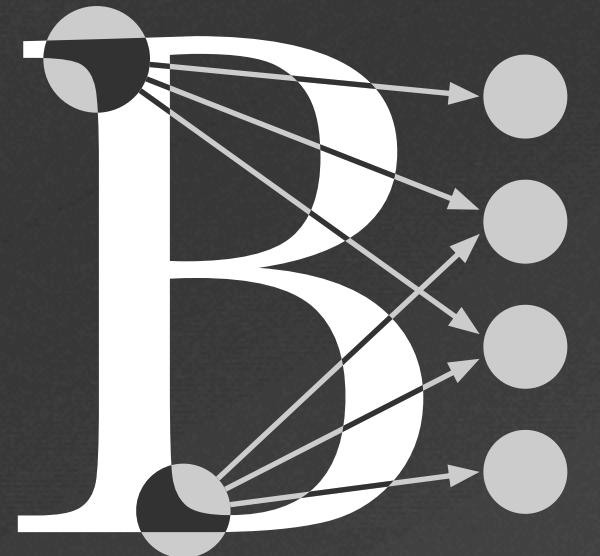


Baskerville - Accelerate your research with GPUs

Accelerating Data Engineering Pipelines

Tuesday 7th February 2023



Why is data engineering important?

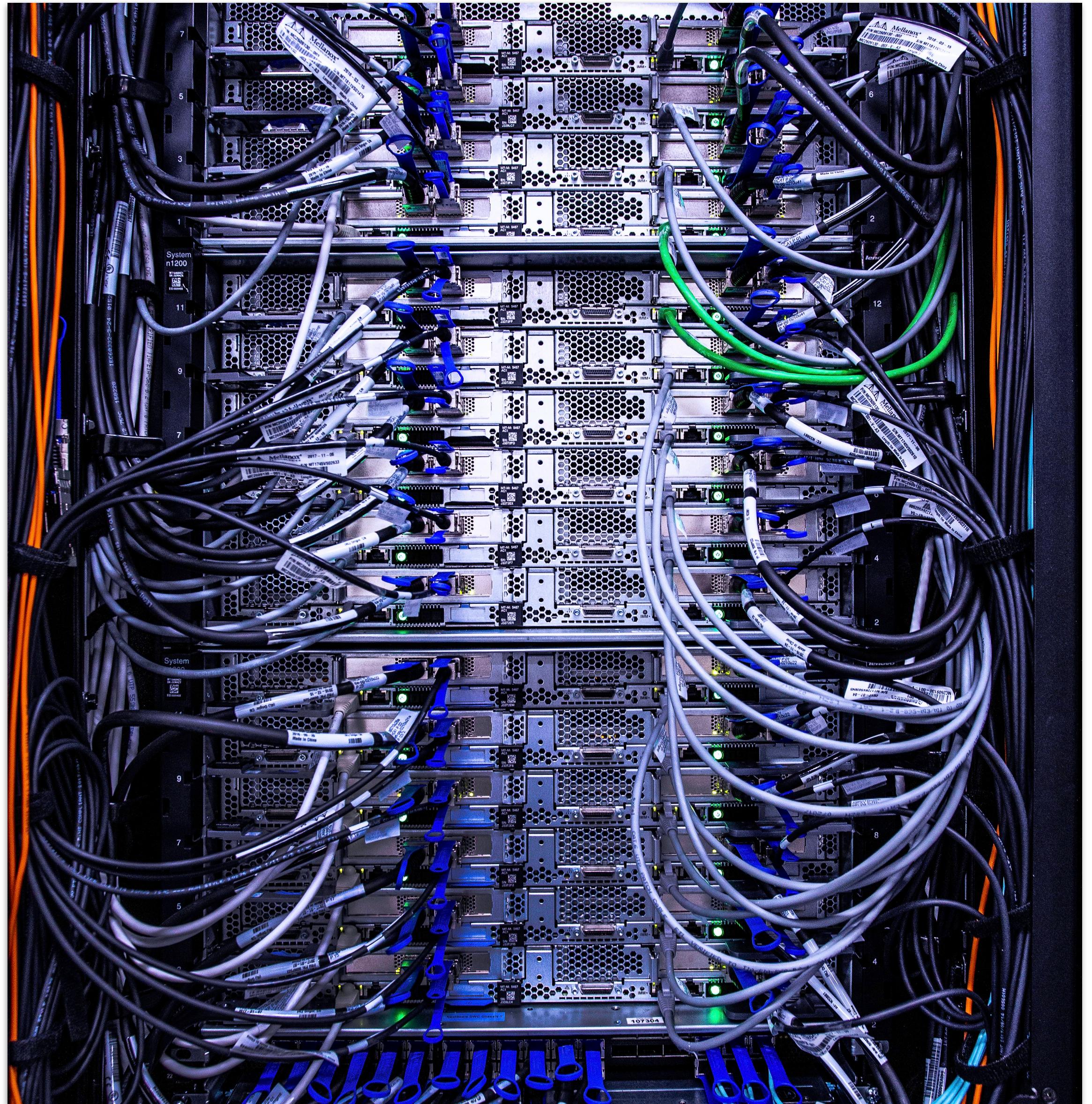
79 Zettabytes

of data generated world wide in 2021

(that's $79 \times 10^{21} = 79\,000\,000\,000\,000\,000\,000$ bytes!)

Data Engineering

- ◆ Storing, analysing and visualising large volumes of data is not fast enough using traditional methods (SQL, CPUs)
- ◆ Essential to accelerate and parallelise processes using multiple GPUs
- ◆ This workshop will guide you through the tools to manipulate large datasets and visualise results using
 - ◆ cuDF
 - ◆ Dask
 - ◆ Plotly



Baskerville - a national accelerated compute system funded by EPSRC

cuDF

GPU dataframe library

- ◆ Pandas-like API
- ◆ Built on the Apache Arrow columnar memory format
- ◆ Uses CUDA under the hood so you don't have to learn C/C++/Fortran
- ◆ For workflows on a single GPU or if your data fits in memory on a single GPU
- ◆ Multi-GPU support with Dask





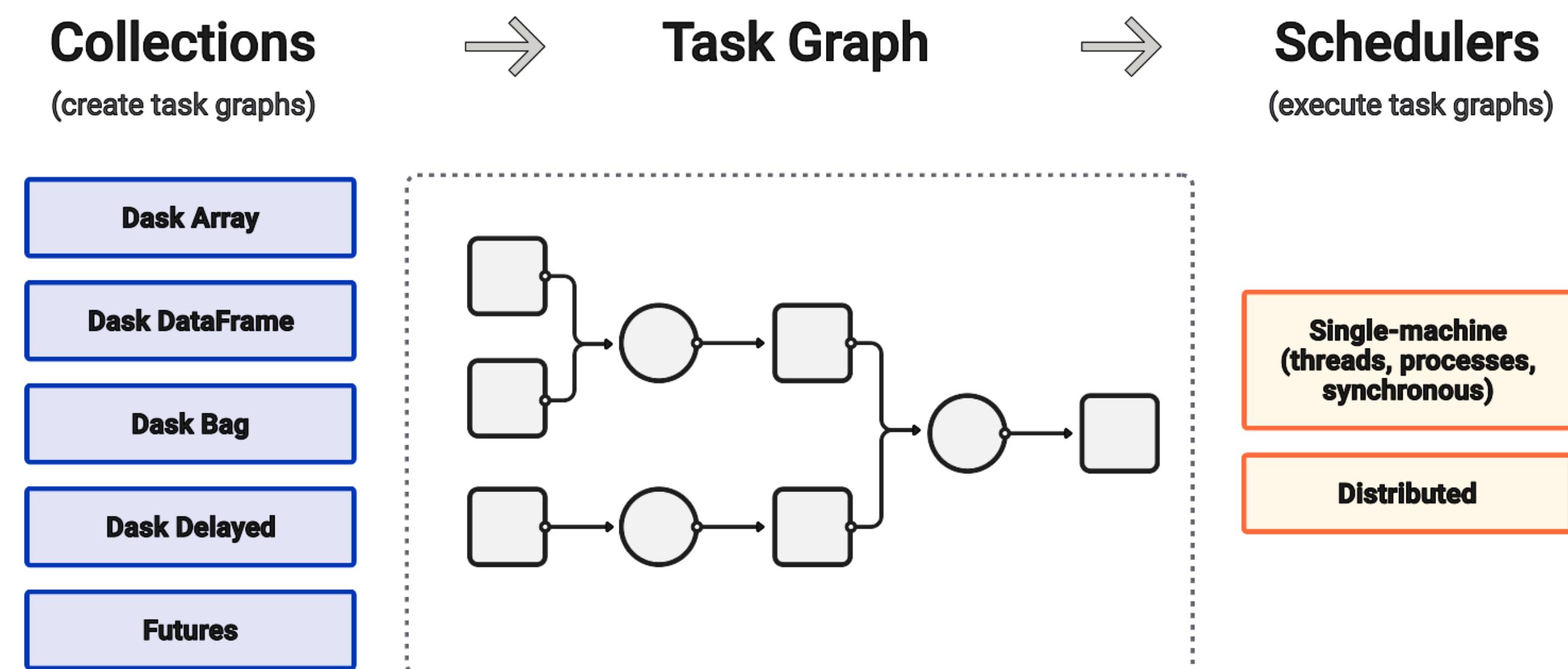
Flexible parallel computing

- ◆ Dask for CPUs and Dask-cuDF for GPUs
- ◆ Stages of computation

- ◆ “Lazy” = calculation computed only when needed

- ◆ Operations on dataframe are “queued-up” and built into task graphs

- ◆ Run with `.compute()` or `.persist()`





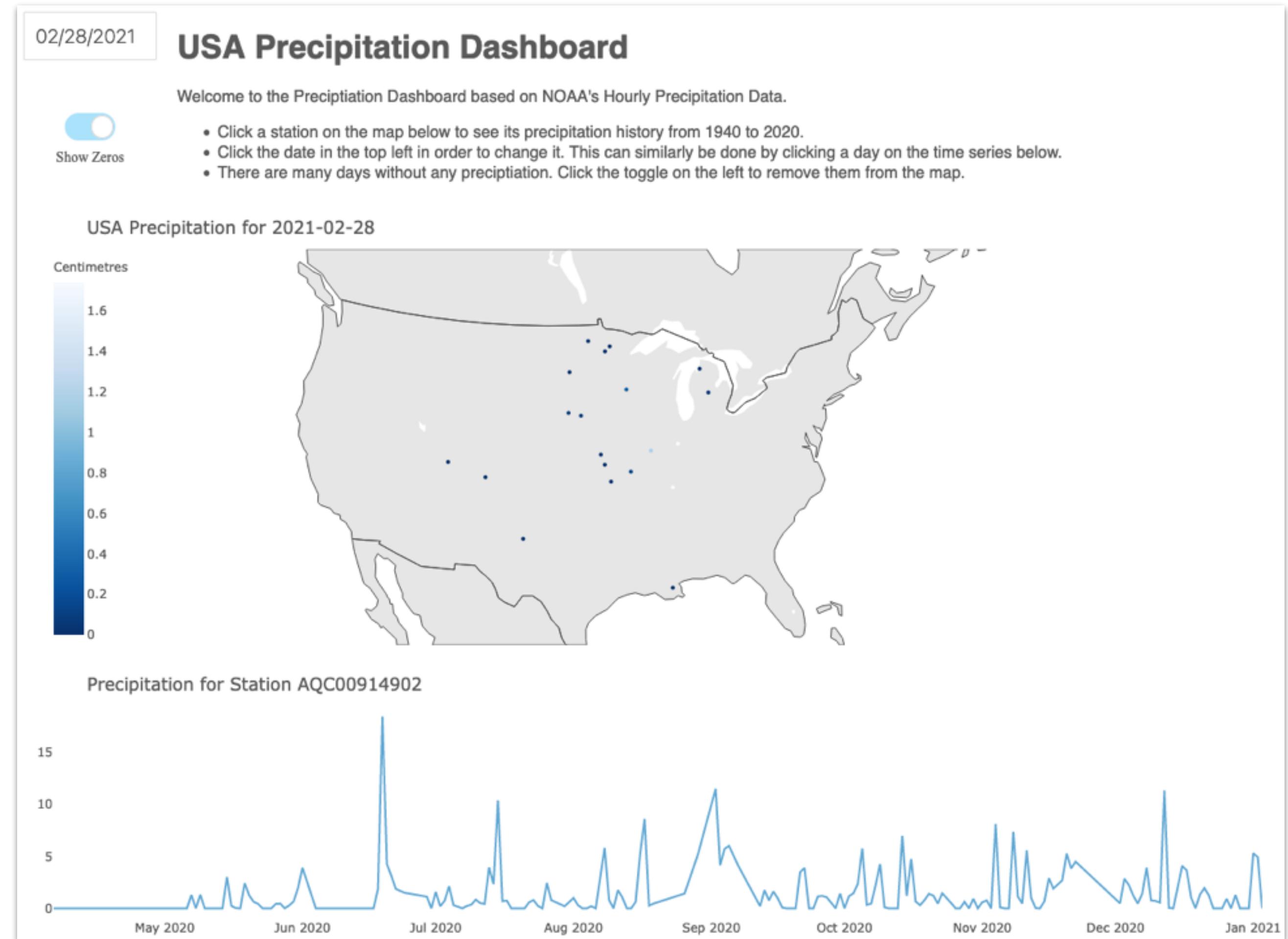
Interactive plotting library

- ◆ Build interactive web-based visualisations
- ◆ Rendered using JavaScript under the hood
- ◆ Export static images with Kaleido for non-web plots
- ◆ Integrate into Dash applications



Workshop Outline

- ◆ You will visualise precipitation data from US NOAA
- ◆ You will accelerate and parallelise a “colleague’s” unfinished notebook
- ◆ You will use Jupyter Notebooks on Baskerville Portal
- ◆ Each account is limited to 2 GPUs each





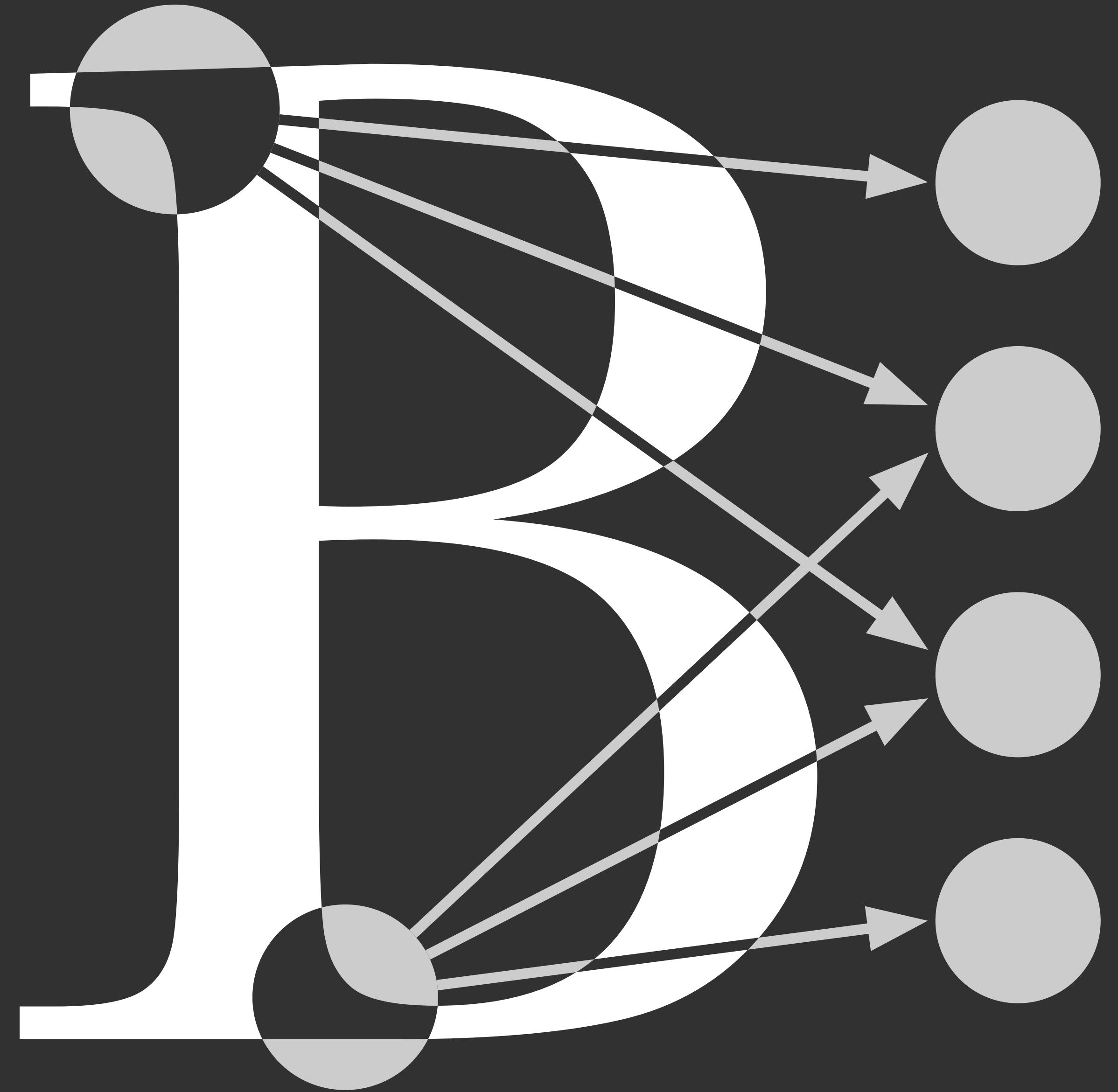
The Baskerville portal provides web-based access to the Baskerville Tier 2 system

This service is operated by Advanced Research Computing at the University of Birmingham and is funded by EPSRC
Grant EP/T022221/1

Workshop Setup

- 1) Create a symlink from the project to your home folder with ***In /bask/projects/w/wongj-bham-training ~***
- 2) Create your user folder with
cd ~/wongj-bham-training/users && mkdir \$USER
- 3) Setup your environment with ***source ./create_participant.sh*** (takes a while)
- 4) Close and re-launch the JupyterLab server (make sure ‘Show Conda Environments’ is ticked)
- 5) Work through ***users/\$USER/info_data_engineering/challenge_instructions.ipynb***
- 6) Challenge yourself with ***users/\$USER/info_data_engineering/challenge_notebook.ipynb***
- 7) Results are collected at 16:30

Will your notebook feature in the Top 5?



Collecting results...