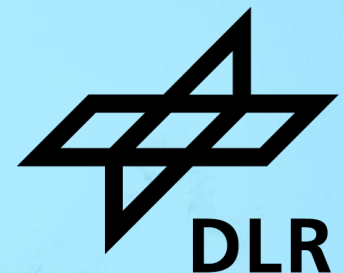


Implementing the Singular Value Decomposition in the Helmholtz Analytics Toolkit HeAT

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What is the Singular Value Decomposition (SVD)?



Singular Value Decomposition: Given $A \in \mathbb{R}^{m \times n}$ there are uniquely determined orthonormal $U \in \mathbb{R}^{m \times r}$, $V \in \mathbb{R}^{n \times r}$ and $\Sigma = \text{diag}(\sigma_1, \dots, \sigma_r) \in \mathbb{R}^{r \times r}$, $\sigma_1 \geq \dots \geq \sigma_r > 0$ ("singular values"), such that

$$A = U \cdot \Sigma \cdot V^T$$
A diagram illustrating the SVD equation $A = U \cdot \Sigma \cdot V^T$. The matrix A is represented by a large gray rectangle. The matrix U is a tall, narrow gray rectangle. The matrix Σ is a small square with a diagonal line from the top-left to the bottom-right. The matrix V^T is a wide, short gray rectangle.

...and why is it important for Machine Learning/Data Science?

SVD (also known as **PCA** – "Principal Component Analysis") often serves as an **important pre-processing step**

- Determines "*optimal linear coordinate system*" for a given data set
- *Compression/Reduction to the main features* due to low-rank best-approximation property of truncated SVD

The Helmholtz Analytics Toolkit HeAT



HEAT
Helmholtz Analytics Toolkit



HELMHOLTZ
Analytics Framework

- Developed by
 - **KIT** (Markus Götz, Daniel Coquelin, Charlotte Debus)
 - **Jülich Supercomputing Center (JSC)** (Claudia Comito, Kai Krajsek, Michael Tarnawa, Björn Hagemeyer)
 - **DLR SC-HPC** (Philipp Knechtges, Martin Siggel, Fabrice von der Lehr, Alexander Rüttgers)
 - and (as usual) plenty of student workers.
- MIT-licensed and available on GitHub: <https://github.com/helmholtz-analytics/heat>
- Originated from the Helmholtz Analytics Framework (HAF) project.
- Currently available in version 1.2.
- **Scope:** High-level library with a numpy-like interface that allows averaged numpy/pytorch-experienced ML users to port their code to HPC systems.

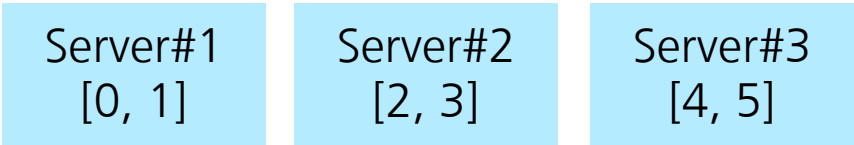
Reference: Götz, Debus, Coquelin, Krajsek, Comito, Knechtges, Hagemeyer, Tarnawa, Hanselmann, Siggel, Basermann, Streit. *HeAT - a Distributed and GPU-accelerated Tensor Framework for Data Analytics*. In 2020 IEEE International Conference on Big Data (Big Data) (pp. 276-287).

Data Distribution in HeAT

- A distributed HeAT tensor („**DNDarray**“) is composed of multiple local **torch.Tensor**'s
- The tensor can be split along an arbitrary axis

```

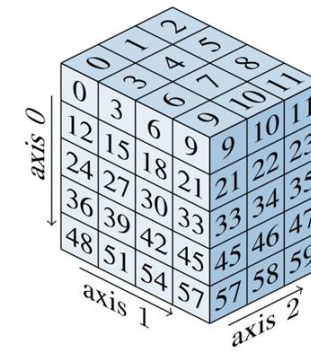
Example:  import heat as ht
          # construct a range tensor
          >>> range_data = ht.arange(6, split=1)
    
```



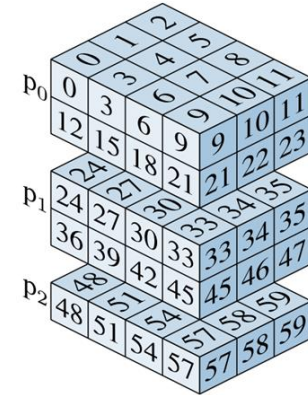
```

>>> range_data.mean()
2.5
>>> range_data.argmax()
5
    
```

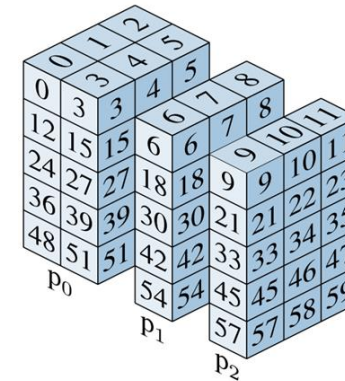
- Parallelization utilizing **MPI** (“Message Passing Interface“) and **mpi4py**, GPU-support inherited from **torch**



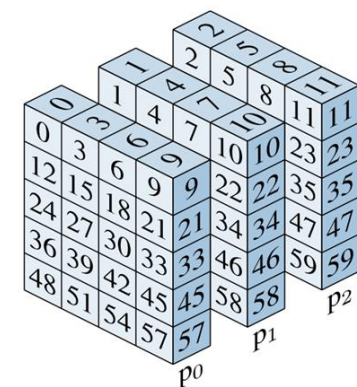
(a) split=None



(b) split=0



(c) split=1



(d) split=2

What is available?

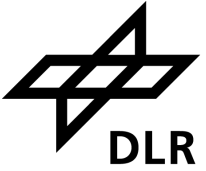
(Excerpt)



All relevant options are available on **both CPU and GPU**

- *Global functions:*
 - Global and local setters and getters
 - mean, std, var
 - max, argmax, min, argmin
 - transpose
 - sort
- *Elementwise functions:*
 - cov
 - where
 - nonzero
- *Parallel IO using HDF5*
- *Linear Algebra:*
 - Matrix-matrix multiplication
 - Matrix-vector multiplication (dot)
 - QR
- *Analysis / Regression:*
 - K-means clustering
 - Spectral clustering
 - Lasso regression
 - DASO (data-parallel NN)
- *Basic building blocks:*
 - concatenate
 - diag
 - arange
 - Distributed random numbers
 - ...

Currently being developed



- **SVD (PCA)**
 - **work in progress: distributed hierarchical SVD** (hSVD, available as branch on github)
 - Future work: Zolotarev-SVD*
- Autograd / Automatic Differentiation
- K-nearest neighbours
- Logistic Regression
- Neural Networks
- Random Decision Trees
- SVM
- DBSCAN
- GMM
- ICA

*Nakatsukasa, Freund. *Computing fundamental matrix decompositions accurately via the matrix sign function in two iterations: The power of Zolotarev's functions*. SIAM Rev. 58, 3 (2016), 461--493.

Distributed hierarchical SVD

Distributed hierarchical SVD



- „**Big Data is Low Rank**“* – in many applications, it suffices to compute the first (i.e. largest) „few“ singular values and vectors
- Exploit the specific structure of **DNDarray**'s: apply algorithms that can make use of **torch**-routines on the local arrays *in MPI-parallel* and somehow „merge“ the results

The „distributed hierarchical“ idea in a nutshell – Example: Computation of the mean value on 2 processes

Process 1

numbers $x_1, x_2, \dots, x_{m-1}, x_m, x_{m+1}, x_{m+2}, \dots, x_{m+n-1}, x_{m+n}$

Compute „local“ mean $\bar{x}^{(1)}$

Receive „local“ mean $\bar{x}^{(2)}$

„Merge“:

Compute global mean by $\bar{x} = \frac{m}{m+n} \bar{x}^{(1)} + \frac{n}{m+n} \bar{x}^{(2)}$

Process 2

Compute „local“ mean $\bar{x}^{(2)}$

Send

* see, e.g.: Udell, Townsend. *Why are big data matrices approximately low rank?* SIAM J. Math. Data Sci., 1 (2019).

Distributed hierarchical SVD



Example: Compute *truncated* SVD on 2 processes

Process 1

Matrix $[A_1 | A_2]$ (split along columns)

Process 2

Compute „local“ SVD $A_1 = U_1 \Sigma_1 V_1^T$
Truncate to rank r and form $U_1[:, :r] \Sigma_1[:, :r]$

Compute „local“ SVD $A_2 = U_2 \Sigma_2 V_2^T$
Truncate to rank r and form $U_2[:, :r] \Sigma_2[:, :r]$

Receive $U_2[:, :r] \Sigma_2[:, :r]$

Send

„Merge“:

Compute the SVD of the concatenation
 $[U_1[:, :r] \Sigma_1[:, :r] | U_2[:, :r] \Sigma_2[:, :r]]$
and *truncate* to rank r

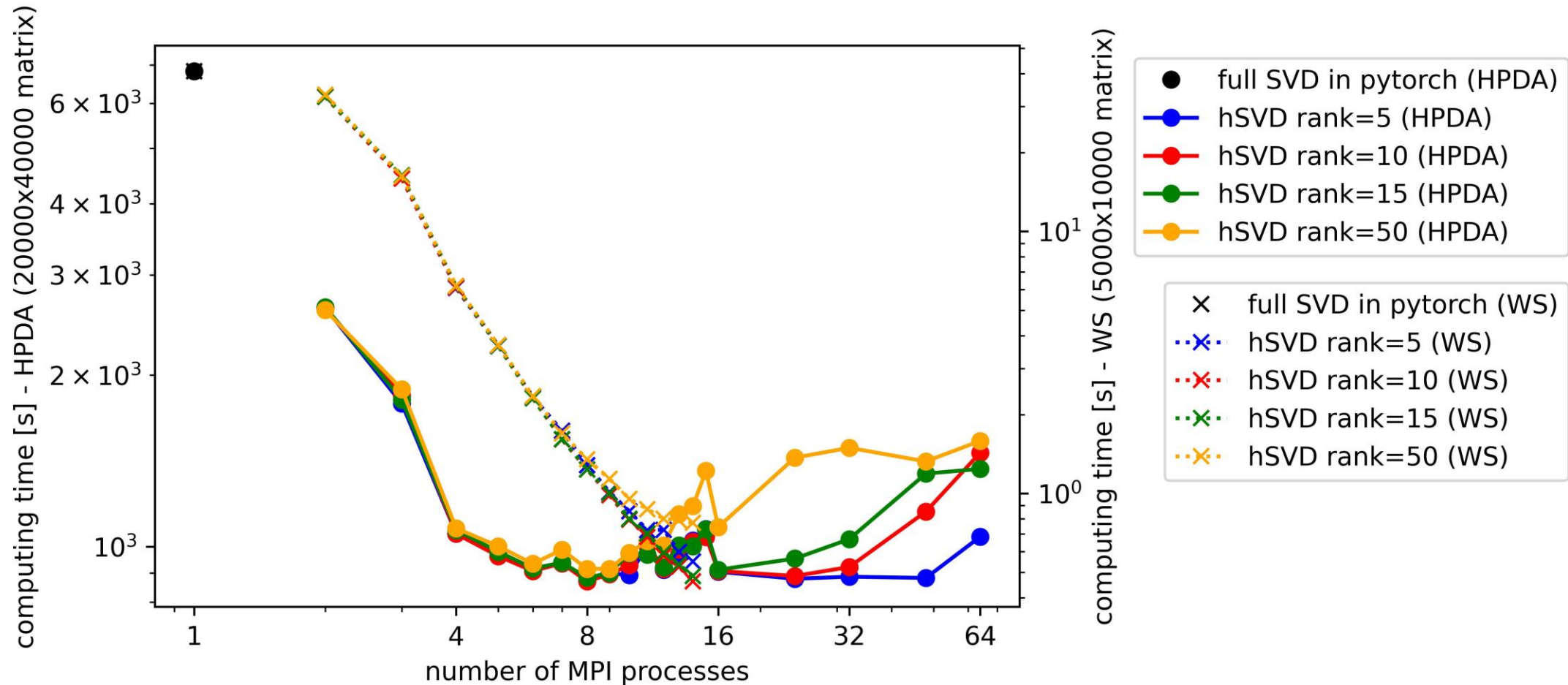
- Generalization towards „merging“ along more complicated tree-structures is straightforward
- Theoretical analysis including error estimation/control is available

References:

Iwen, Ong. *A distributed and incremental SVD algorithm for agglomerative data analysis on large networks*. SIAM J. Matrix Anal. Appl., 37(4), 2016.
Himpe, Leibner, Rave. *Hierarchical approximate proper orthogonal decomposition*. SIAM J. Sci. Comput., 40 (5), 2018.

Numerical Experiment: hSVD on CPU

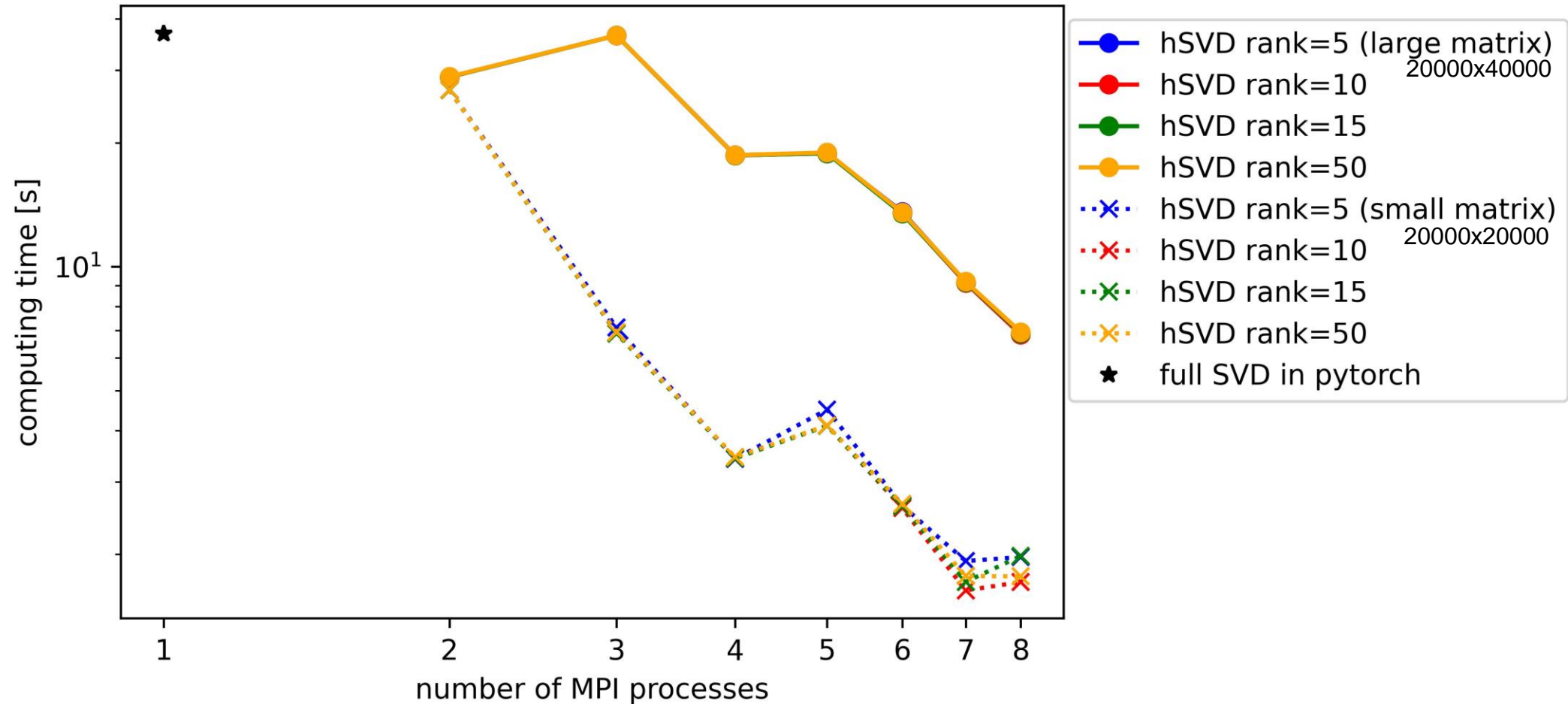
(Strong scaling)



Data: synthetic data set with rapidly decaying singular values [rel. truncation errors (in Frobenius norm): ~10% for rank 5, ~1% for rank 10 and < 0.002% for rank 15...]
Hardware: HPDA-Cluster SC-HPC, BS (=HPDA) and Linux Workstation, KP (=WS)

Numerical Experiment: hSVD on GPU

(strong scaling)



Data: synthetic data set with rapidly decaying singular values [rel. truncation errors (in Frobenius norm): ~10% for rank 5, ~1% for rank 10 and < 0.002% for rank 15...]

Hardware: HPDA-Cluster SC-HPC, BS

Thank you for your attention!