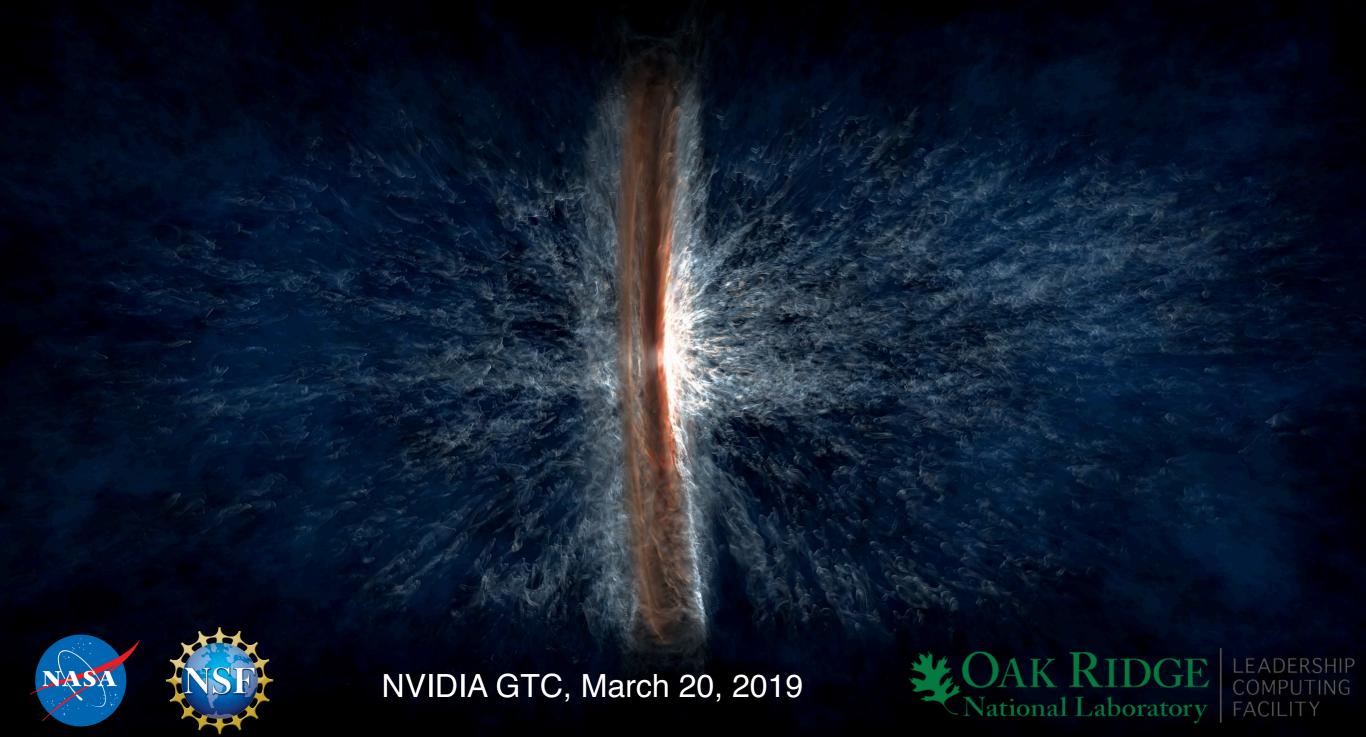
Advancing Astrophysics with the GPU-Native, Massively-Parallel Code, Cholla

Evan Schneider, Princeton University



Why did we need a new astrophysics code?



Starburst Galaxy M82 Image credit: NASA

Starburst Galaxy M82

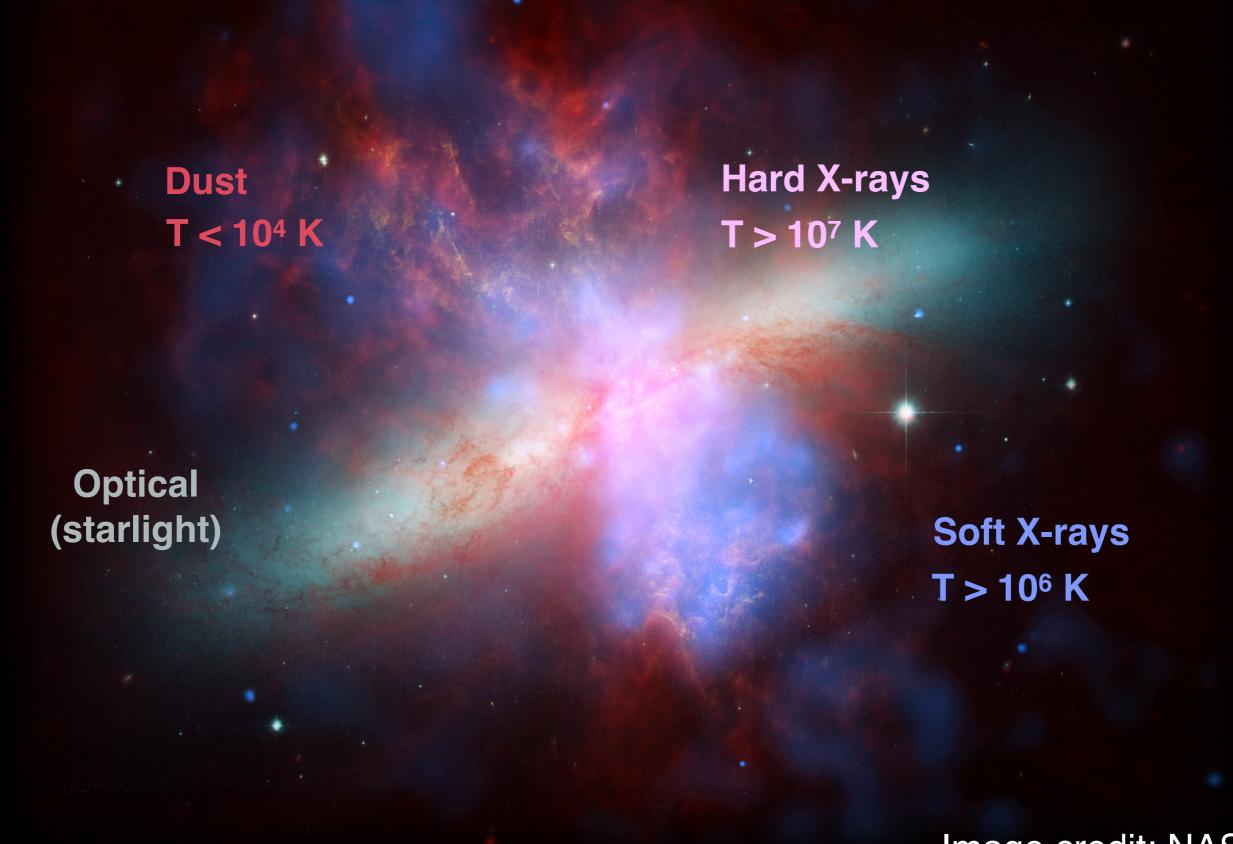


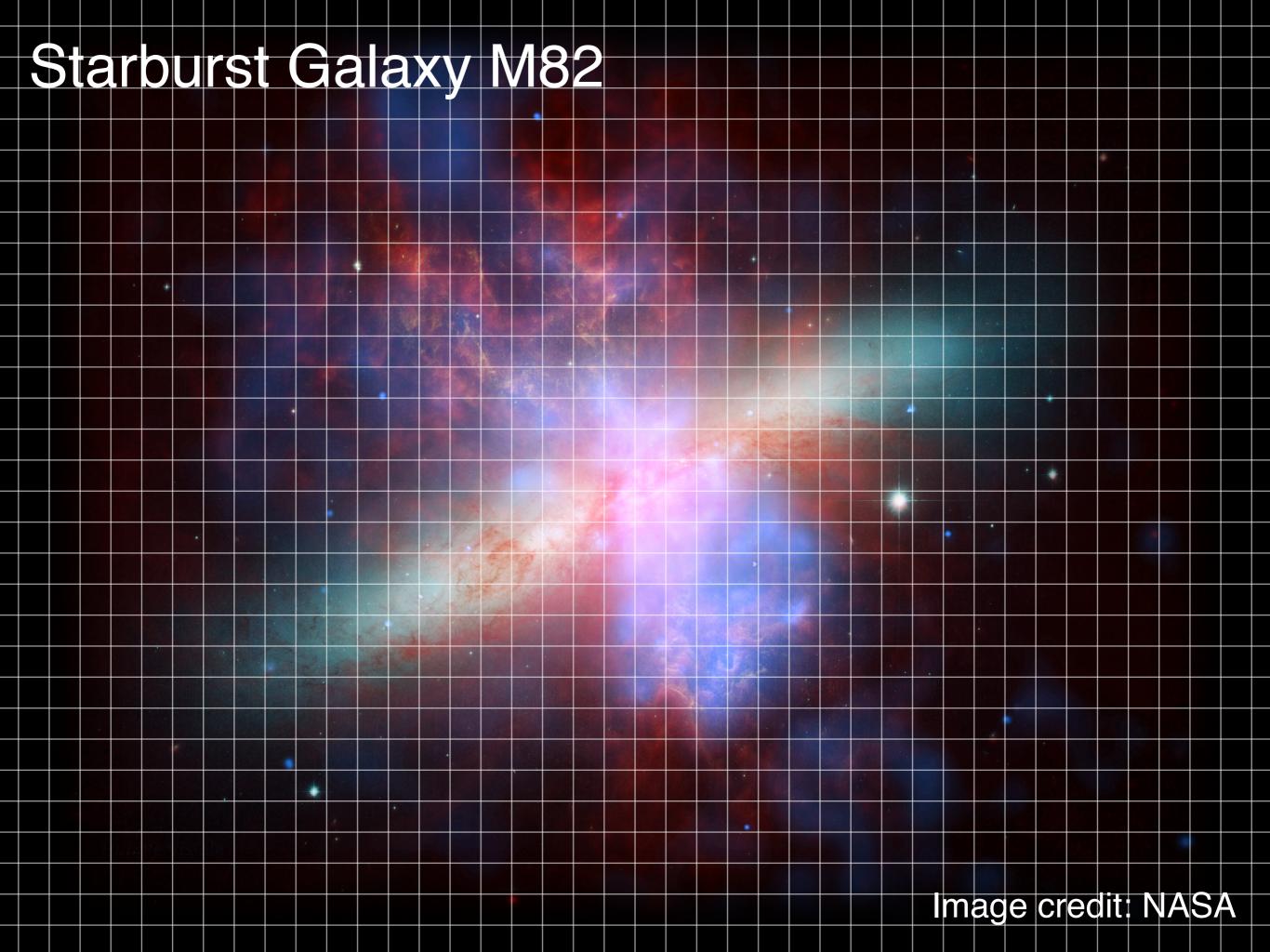
Image credit: NASA

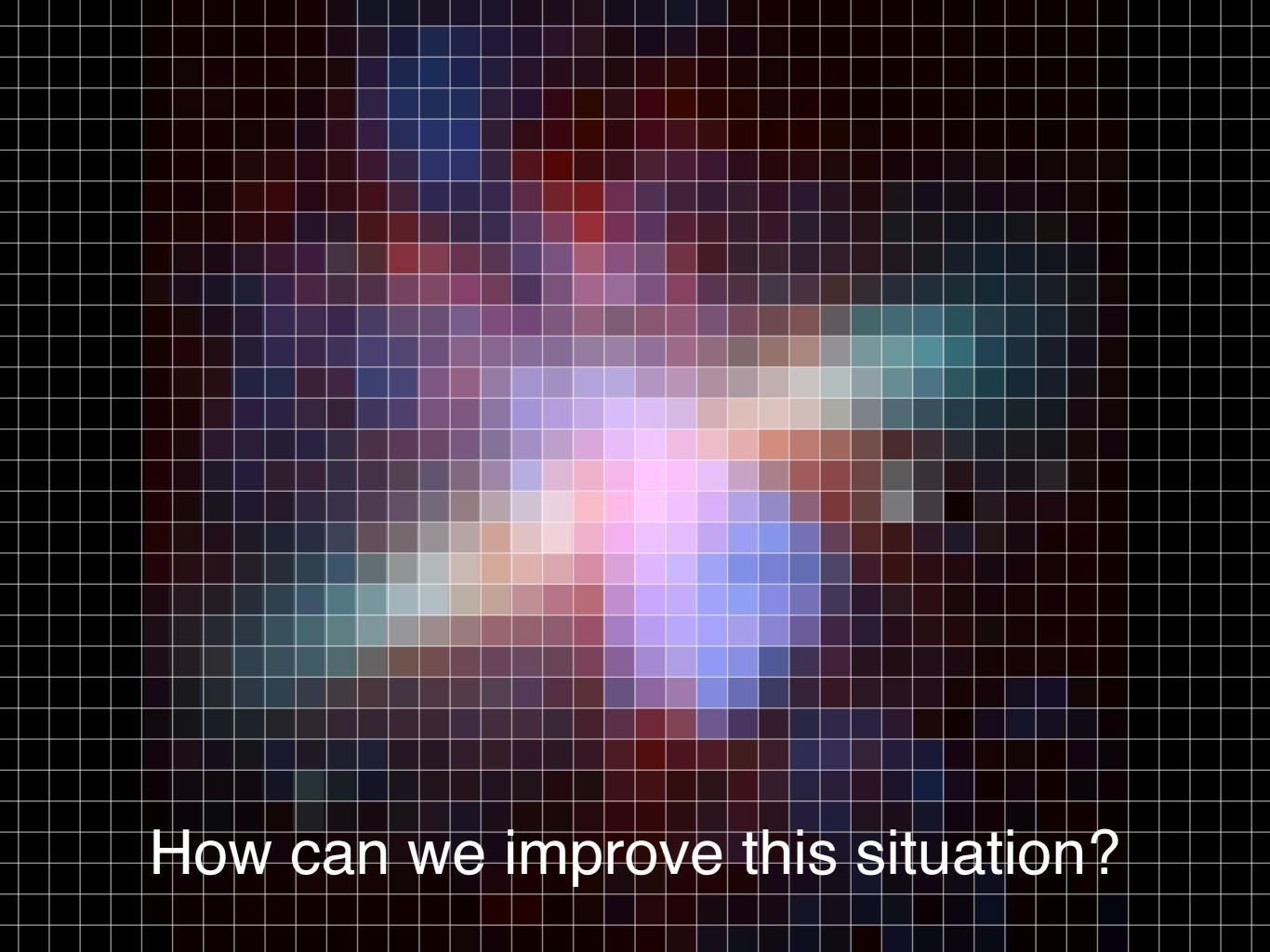
Simulating These Systems Is Computationally Challenging

~1K light-years ~10K light-years (winds generated) (galaxy disk)

~1M light-years (galaxy halo)

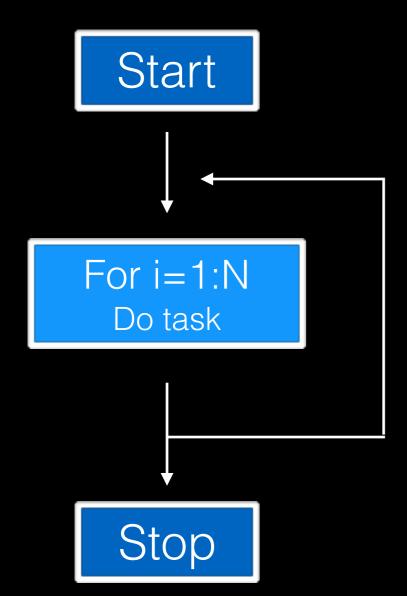
The scales involved in galactic wind evolution range from 10 light-years (cooling radius of supernova bubbles) to 1 million light years (extent of galaxy halo).



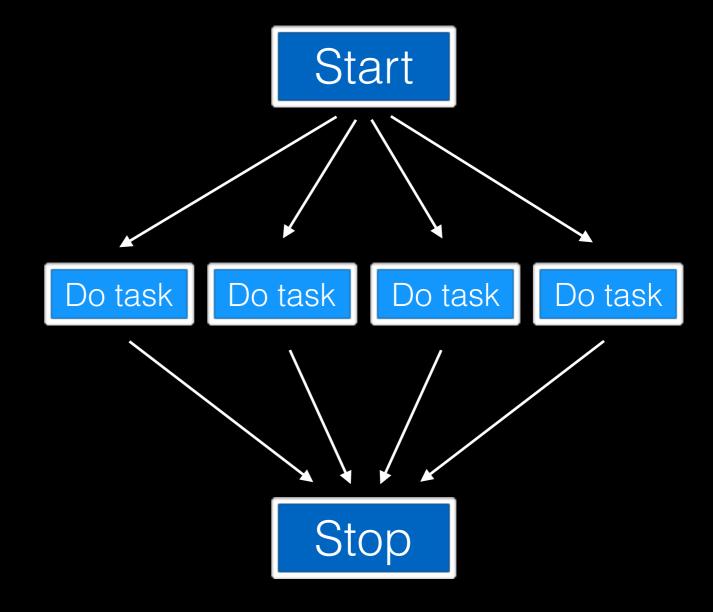


Computer Architectures Have Changed

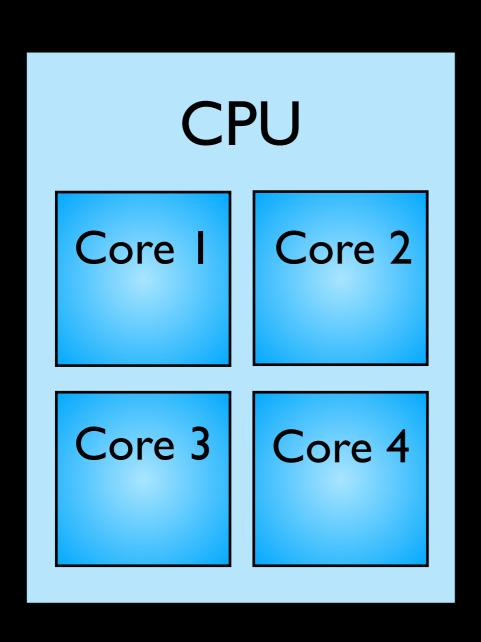
Serial Approach

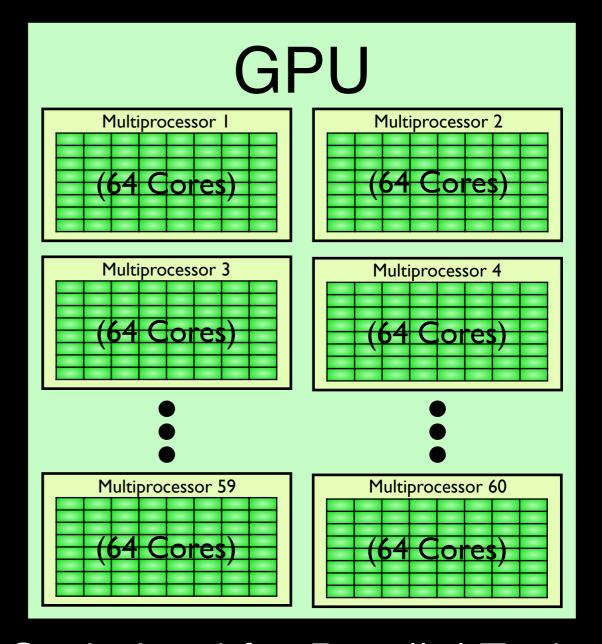


Parallel Approach



Computer Architectures Have Changed



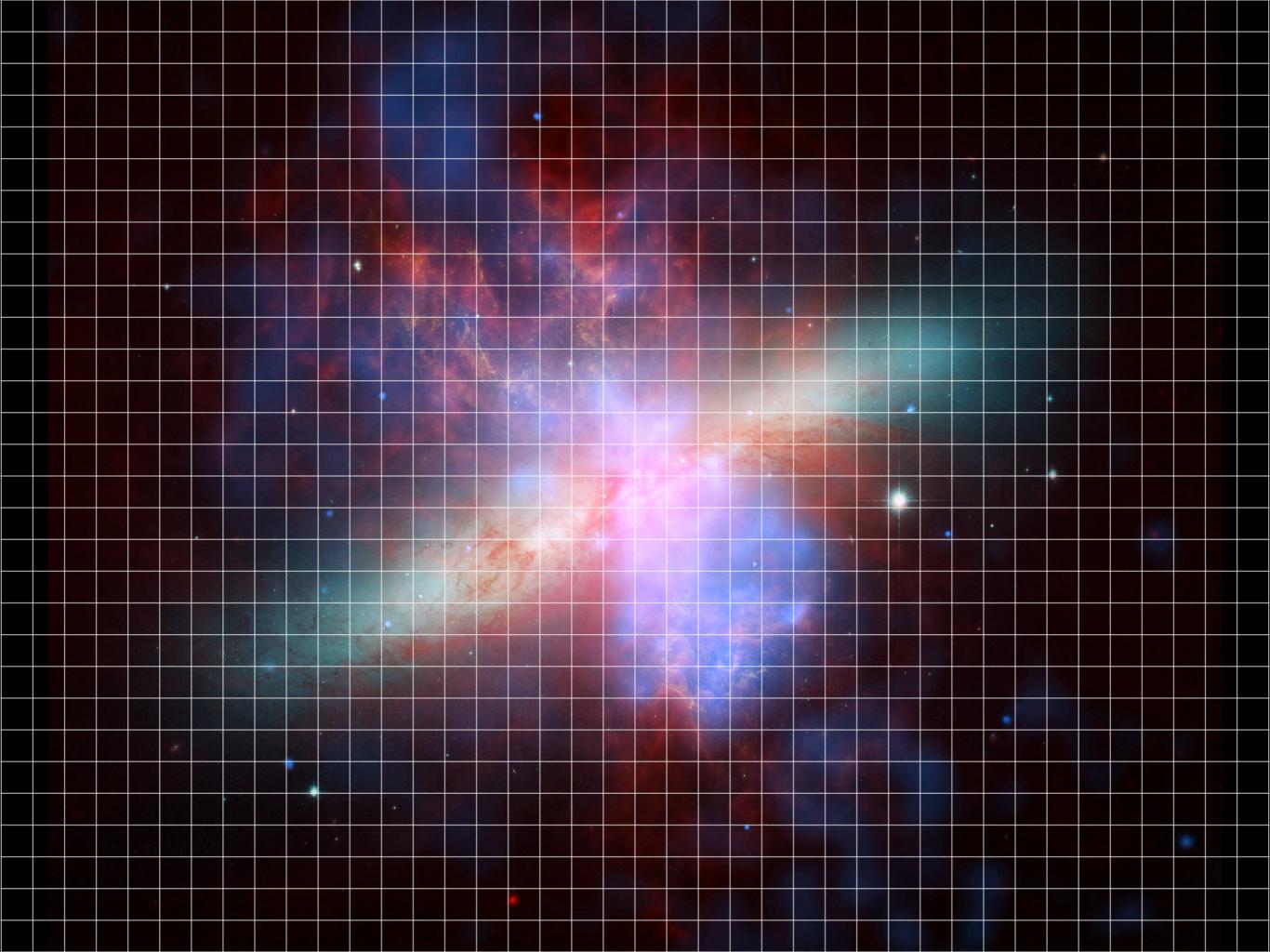


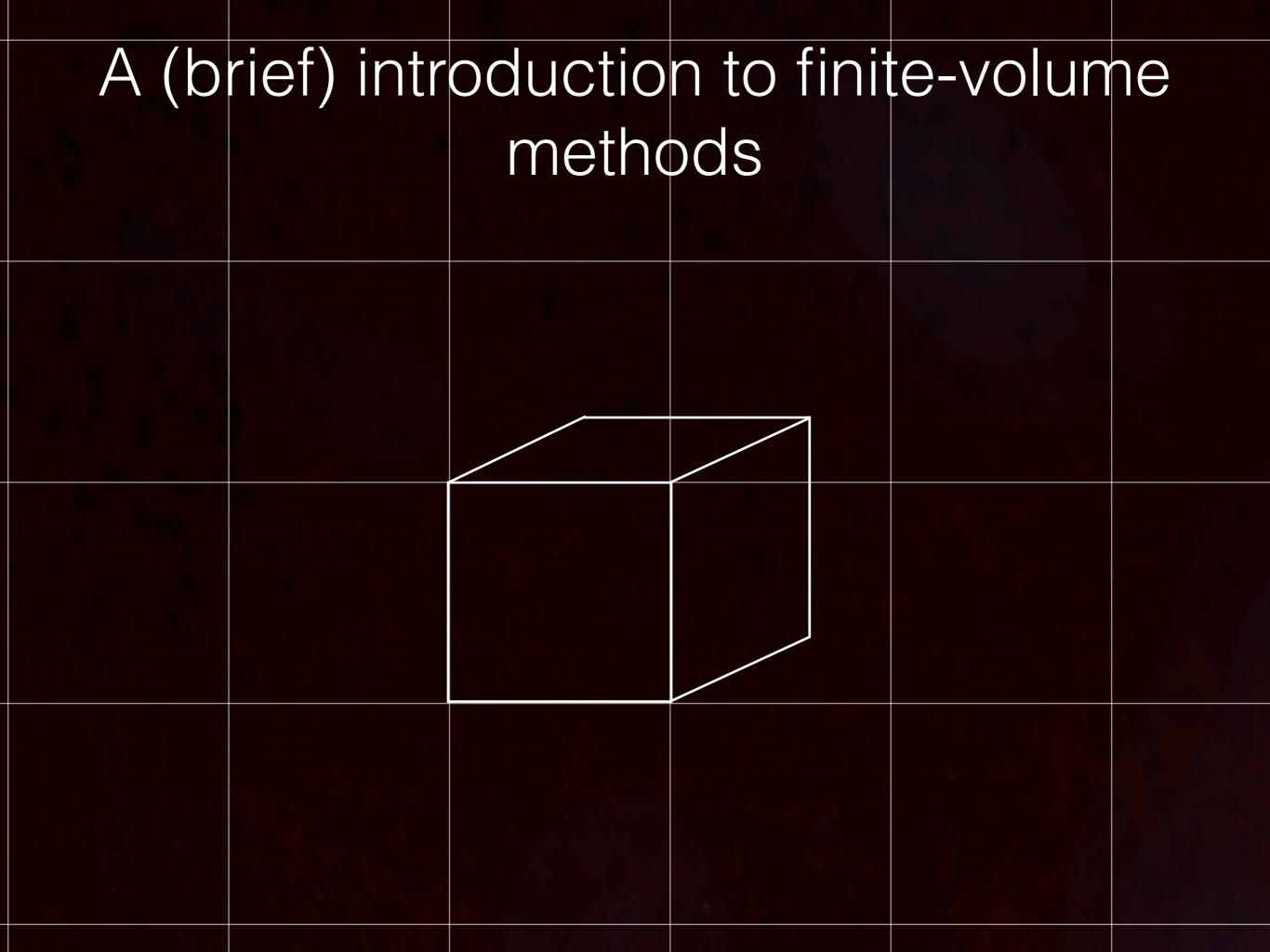
Optimized for Serial Tasks

Optimized for Parallel Tasks

So, the goal was to build a *new* code, that could:

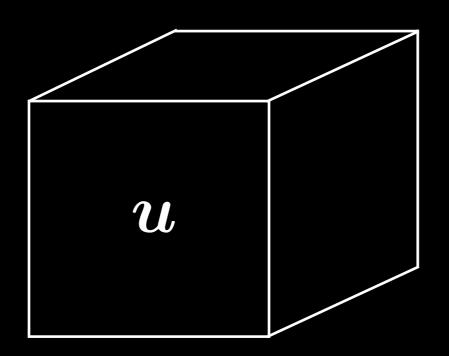
- achieve high resolution throughout the simulation volume (run simulations with large numbers of cells)
- take full advantage of new computing architectures
- address the limitations of the previous generation of astrophysics codes.





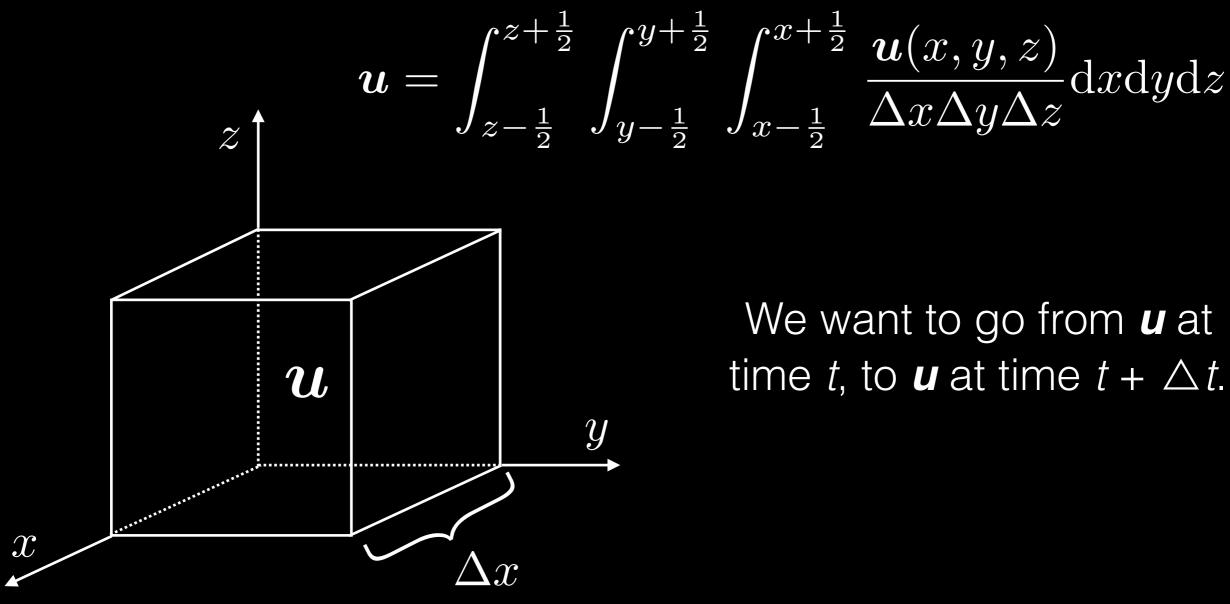
A (brief) introduction to finite-volume methods

 $m{u} = [
ho,
ho u,
ho v,
ho w, E]^{\mathrm{T}}$ a vector of conserved quantities



A (brief) introduction to finite-volume methods

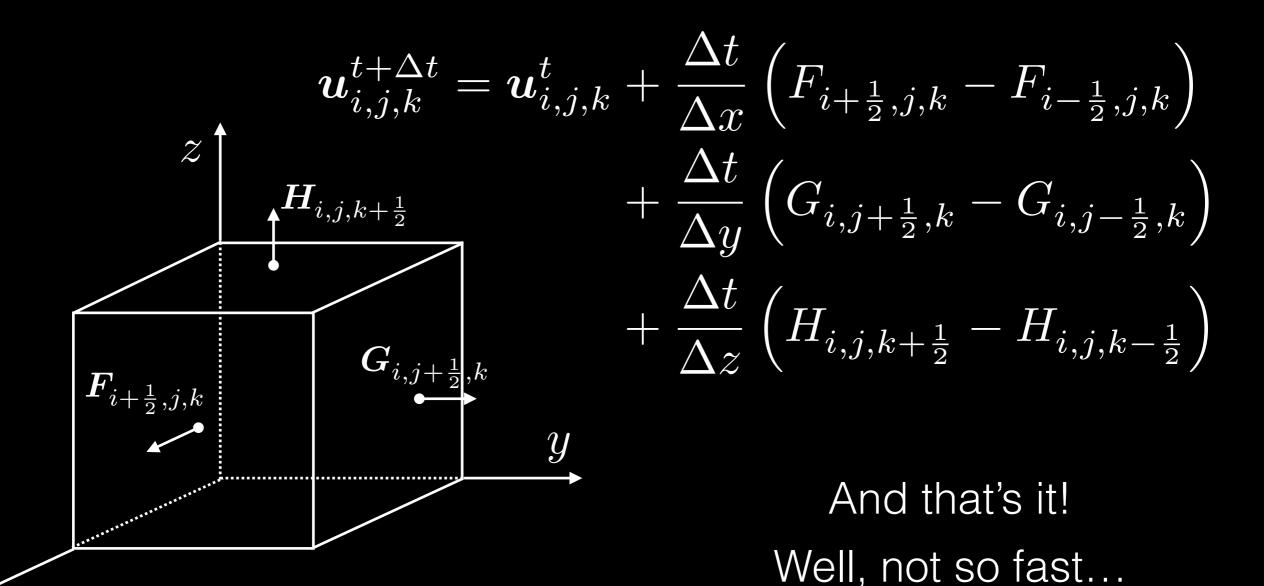
 $\mathbf{u} = [\rho, \rho u, \rho v, \rho w, E]^{\mathrm{T}}$ a vector of conserved quantities



We want to go from **u** at time t, to \boldsymbol{u} at time $t + \triangle t$.

A (brief) introduction to finite-volume methods

 $\boldsymbol{u} = [\rho, \rho u, \rho v, \rho w, E]^{\mathrm{T}}$ a vector of conserved quantities



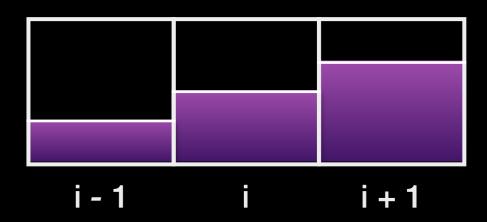
 \mathcal{X}

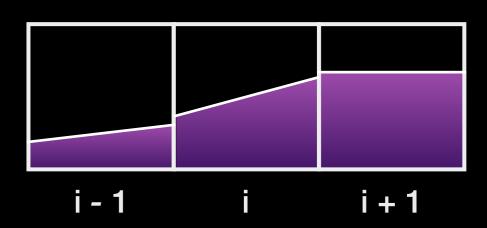
How do we do it?

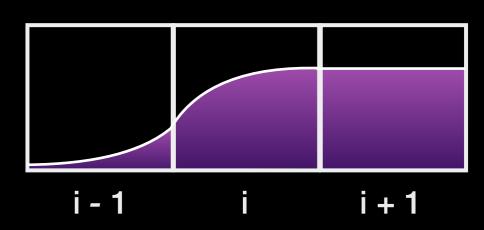
Reconstruct interface values using cell averages.

Choose either piecewise constant, piecewise linear, or piecewise parabolic reconstruction.

Piecewise linear and piecewise parabolic reconstruction can be done in either the primitive variables or the characteristic variables.

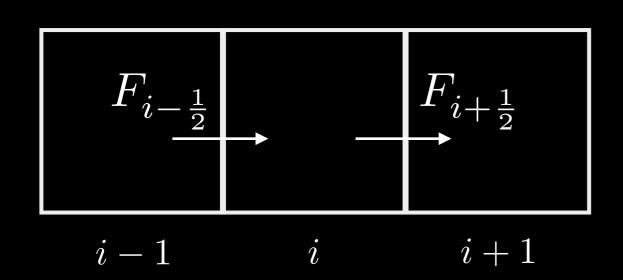


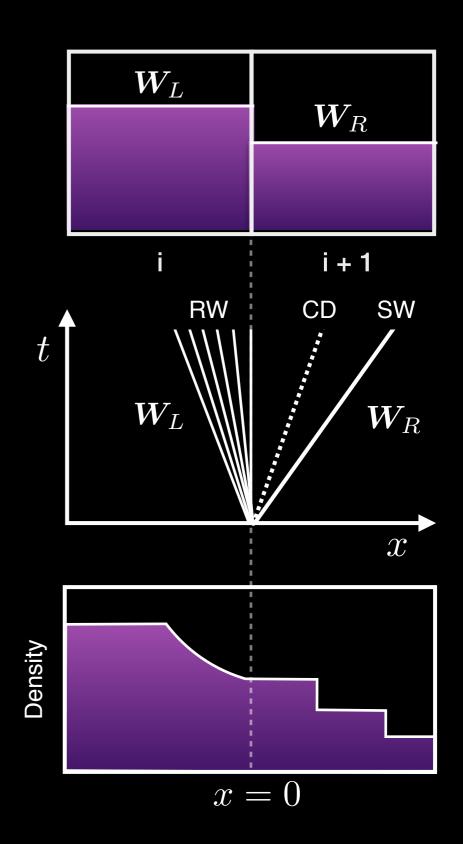




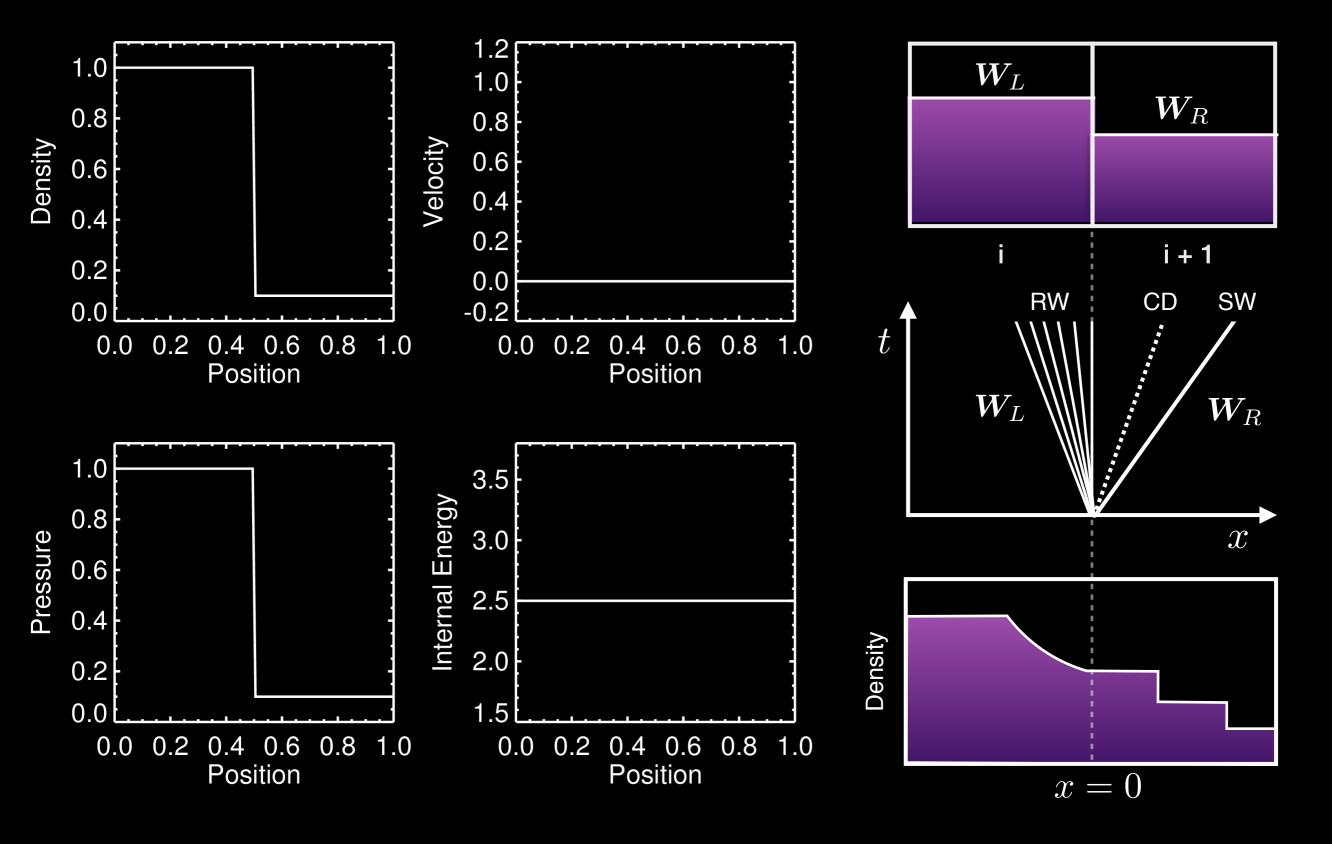
How do we do it?

Calculate fluxes across cell interfaces using reconstructed interface values.



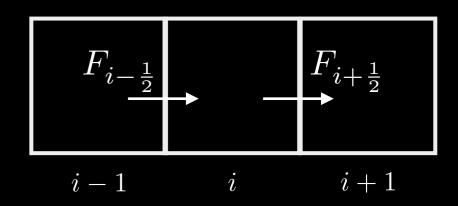


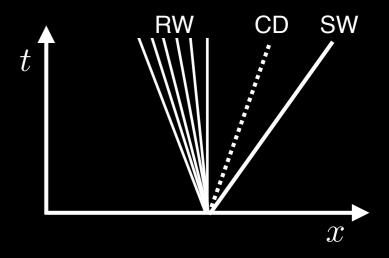
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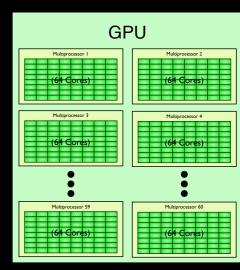


What's the GPU advantage?

- Grid-based hydro codes are eminently parallelizable - each cell needs data from only a few nearby cells to reconstruct interface values, calculate fluxes across interfaces, and update conserved quantities.
- Hydro solvers are computationally expensive. Many unsplit algorithms require 6 Riemann problems per cell, per timestep (in 3D).
- With GPUs, we massively parallelize the calculation across many cores, allowing us to speed up computation by an order of magnitude as compared to similarly intensive CPU codes.

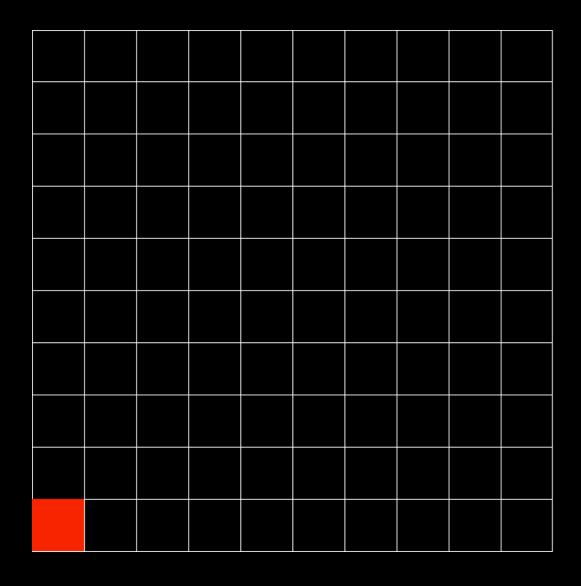




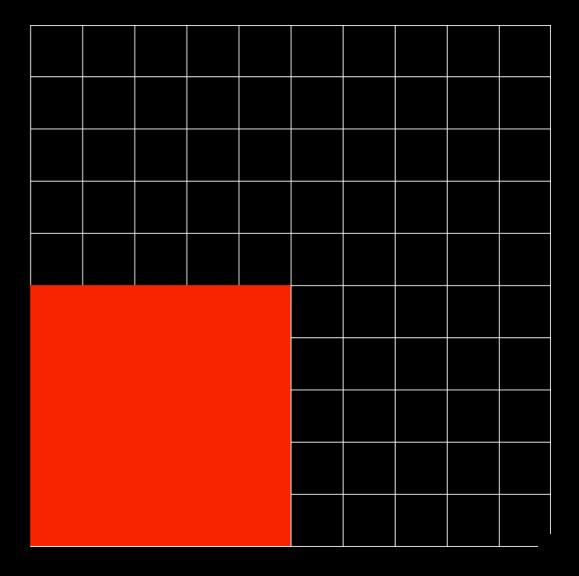


What's the GPU advantage?

Grid code on a CPU

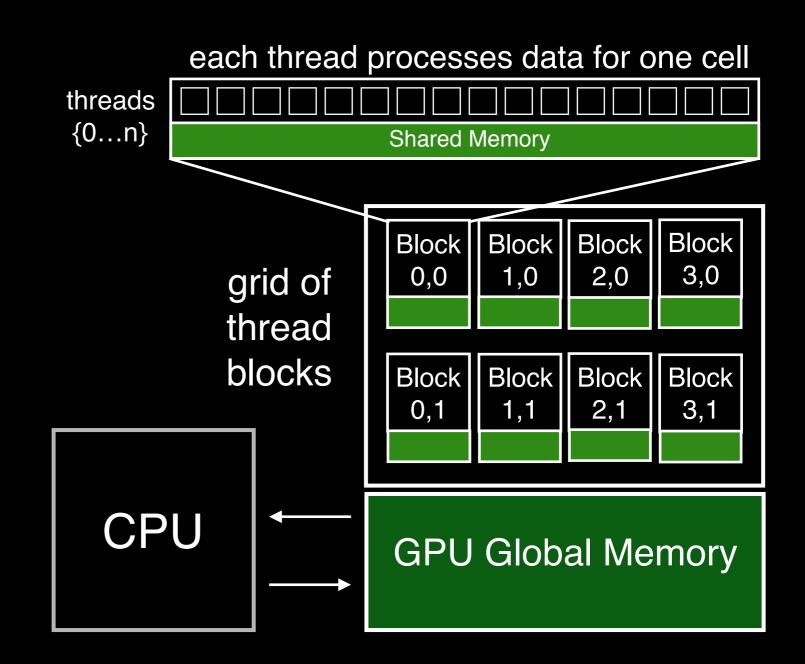


Grid code on a GPU



How does it work?

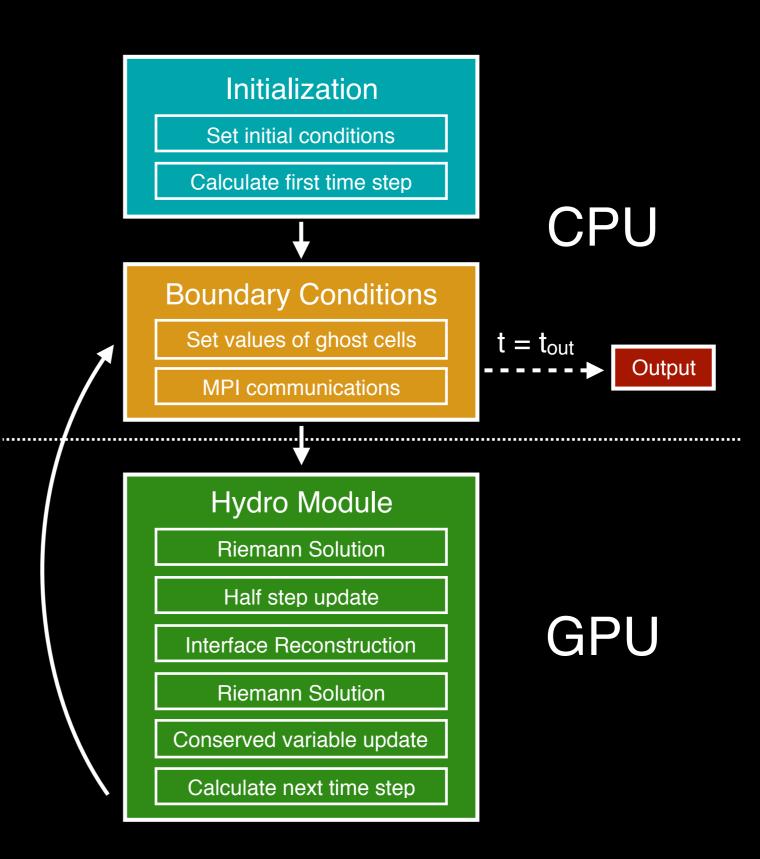
- GPU functions execute as CUDA kernels on a grid of thread blocks each cell in the simulation is mapped to a single thread.
- Our design minimizes memory transfers between the CPU and GPU, reducing computational overhead.



How does it work?

Serial parts of code execute on the CPU

Parallel portions execute on the GPU

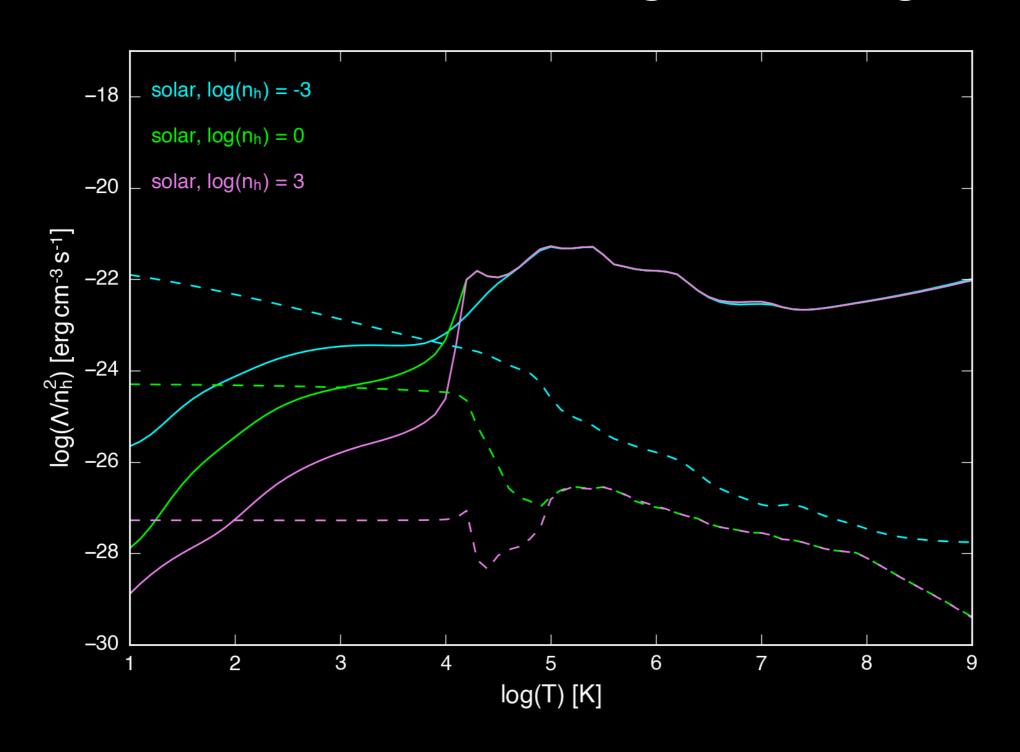


An aside: using texture mapping to accelerate cooling calculations.

What is texture mapping?



Many astrophysical cooling calculations rely on multidimensional table lookups to calculate radiative cooling / heating rates.



Texture mapping is a natural fit to the cooling implementation.

- 1. Copy 2D cooling tables to texture memory on the GPU
- 2. Calculate density and temperature of gas
- 3. "Fetch" cooling and heating rates from the texture bilinear interpolation comes for free!

How does it compare to the CPU version?

Loops/ Threads	GSL (CPU)	CUDA (GPU)
102	0.006 ms	0.023 ms
104	0.31 ms	0.023 ms
106	30 ms	0.023 ms

So basically, texture mapping makes our cooling calculations free.

Cholla: Computational hydrodynamics on

architectures



Cholla are also a group of cactus species that grows in the Sonoran Desert of southern Arizona.

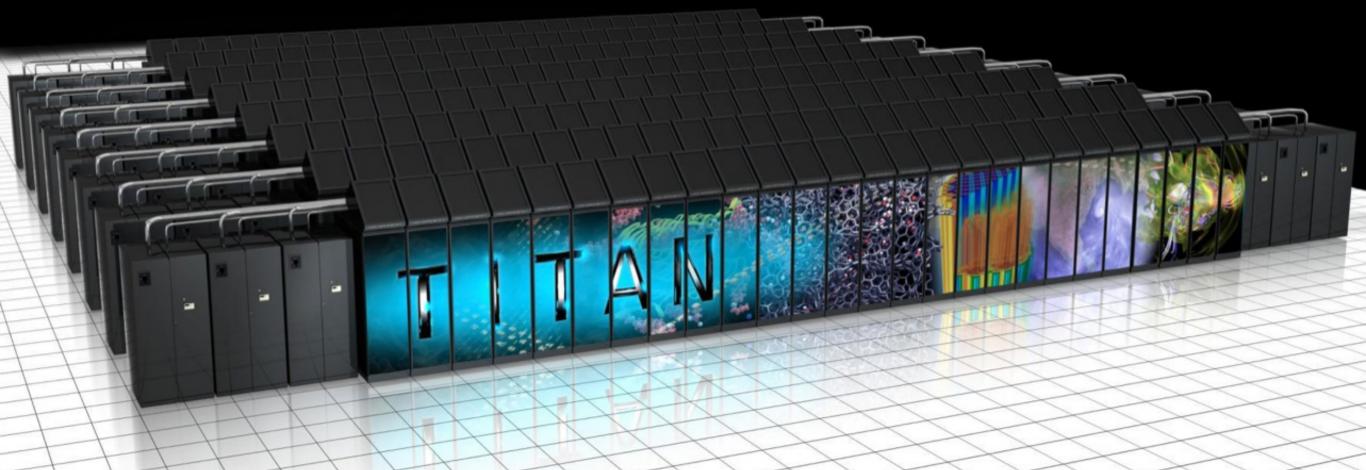
- A GPU-native, massivelyparallel, grid-based hydrodynamics code (publicly available at github.com/chollahydro/cholla)
- Incorporates state-of-the-art hydrodynamics algorithms (unsplit integrators, 3rd order spatial reconstruction, precise Riemann solvers, dual energy formulation, etc.)
- Also includes optically-thin cooling and photoionization heating based on precomputed rate tables, and static gravity.

Schneider & Robertson (2015, 2017)

Cholla takes advantage of the world's most powerful supercomputers.

Titan: Fastest Open Science Supercomputer in the US

Flagship accelerated computing system | 200-cabinet Cray XK7 supercomputer | 18,688 nodes (AMD 16-core Opteron + NVIDIA Tesla K20 GPU) | CPUs/GPUs working together – GPU accelerates | 20+ Petaflops



Cholla takes advantage of the world's most powerful supercomputers.

Summit:

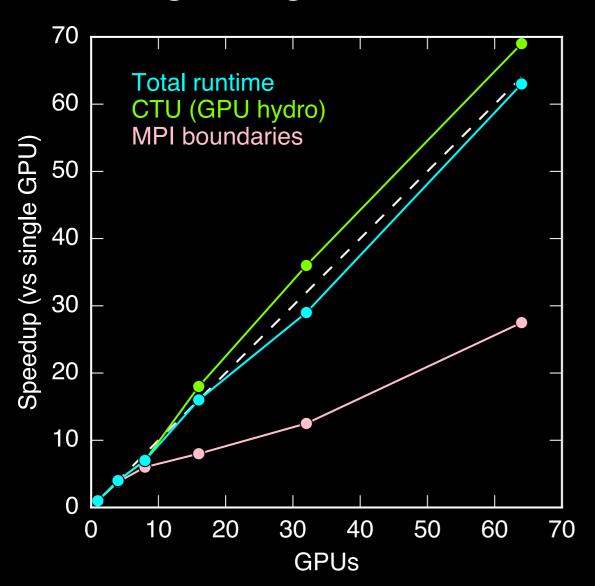
Fastest Open
Science
Supercomputer
in the World

- 200 petaflops
- 4600 nodes
- 27,600 NVIDIA
 V100 GPUs

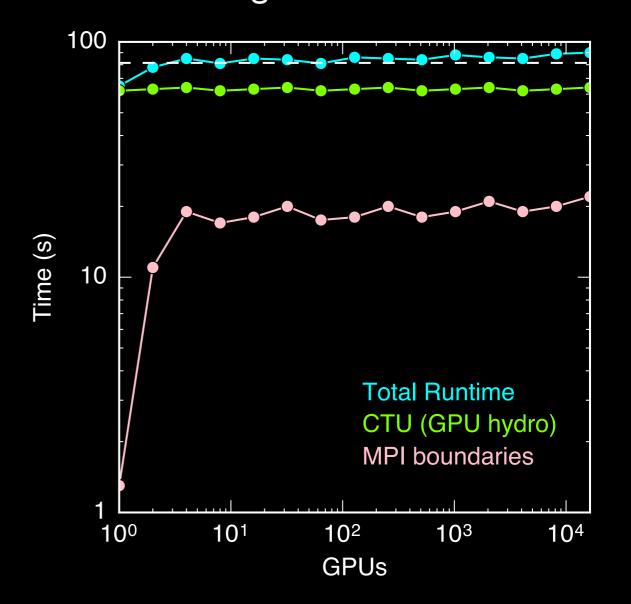


Cholla Achieves Excellent Scaling





Weak Scaling test, ~3223 cells / GPU



Schneider & Robertson (2015)

<u>Advantages</u>

- Optimized for fast execution of parallel tasks
- Blocked architecture easily transitions to new hardware models
- Specialized hardware functions are FAST
- Offloading computation to GPU leaves CPU free to perform other tasks
- Energy efficient!

Challenges

- Limited memory on GPU
- Need lots of computation to make up for data transfer and memory latency
- Blocked architecture not optimal for some problems

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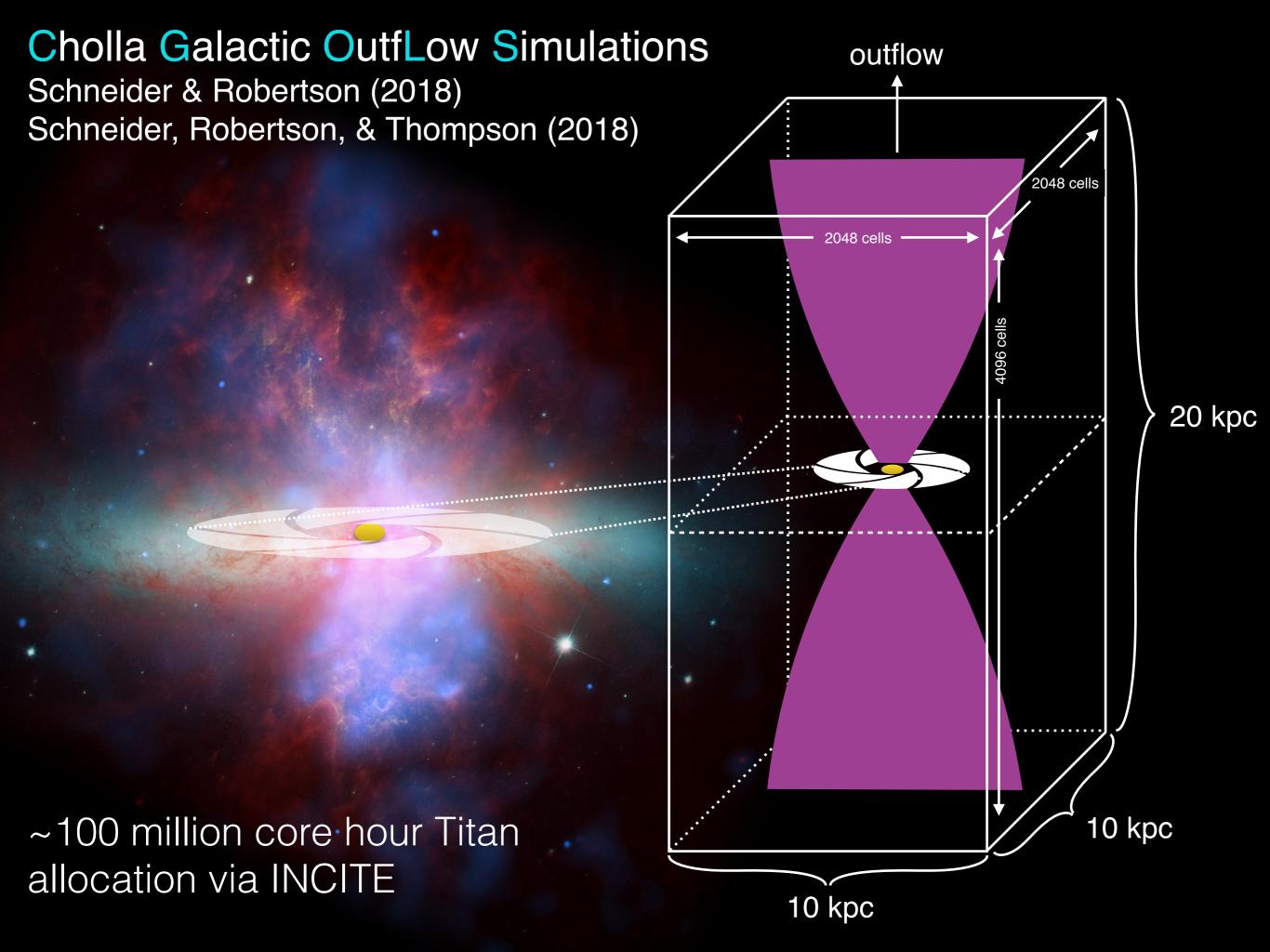
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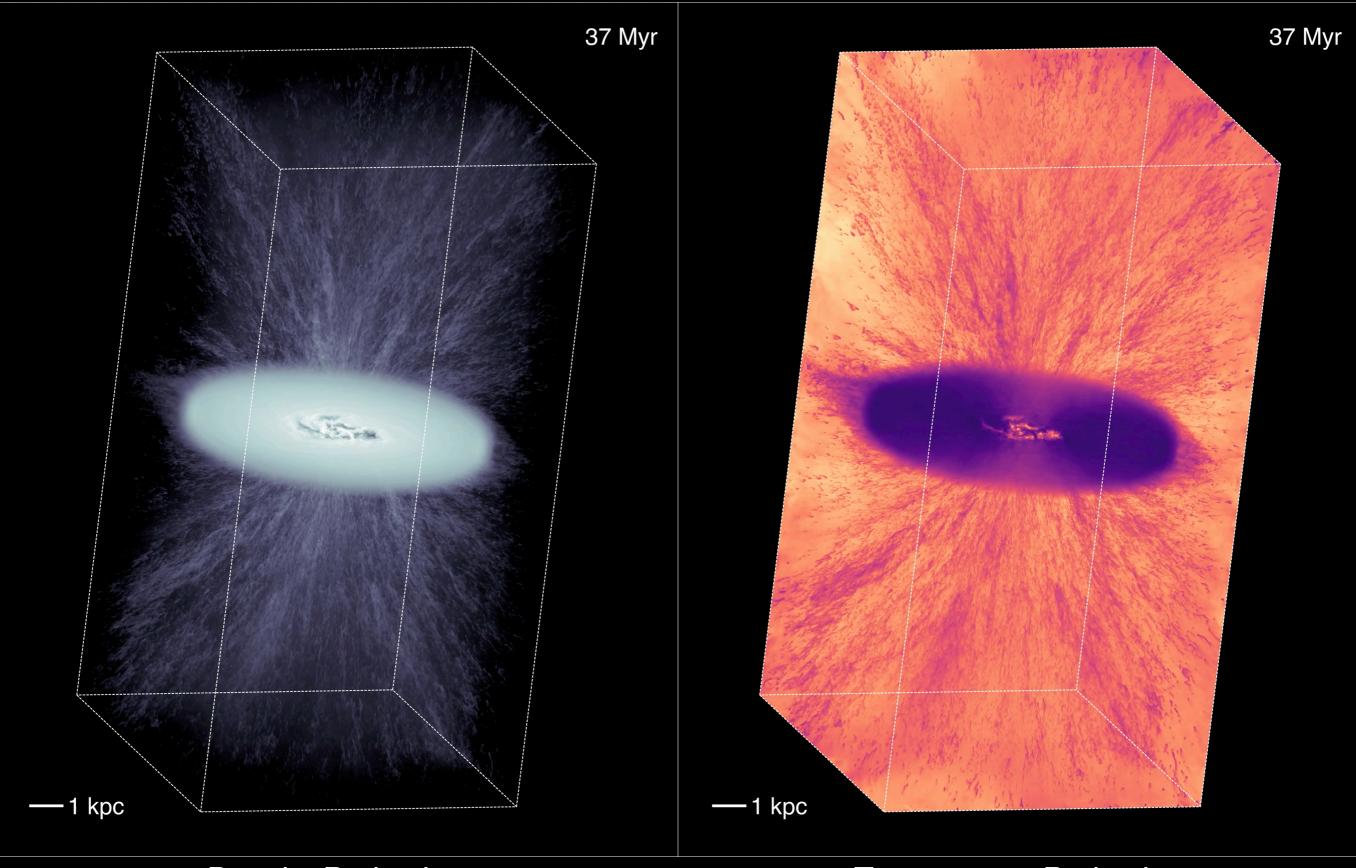
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Starburst Galaxy M82 Image credit: NASA



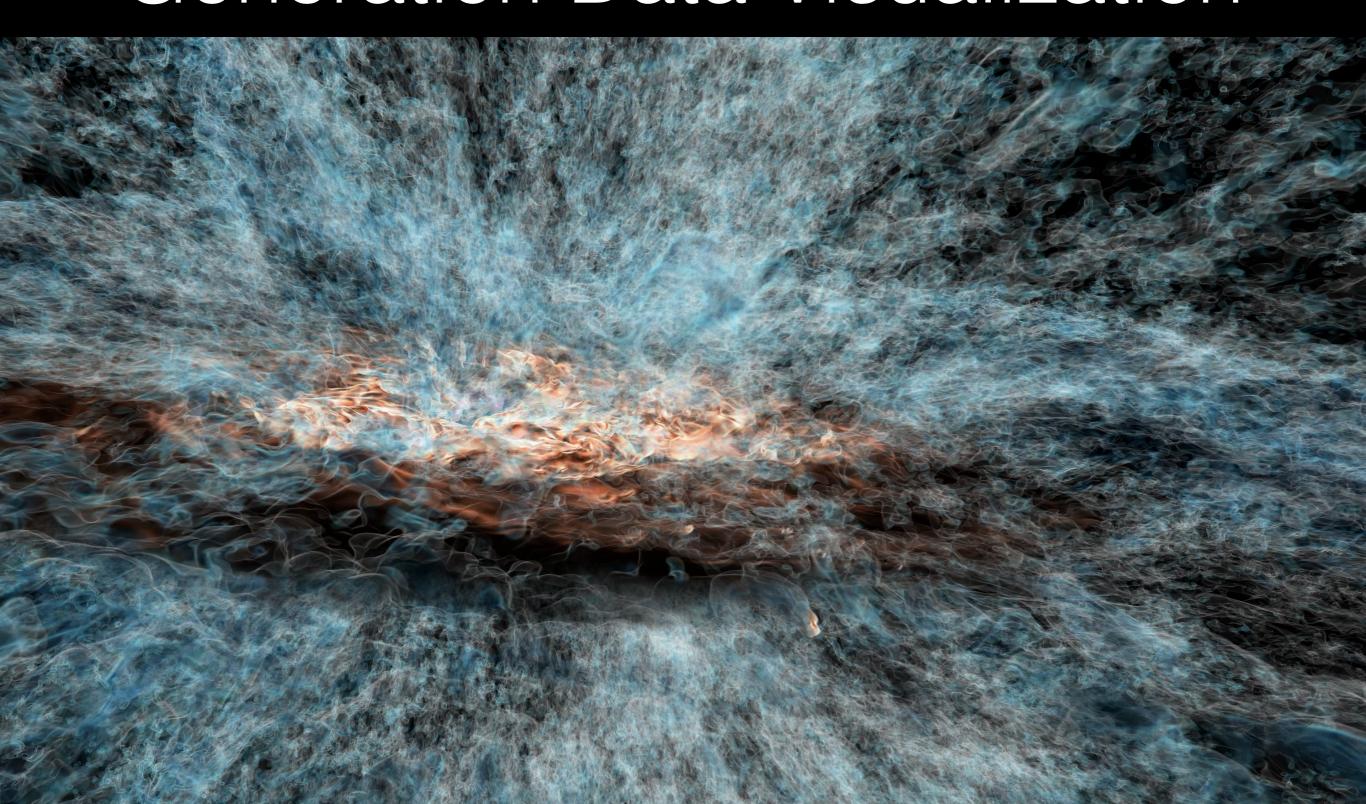
CGOLS Simulation - 17 billion cells



Density Projection

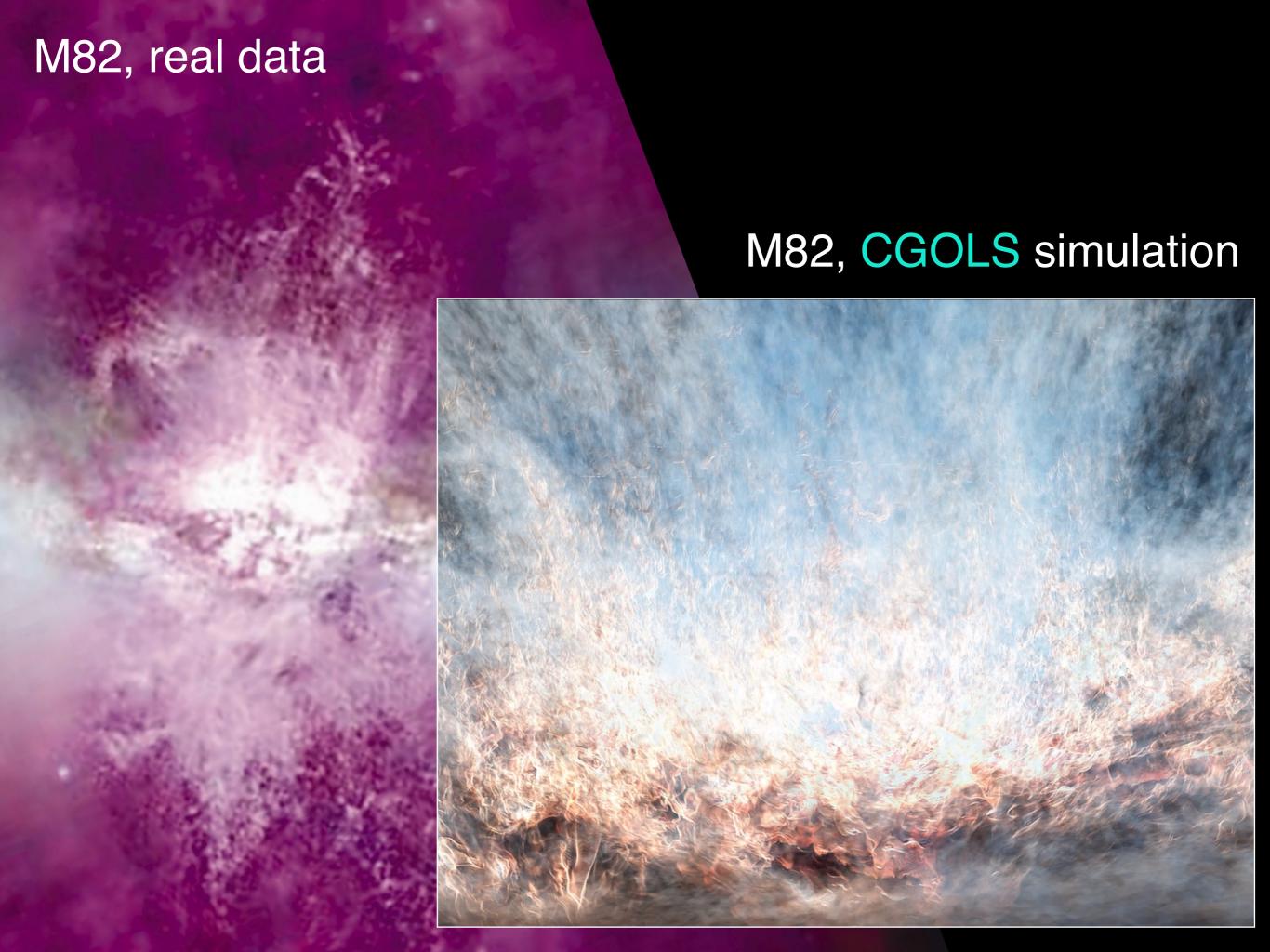
Temperature Projection

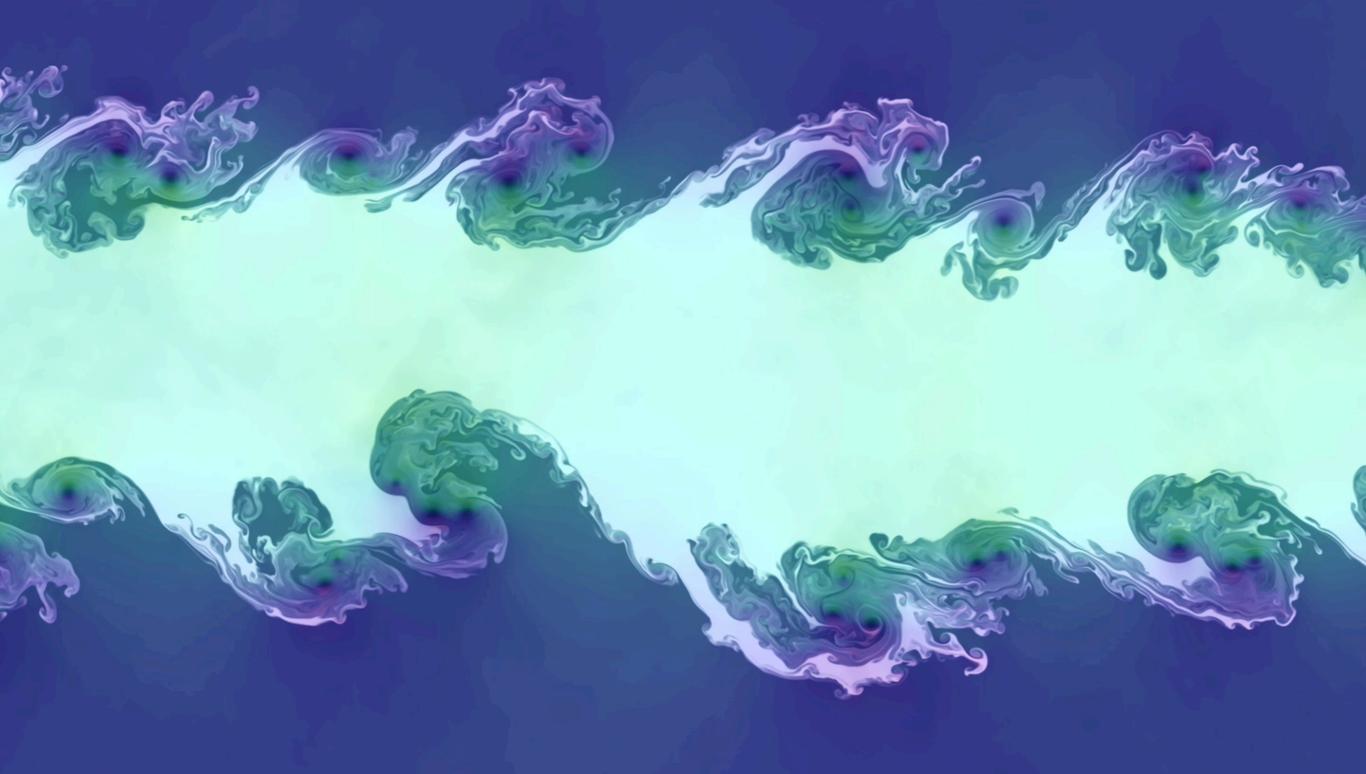
Cholla plus IndeX - Next Generation Data Visualization



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Thanks!