

RUHR-UNIVERSITÄT BOCHUM

STREAMLINING THE ANALYSIS OF MAGNETIC RECONNECTION SIMULATIONS USING MACHINE LEARNING METHODS

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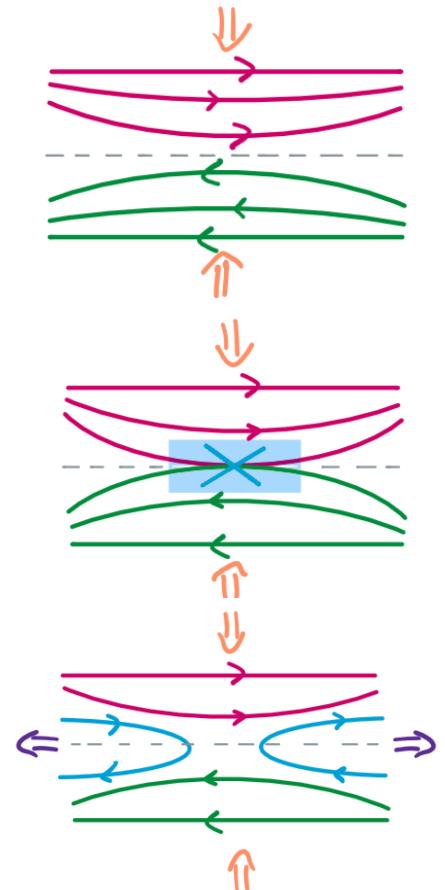
General Assembly SFB 1491 – Theoretische Physik 1 (Jun.-Prof. Dr. Maria Elena Innocenti)

Motivation | Magnetic Reconnection

- How are particles **preaccelerated** to reach high energies observed in **astrophysical contexts**? – possible answer: magnetic reconnection
- Magnetic field lines ‘snap open’ and reconnect in new topology
- **magnetic energy is converted** to heat and non-thermal acceleration

Fundamental for understanding:

- analyse and **classify** data from spacecrafts and simulations



Motivation | Machine Learning as Solution

Goal:

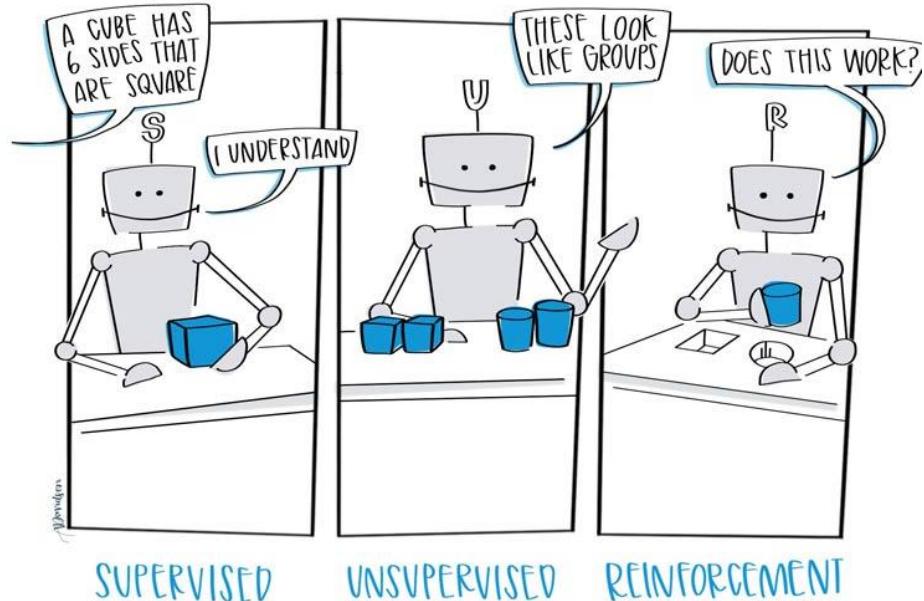
- Streamline identification of physically distinct regions of magnetic reconnection

Problems:

- huge amounts of data
- unconscious bias

Solution (?):

- Unsupervised machine learning



<https://www.ceralytics.com/3-types-of-machine-learning/>

Background Clustering and Self Organizing Maps (SOMs)

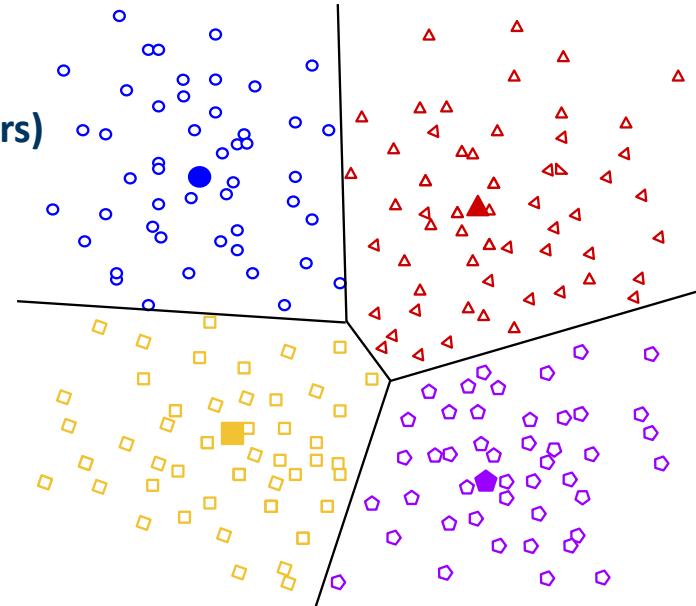
Background | Unsupervised learning methods

Clustering:

- Process of segmenting unlabeled data into subgroups (clusters) of similar input (Jo, 2021)
- Centroid clustering: Finds the prototypes of clusters
- **K-means** is the most widely used algorithm for centroid clustering

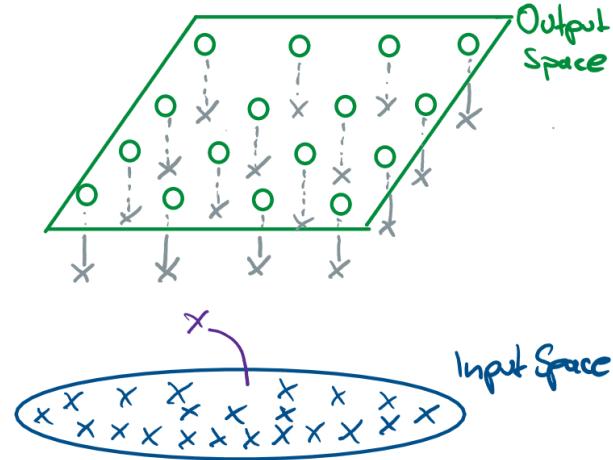
Self-organizing maps (Kohonen, 1982):

- Form two-dimensional maps of high-dimensional data
- **Preserve the topology** of input data
- Can be used for clustering but also offer powerful visualizations



Background | SOM training

Goal: Find weight values, such that adjacent units have similar values



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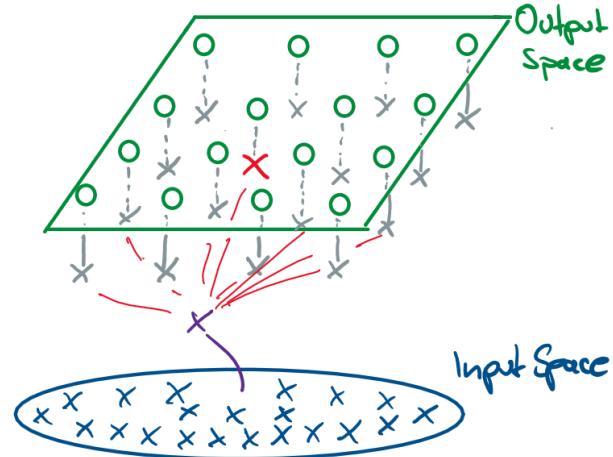
- Inputs are assigned to unit (-weights) that are most similar to them – Best Matching Units (BMU)
- The weights of the most similar unit and its neighbours get activated for weight update
- The activated unit's weights get updated

The SOM training is separable in three phases (van Hulle, 2012):

Competition

Collaboration

Weight Updates



Background | SOM training

Goal: Find weight values, such that adjacent units have similar values

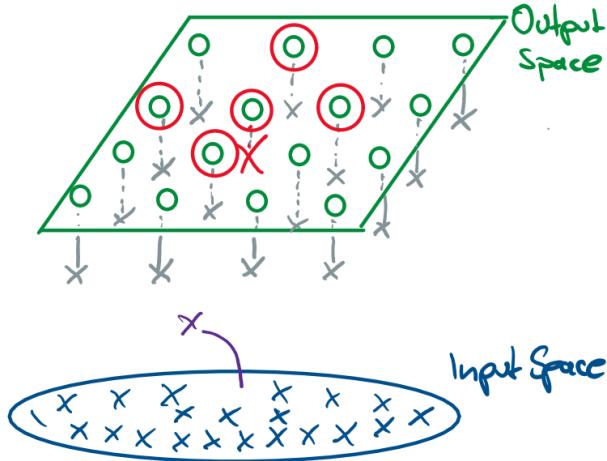
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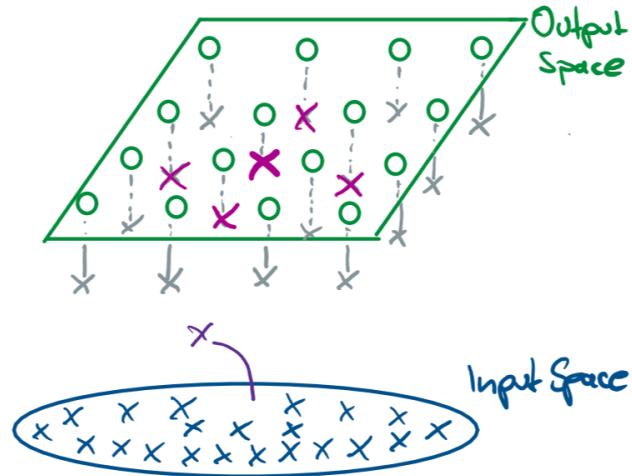
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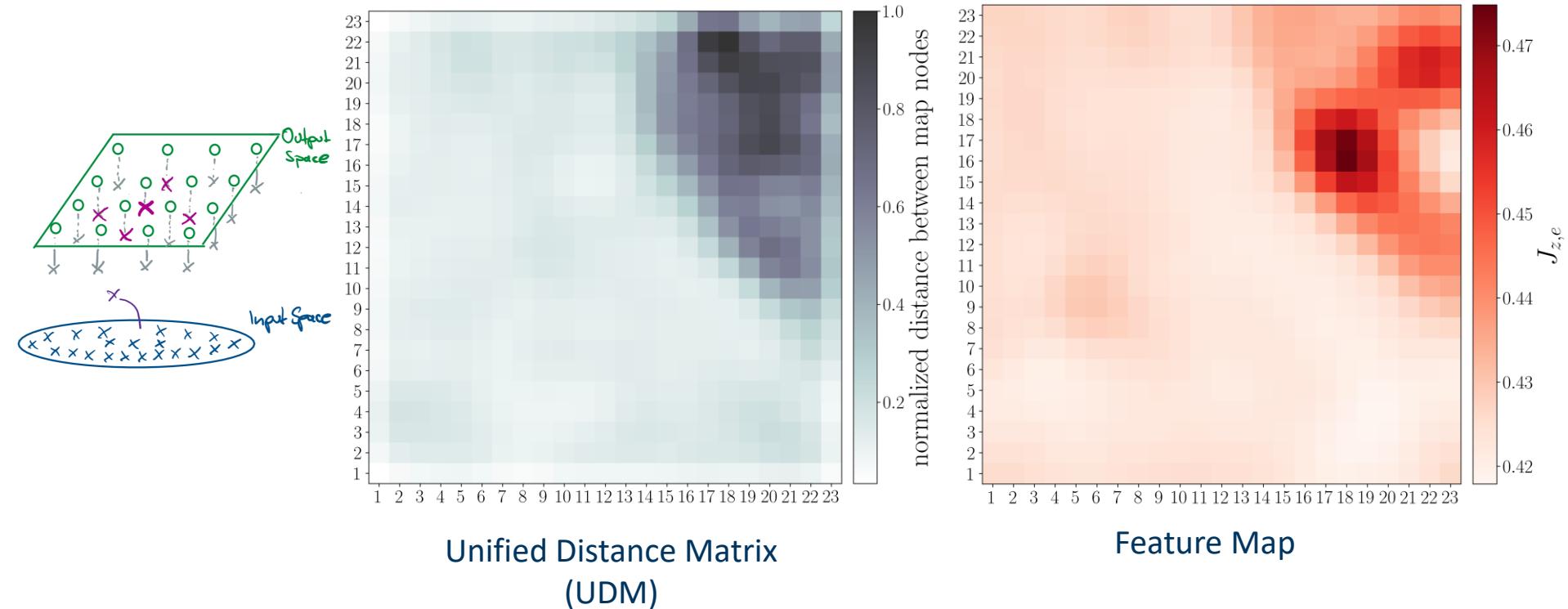
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Background | Visualizations of the SOM

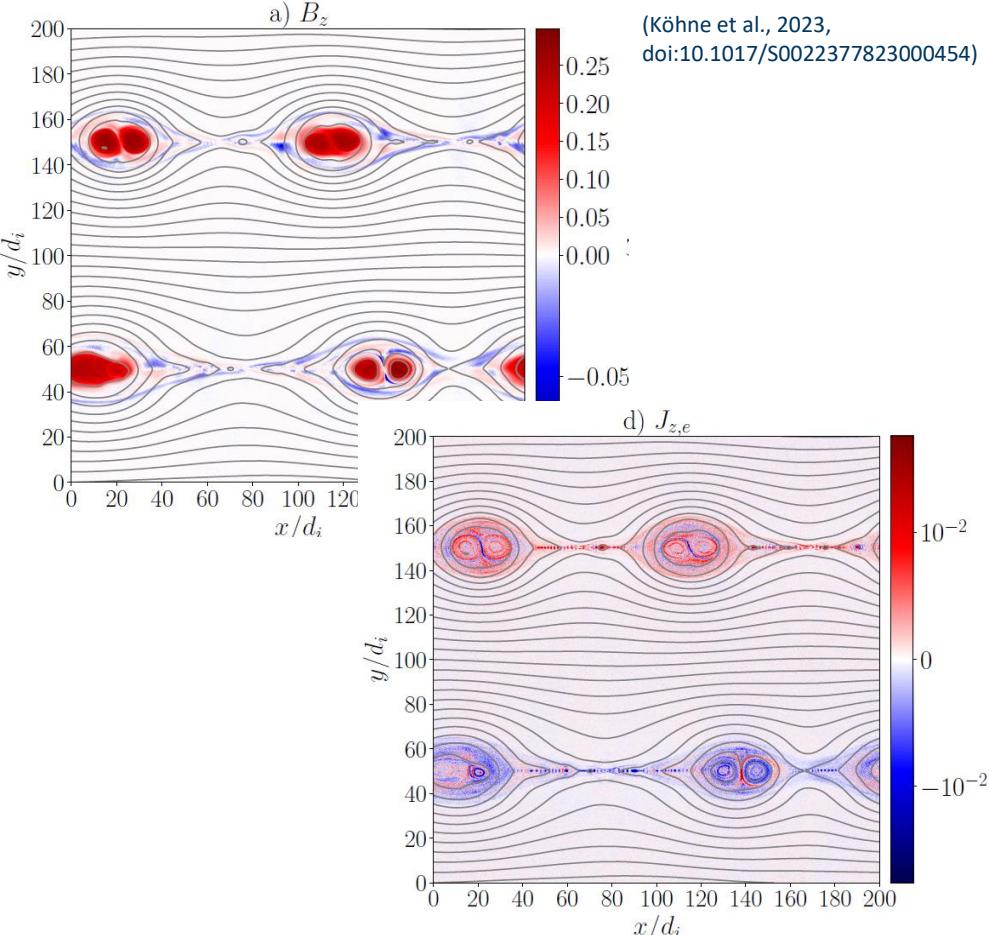


Data and Methodology

Data | Simulation

Data points produced by fully kinetic simulation
of plasmoid instability

- semi-implicit, energy conserving PIC code
ECsim (Lapenta et al., 2017)
 - Force free initial conditions, periodic boundary conditions, collisionless regime
 - Reduced mass ratio $m_r = 25$
- 4528384 samples with 26 features each



Method | How results were obtained

1. Preprocessing
2. Initialise SOM
3. SOM training on preprocessed data
4. k-Means clustering of SOM
5. Visualize results

Method

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Scaling the data

- Scale feature values according to defined rule
- Assures that all features have same level of influence on model
- Most common: scale to interval, e.g. [0,1]

Method

1. Preprocessing
2. Initialise SOM
3. SOM training on preprocessed data
4. k-Means clustering of SOM
5. Visualize results

SOM implementation used:

<https://github.com/JustGlowing/minisom>
(serial implementation)

<https://github.com/mistrello96/CUDA-SOM>
(parallel implementation)

(Hyper-) Parameters (Kohonen, 2014):

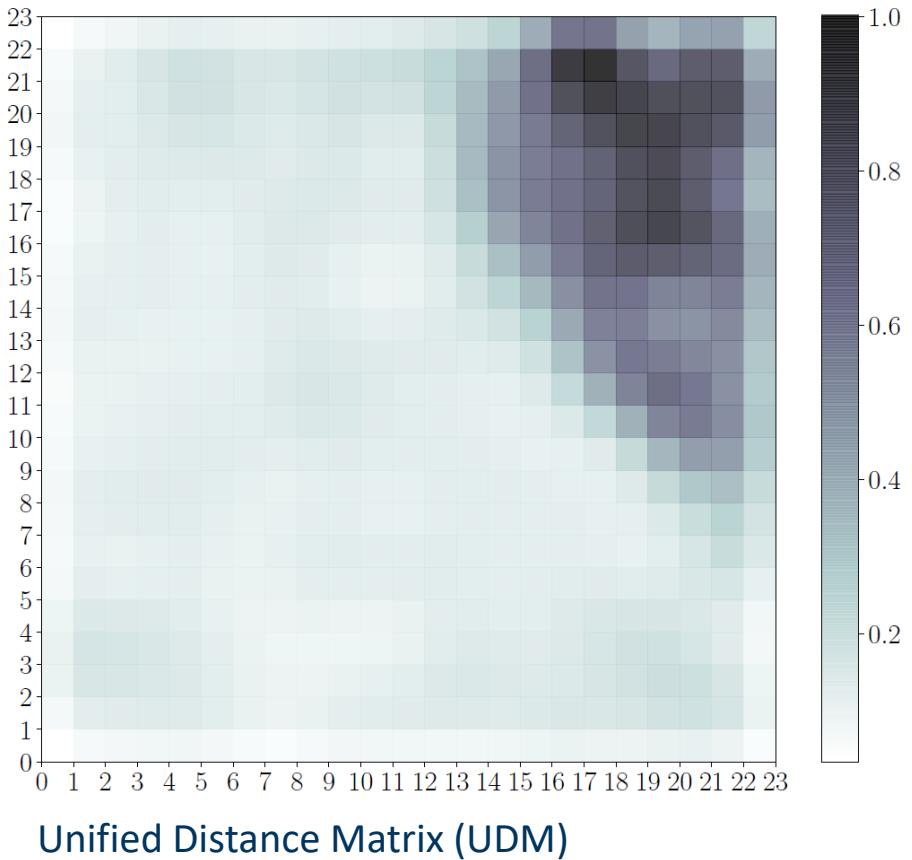
- Number of neurons $n \approx 5\sqrt{N}$ (J. Tian et al., 2014)
- Aspect ratio of x,y dimensions is the same as the ratio of the two first principal components (J. Tian et al., 2014)
- Initial neighborhood radius $\sigma_0 = 0.2 * \max\{x, y\}$
- Initial learning rate $\eta_0 = 0.5$
- # of iterations: $n_{iter} = 5N$
- Neighborhoodfunction: gaussian
- Neighborhood distance function: euclidean
- Decay function: asymptotic
- Weights initialization: random samples

Method

1. Preprocessing
 2. Initialise SOM
 3. SOM training on preprocessed data
 4. **k-Means clustering of SOM**
 5. Visualize results
- $k \in [2,8]$
 - Optimal cluster number was determined using Satopaa kneedle method (Satopaa et al., 2011)

Method

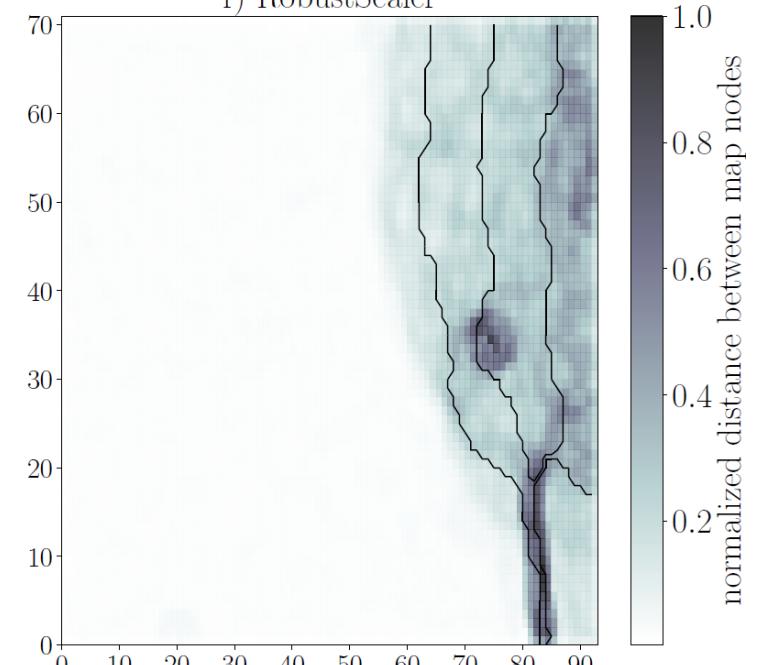
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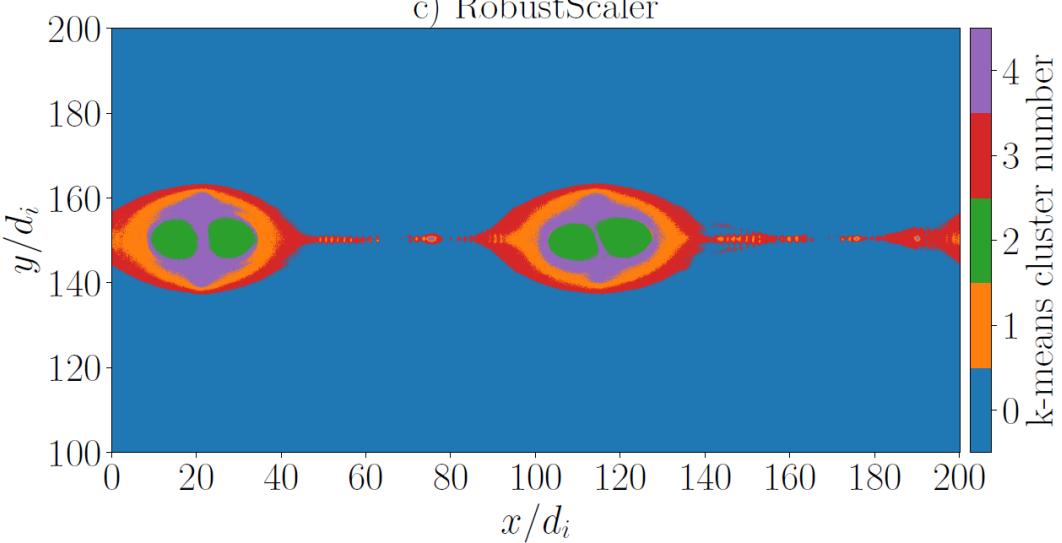
Clustering results

Results | Classification

f) RobustScaler

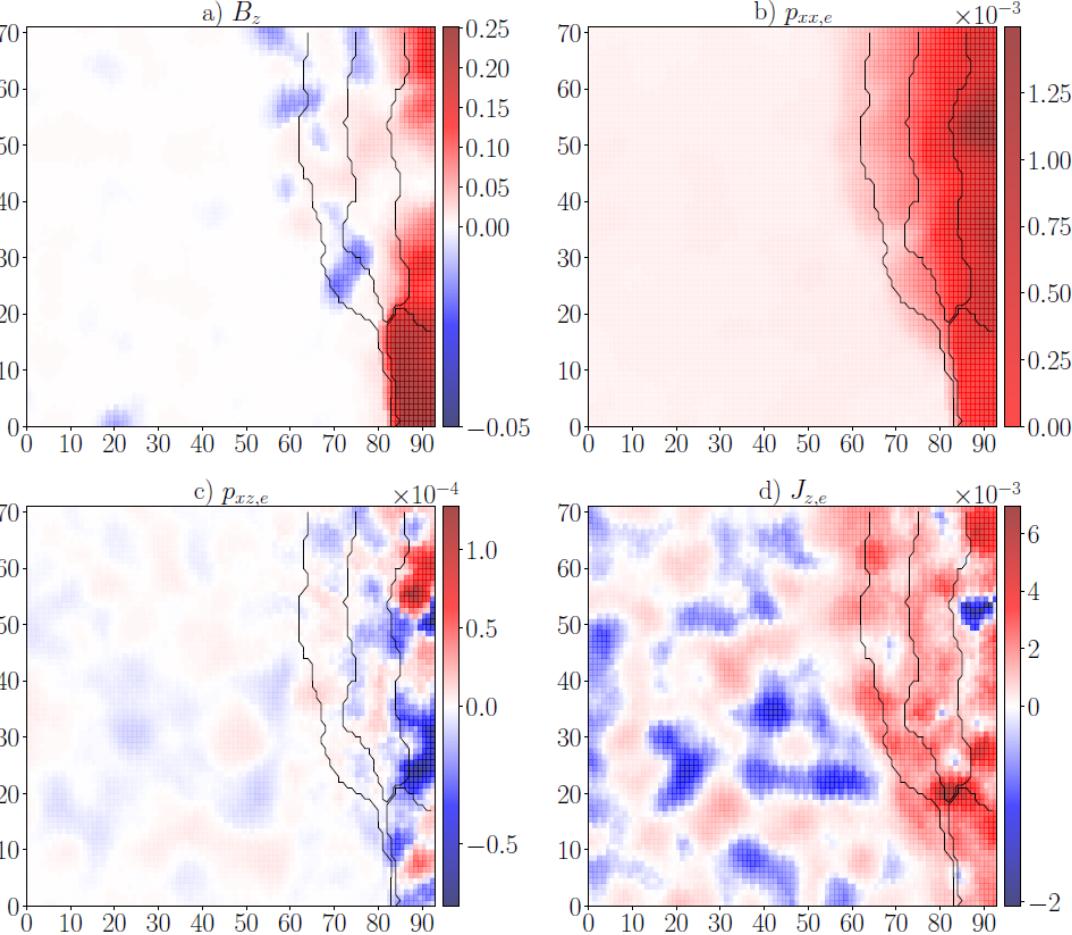
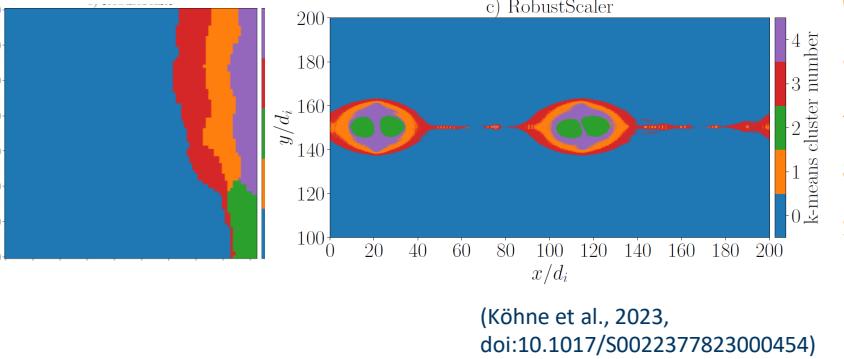


c) RobustScaler



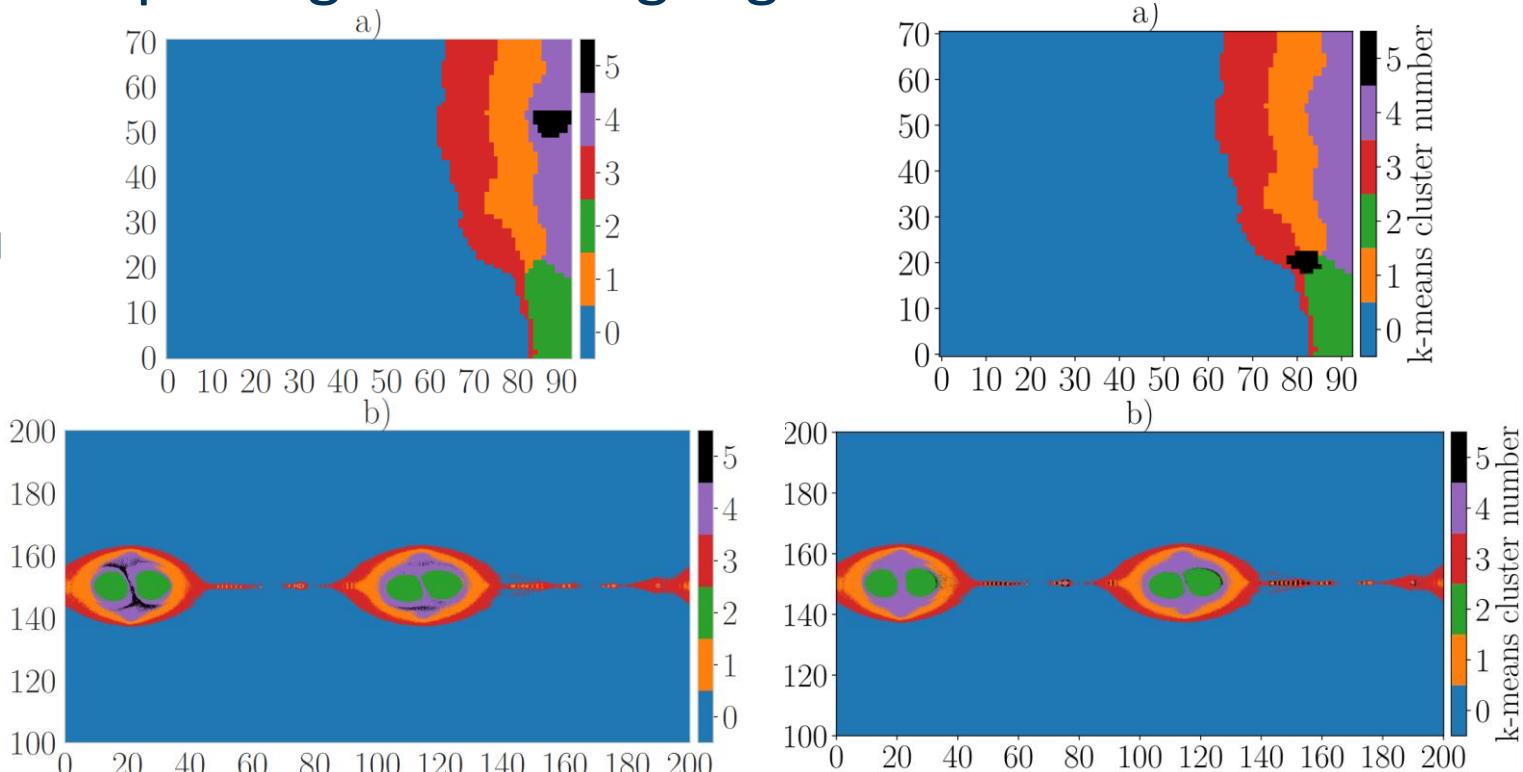
Results | Feature Maps

- show values of selected features in the weights of each node
- possibility to investigate peculiarities



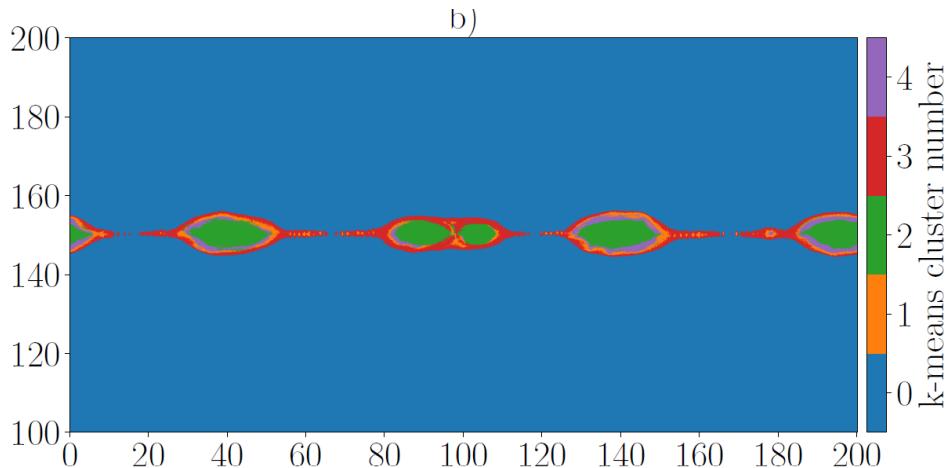
Results | Handpicking interesting regions

- extreme $J_{z,e}$ inside of purple cluster – **plasmoid merging regions**
- High-value $J_{z,e}$ at intersection of plasmoid clusters – **mini-plasmoids** at x-line



Results | Robustness of the results

- **Hyperparameter variation:** number of epochs, initial neighborhood radius, initial learning rate, initialization seed, number of nodes
 - Plasmoid classification stays very stable: run with largest deviation matches reference run to 86 %
 - most unstable clusters: intermediate plasmoid region (orange and purple)
- **Temporal variation:** classify data from earlier timestep of simulation using SOM trained on later timestep
 - General classification stays sensible
 - Plasmoid merging regions shrink



Summary and Outlook

Summary & Outlook

- To understand magnetic reconnection analysis of huge amounts of data is needed
- **Combination of unsupervised machine learning methods:** k-means and SOMs
 - Can identify **physically distinct regions** in fully kinetic simulations
 - SOMs offer many **intuitive investigation** possibilities

Further steps:

- Analysis and classification of **observational data** (Amaya et al., 2020)
- Usage in simulations to e.g. switch from one numerical method to another according to cluster

Unsupervised classification of fully kinetic simulations of plasmoid instability using self-organizing maps (SOMs)

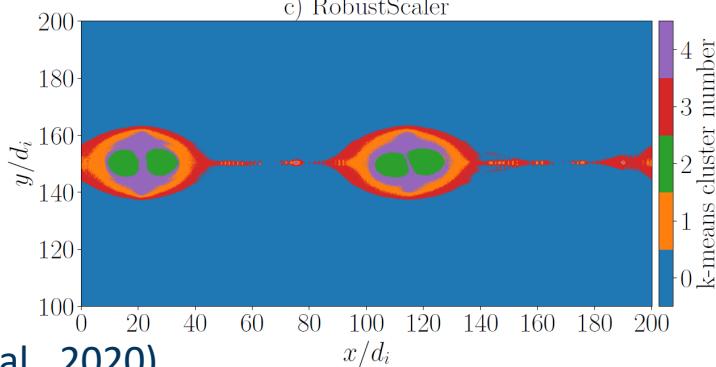
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Additional Slides

Clustering | K-means

Input:

number of clusters K
dataset $x_i, i \in (1, \dots, n)$

Output:

Set of K clusters

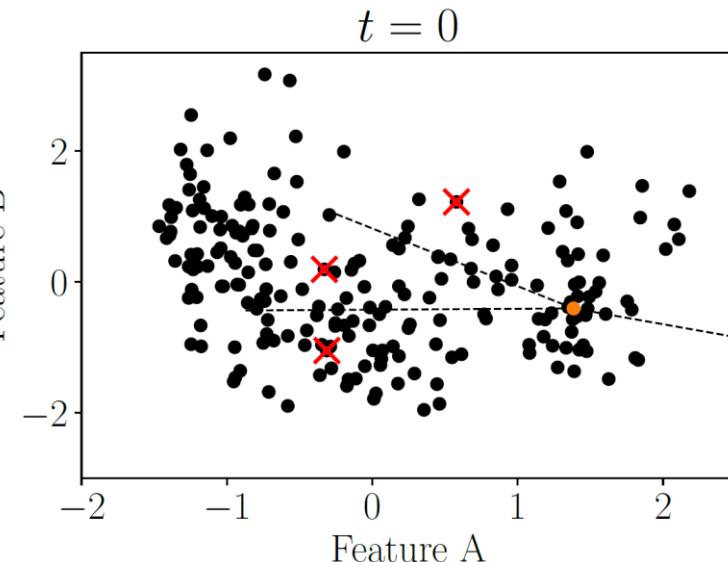
Algorithm:

1. Place k initial centroids randomly

2. Assign each data point to its closest centroid, minimizing:

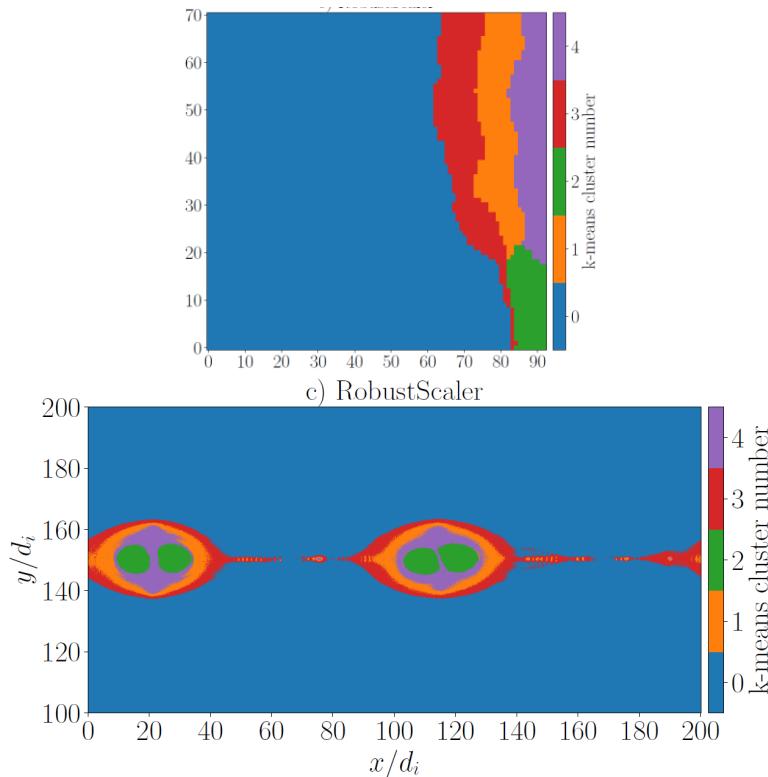
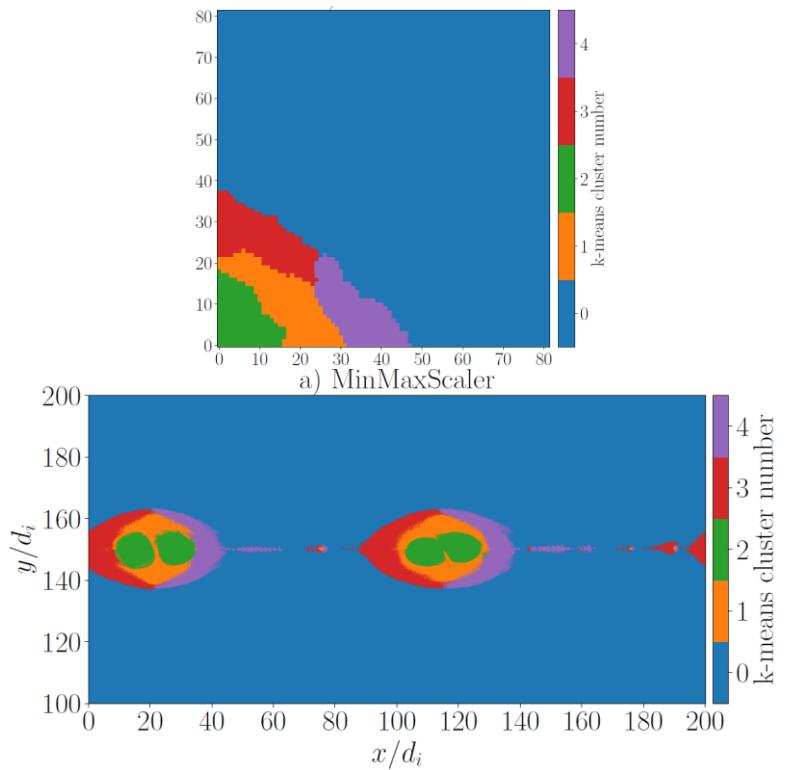
$$J = \sum_{j=1}^K \sum_{i=1}^n \|x_i - c_j\|^2$$

3. Update centroids - move them to mean of all x_i assigned to them

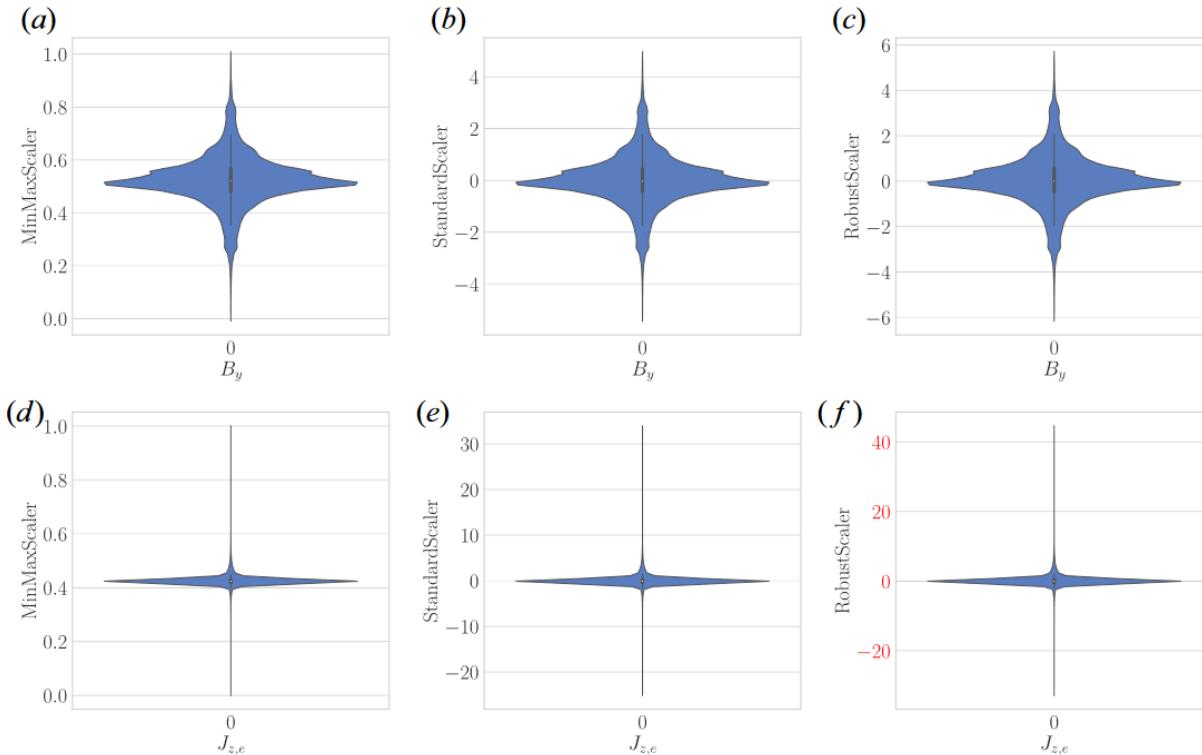


Break if convergence criterion is matched

Results | Importance of Scalers



Results | Importance of Scalers



SOM | Convergence

Two convergence phases: Topographic ordering of weights, Convergence of those weights

- Both are strictly speaking not mathematically proven (Cotrell, Fort, et al., 1998)
- SOMs do not have an objective function J where $\Delta w_i \propto \frac{\partial J}{\partial w_i}$

Practical convergence criteria:

- Maximum number of iterations
- Stop when Δw_i is under threshold
- Quantization error: average distance between each input sample and its best matching unit → should be as low as possible
- Topographic error: percentage of samples whose second-best matching unit is not adjacent to their BMUs