVector(u[1], u[2], u[3], 0.0, u[4], 0.0, 0.0, 0.0, 0.0)
coupling_function21 = (x, u, equations_other, equations_own) -> SVector(u[1], u[2], u[3], u[5])
```

- ➔ User can define converter functions.
- ➔ Any pair of systems can be coupled.

Adaptive Coupling



strong currents and magnetic fields

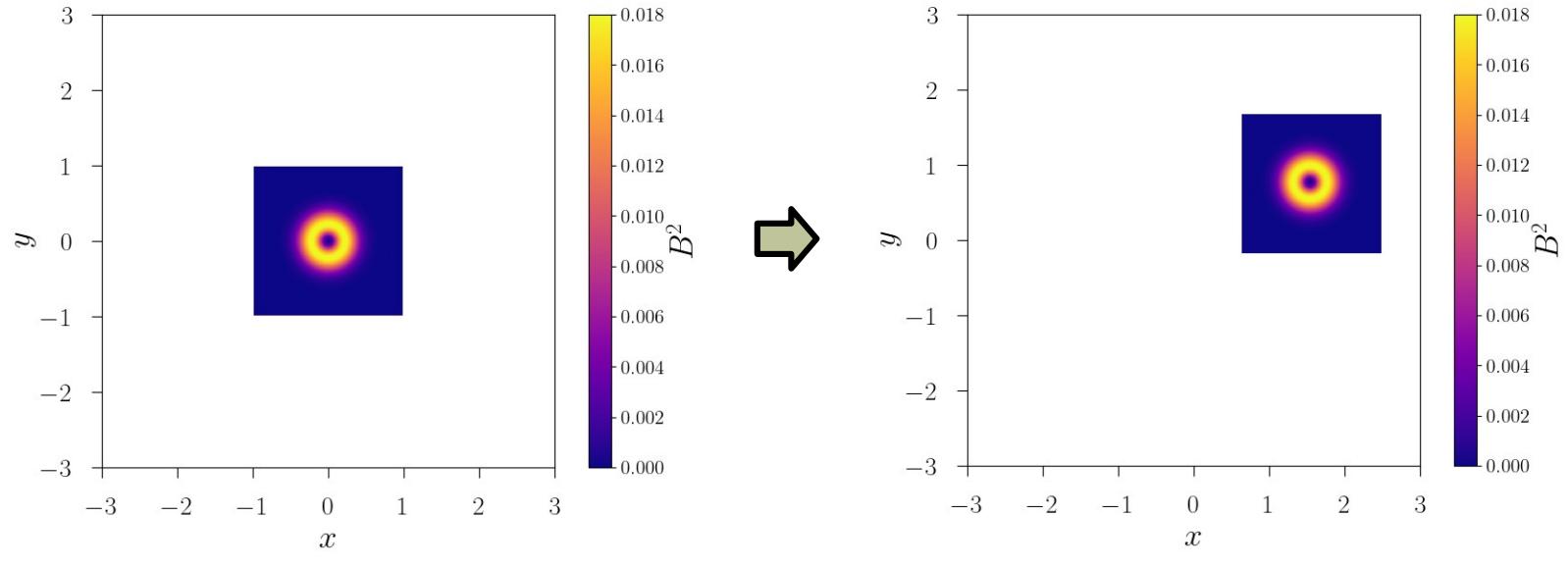


- ➡ Use callback functions to remesh.
- ➡ Use coupling functions to copy data.

Adaptive Model Selection

1. Generate the new grid (mesh views).
2. Write new u-solution vectors
3. Generate new ODE for OrdinaryDiffEq (integrator).
4. Reinitialize ODE integrator with new problem and new solution vector.

Euler and MHD



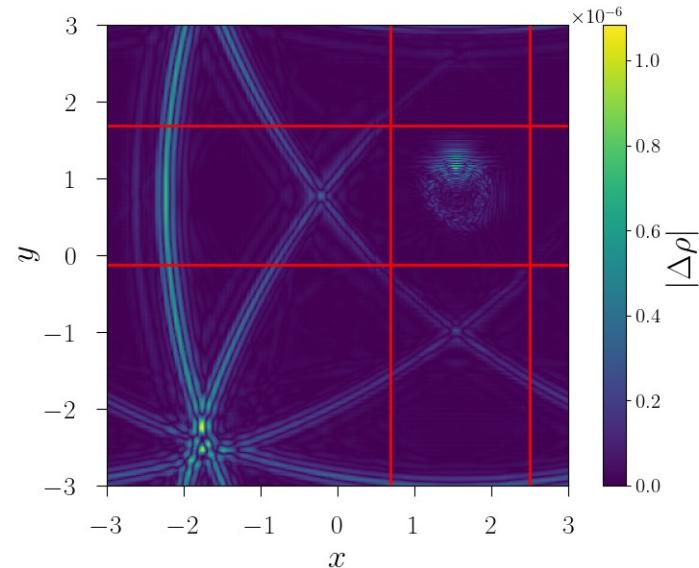
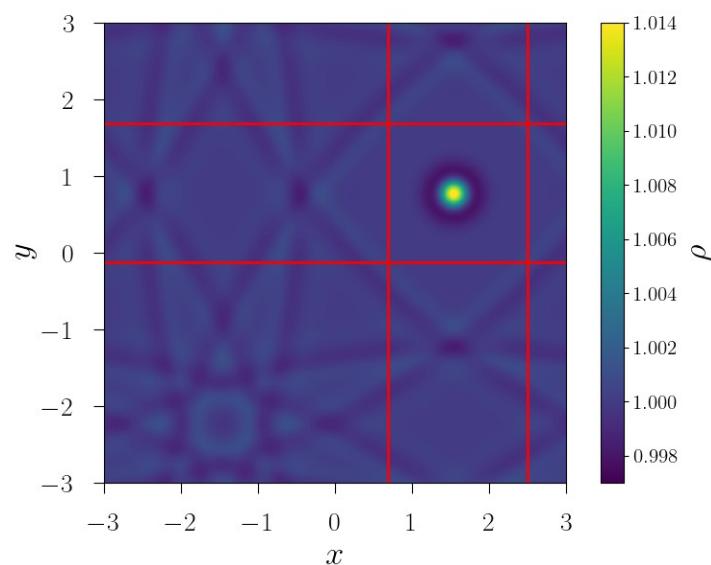
Time spent on coupled boundaries: 0.8%.

Time spent on model selection: 1.08s per model selection

Timings: coupled → 765.6s
full MHD → 1905.3s

Estimated maximum speed up: 3.31x; here: 2.49x

Density Evolution



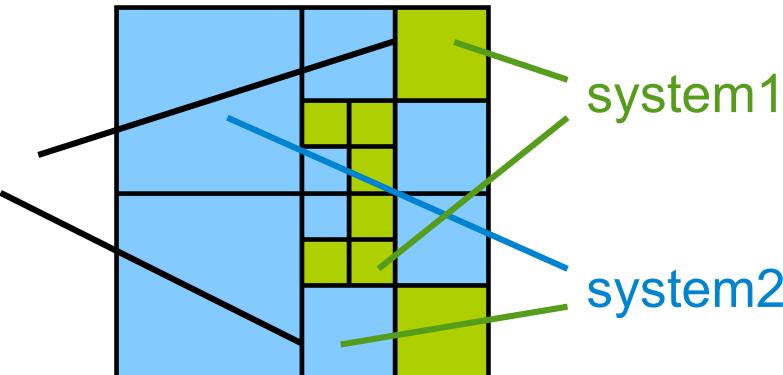
full MHD vs. coupled

→ No artificial discontinuities.

→ Small differences to full MHD simulation.

p4est Meshes

We can define the b.c. for each cell boundary.

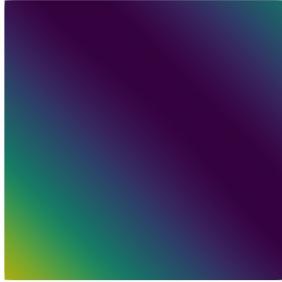
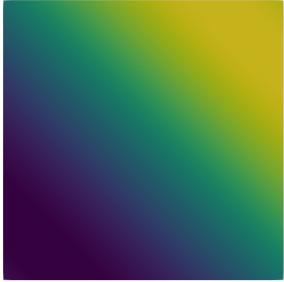


parent mesh

mesh views:



p4est Meshes



- p4est mesh views in Trixi.jl
- arbitrary topology
- views on smallest (cell) scale
- no coupling implemented yet

Conclusion

- ➡ Flexible coupling through converter functions.
- ➡ Free domain definitions.
- ➡ Adaptive coupling with arbitrary criteria.
- ➡ Coupled hierarchy of models.
- ➡ Non-simply connected mesh views in p4est