

# **Visualization on Mistral**

Workflow, applications, future plans

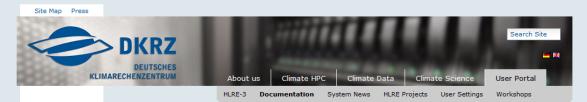
Niklas Röber Deutsches Klimarechenzentrum (DKRZ)



### Hardware

- 12(24) nodes with 2x12 core Haswell CPUs
- 256 GB main memory
- Two Kepler GPUs
- Regular mistral nodes with XAS software tree:
  - avizo/9.0.1
  - paraview/4.3.1
  - simvis/3.4.4
  - vapor/2.4
  - idl/8.5

- ncl/6.2.1
- grads/2.0.2
- ferret/6.9.3
- gmt/5.1.2



#### Documentation

How to get a user account

Mistral

Blizzard

Wizard

HPSS tape archive

Data Processing Visualization

Software

Visualization on Mistral

Remote3D

Filesystems

Cloud Storage

Training

FAQs & known issues

Seminar Rooms

Terms of use

News

Kick-off for ESIWACE and ESCAPE

Sep 29, 2015
Allocations 2016

- request resources Sep 18, 2015

DKRZ@ISC'15 Jul 13, 2015

DKK publishes "Prospects for climate research"

Jun 10, 2015

More news.

 $\mathsf{Home} \to \mathsf{User} \; \mathsf{Portal} \to \mathsf{Documentation} \to \mathsf{Visualization} \to \mathsf{Visualization} \; \mathsf{on} \; \mathsf{Mistral}$ 

#### **Visualization on Mistral**

Our new supercomputer Mistral has 12 GPU nodes, which can be used for 3D visualization, data analysis, and pre/post processing of data. This website explains how to reserve and access a GPU node, and how to run the 3D visualization software for the analysis of your data.

#### Access and reservation of a GPU node

Our supercomputer Mistral includes 12 GPU nodes with four GPUs in each node. Unlike the nodes in our visualization cluster Halo, these nodes are equipped with the same hard and software.

While we might later switch back to our web-based reservation system, currently the reservation has to be done by hand using a console on one of Mistrals login nodes. Furthermore, the vncserver has to be started by hand, as well as cleaned up after your work is done. To access Mistral, simply ssh into the machine.

```
somewhere: ~> ssh user_name@mistral.dkrz.de
```

On Mistral, we have to allocate a GPU node. This is done using the SLURM command "salloc", in which you have to provide your account group ("-A <your project id>"), the number of nodes ("-N 1"), the maximum number of parallel tasks ("-n 24"), as well as the node type ("-p gpu"). Currently, the maximum time allowed is 4 hours and is set automatically. More information on SLURM and salloc can be found in our SLURM documentation.

After login to Mistral, you can allocate a GPU node and automatically ssh into the node reserved.

```
mistral:-> salloc -N 1 -n 24 -p gpu -A project> -- /bin/bash -c 'ssh -X $SLURM_JOB_NODELIST'
salloc: Granted job allocation 284896
user_name@mg100's password:
```

Now you can start a VNC server to connect to the virtual desktop.

```
mg100:-> /opt/TurboVNC/bin/vncserver -geometry 1920x1200
Desktop 'TurboVNC: mg100:1 (user_name)' started on display mg100:1
```

At the first time, you have to supply a password that you later need to access your VNC session remotely. If everything is set, start a vncviewer in the console on your home computer, or using the gui of an application.

```
set, start a vncviewer in the console on your home computer, or using the gui of an application.
```

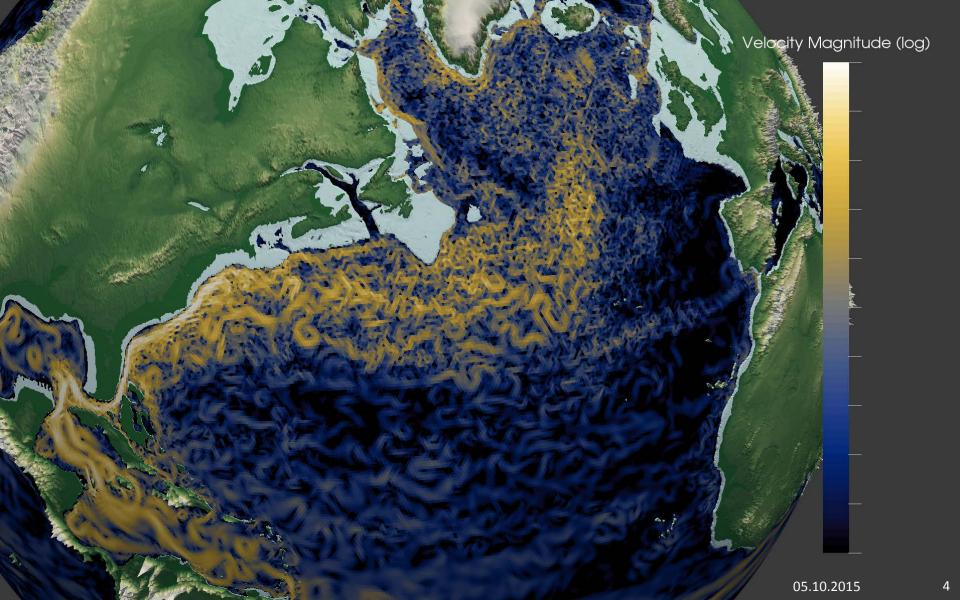
After you are done with your work, please stop the VNC server and clean up the node.

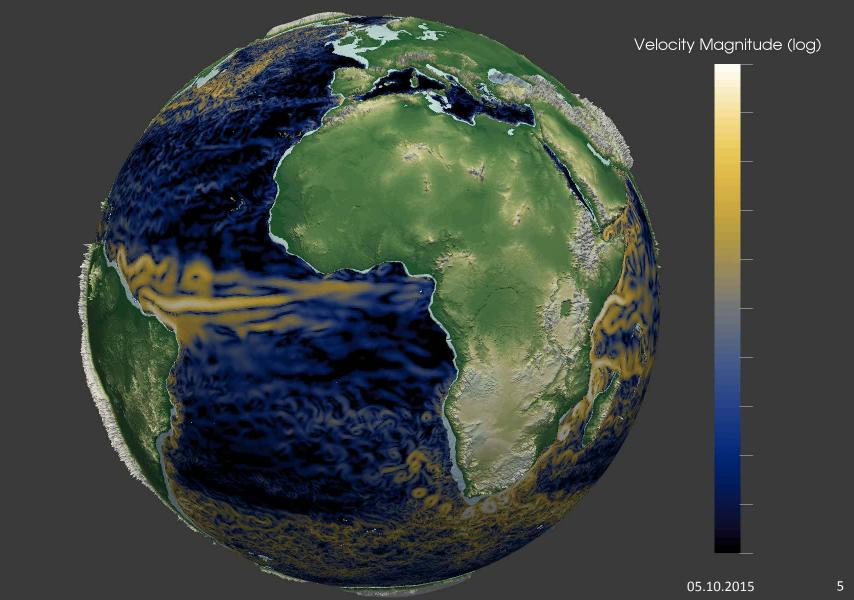
```
mg100:~> /opt/TurboVNC/bin/vncserver -kill :1
```

#### Running visualization applications

somewhere:~> vncviewer mg100.dkrz.de:1

Now a window opens, showing you the virtual X11 session that is running on your GPU node. On Mistral, all our visualization software is made available using modules. The command

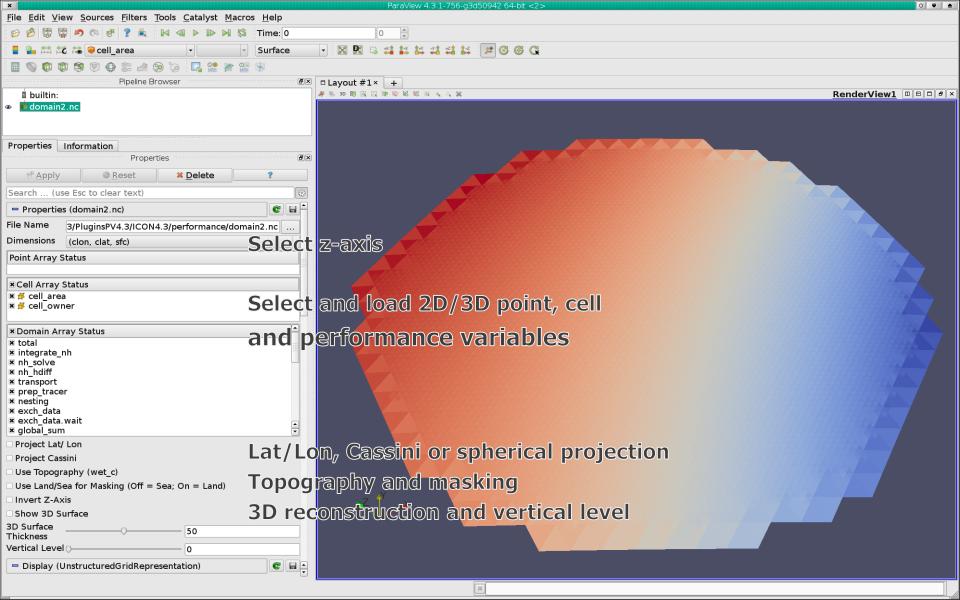






### vtkCDIReader

- Reads ICON/CDI netCDF data
- 2D/3D cell/point data on all levels (depth, depth\_2, ...)
- Runs in Client/Server mode
- 3D reconstruction with correct layer thickness
- Spherical, Lon/Lat and Cassini projection
- Bathymetry masking (wet\_c) and earth topography
- Visualization of performance data
- Plugin submitted to Kitware for official integration



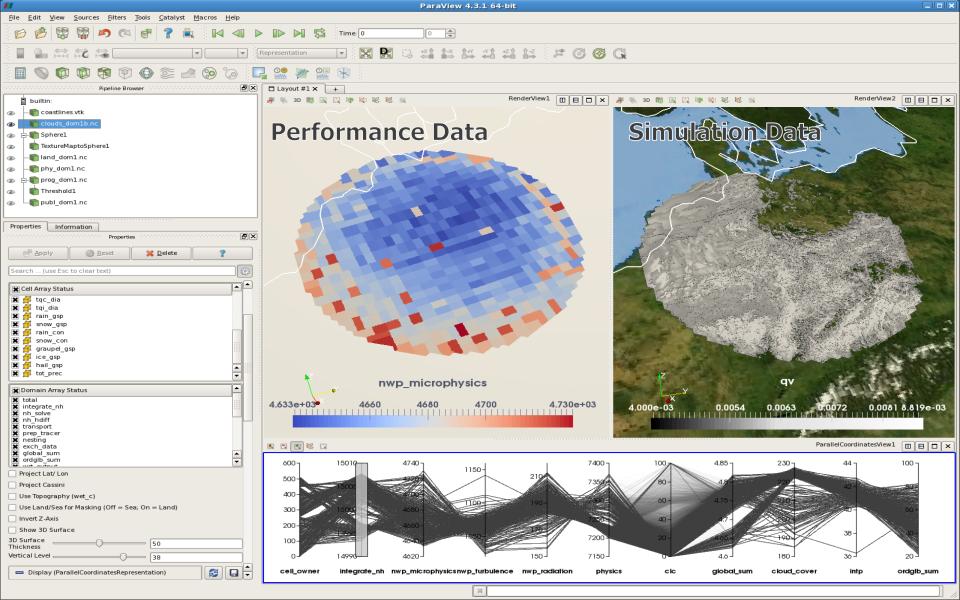


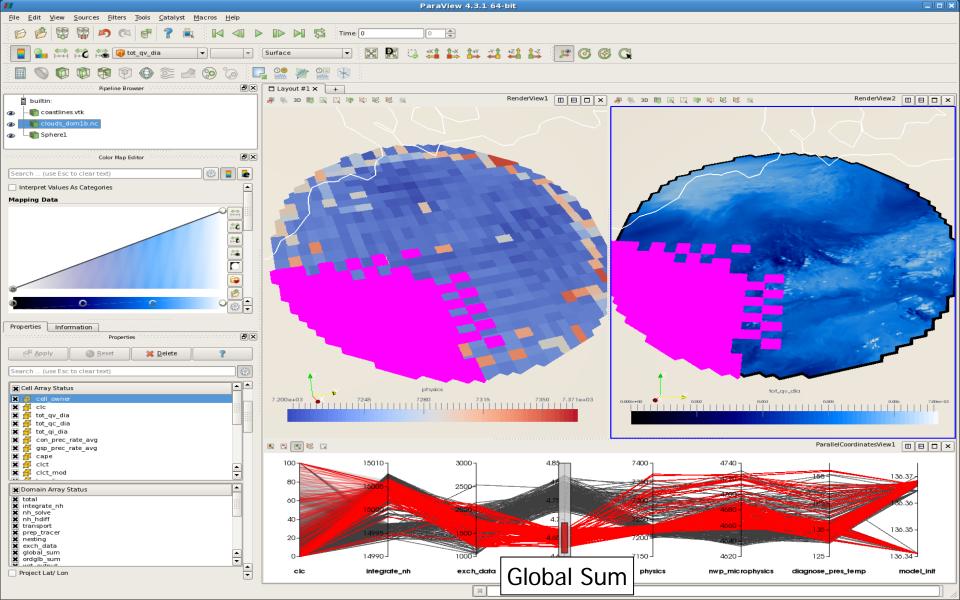
# Visualization of performance data

- ICON data with cell/point variables and performance data
- Select interesting areas and display results using parallel coordinates view
- Integrated into vtkCDIReader
- Requires domain decomposition and performance data

#### **Next Slides:**

HDCP2 Example (350.000 cells; 512 processors)







# InSitu with ParaView/Catalyst

- In-situ (co)processing/visualization using Catalyst extension
- Adaptor required that connects ICON (model) and Catalyst
- Two possibilities for co-processing:
  - Batch-visualization using pre-defined Python scripts
  - Live visualization within a client/server setting
- In development (very simple demo using ICON ocean model)

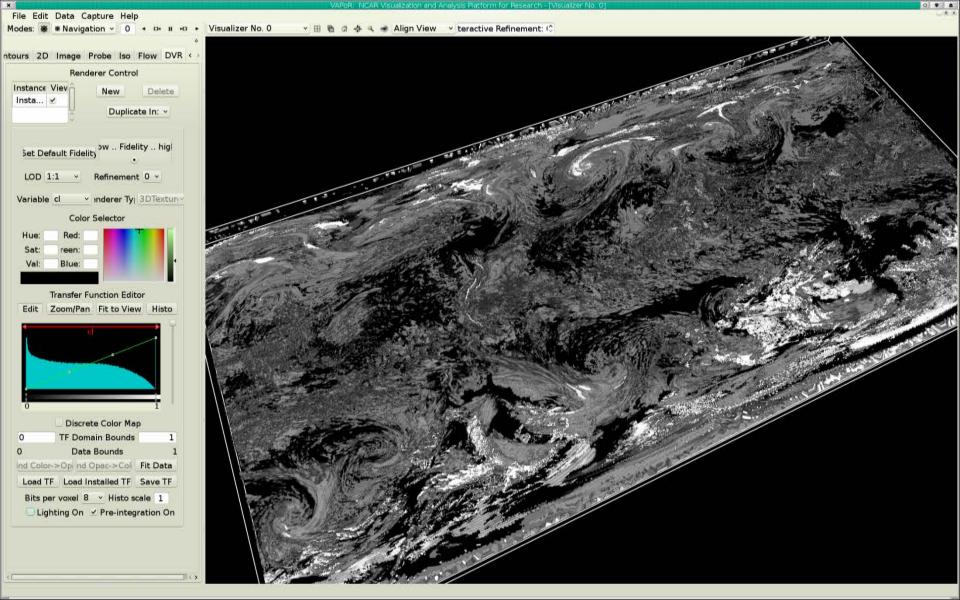


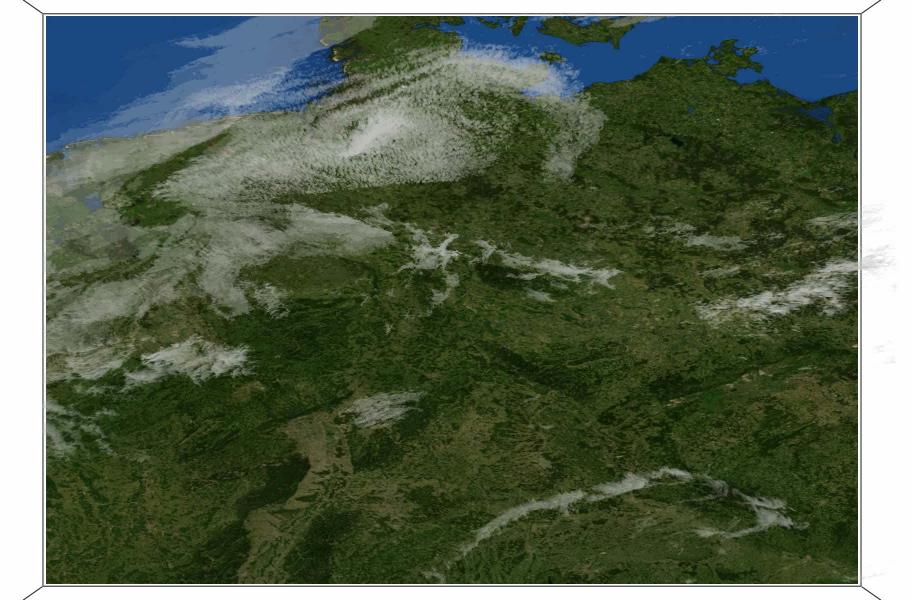
## **ICON** extension for Vapor

- Vapor developed at NCAR (open source)
- Supports wavelet compression and LoD rendering for very large (regular) data sets
- Extension for ICON/MPAS data available
  - Currently only with fast on-the-fly resampling to regular grid
  - - Wavelet compression for ICON/MPAS data

### **Next Slides:**

- ICON HDCP2 (1.3 million cells, CL)
- ICON HDCP2 (350 million cells, CLW)







## Help and Assistance

- DKRZ Website (<u>www.dkrz.de/Nutzerportal-en/doku/vis</u>)
- Tutorials & lectures (Avizo, NCL, ParaView, Vapor)

Niklas Röber (DKRZ) 05.10.2015