#### Analysis of quality measures for dimensionality reduction

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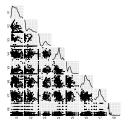
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#### Motivation



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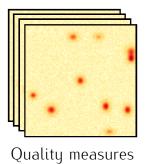


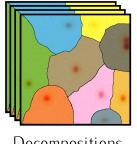
$$Q_{Stress}(x_i) = \sqrt{\sum_{j=1}^{n} (d_{ij} - \delta_{ij})^2 / \sum_{j=1}^{n} \delta_{ij}^2}$$

 $Q_{\text{Stress}}, \ Q_{\text{Residual variance}}, \ Q_{\text{RMSE}}, \ Q_{\text{Spearman}}...$ 

- Is more than one aspect of the data preserved by a method?
- How to compare different quality measures with each other?
- Do multiple quality measures *agree* on the data set?

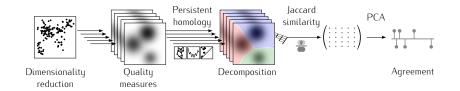
#### Our method





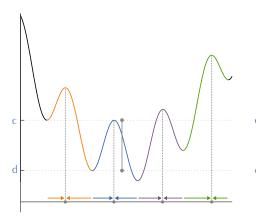
Decompositions

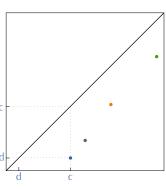
#### Workflow



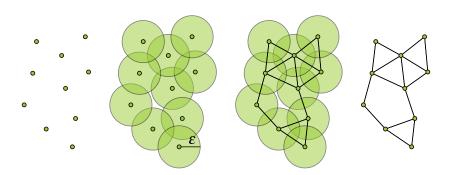
## Persistent homology

Superlevel set filtration





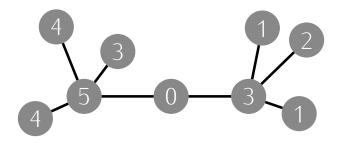
#### High-dimensional data



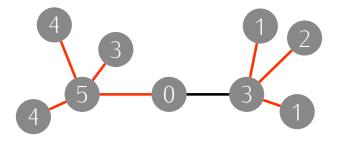
- Rips graph calculation to obtain neighbourhoods.
- Distance threshold estimation controls coarseness of approximation.

## High-dimensional data

Obtaining a decomposition

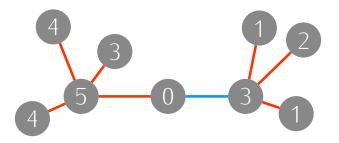


# Example Peak-seeking



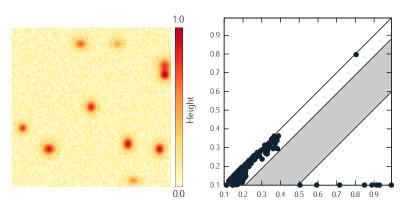
## ${\sf Example}$

Merging

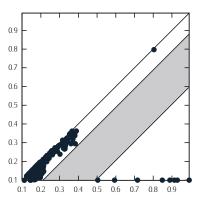


## Merge threshold estimation

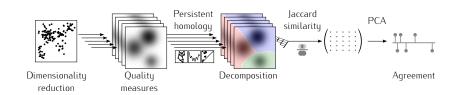
Chazal et al.: If data is sampled sufficiently dense, "relevant" peaks and topological noise are well-separated.



#### In practice

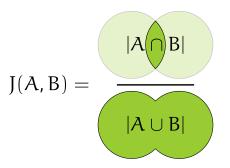


- Relate size of largest empty region to average size of empty regions.
- If ratio is large enough, consider region to be significant.

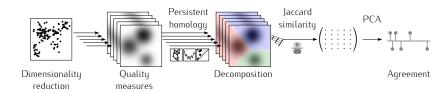


# Decomposition comparison

Jaccard index



#### Visualization



Encode local costs in scatterplot (pointwise).

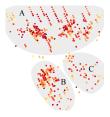
- 1030 cement mixtures with 8-dimensional feature vectors.
- Linear substructures.

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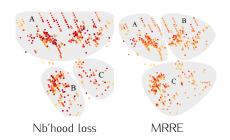




Nb'hood loss

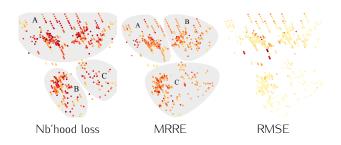
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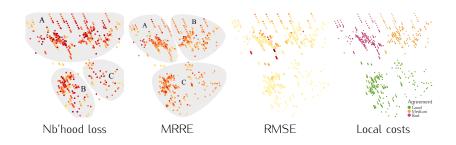
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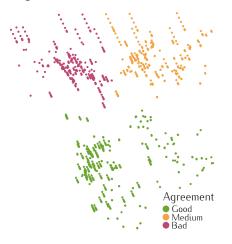




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- Upper region: Rank-based and distance-based measures disagree
- Misrepresentation possible

#### Handwritten digits

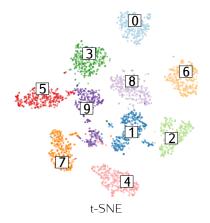
Comparing linear and non-linear dimensionality reduction

- 5620 instances of 64-dimensional feature vectors.
- Handwritten digits of different writers.

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Comparing linear and non-linear dimensionality reduction

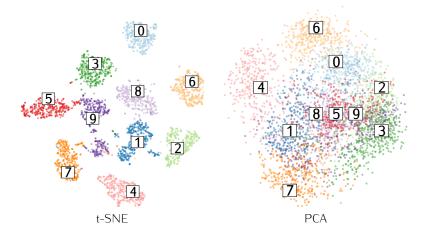
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#### Handwritten digits

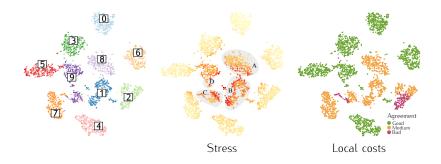
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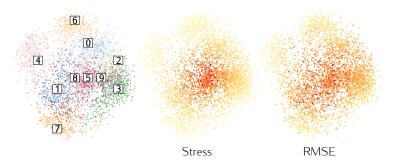


#### t-SNE





#### **PCA**

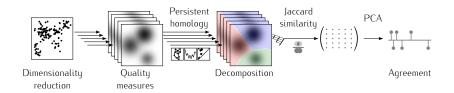


- Quality measures highlight a single persistent peak in the data.
- Peak is centred around the region of digits 5, 8, 9.

## Choosing between PCA and t-SNE

- PCA: Quality measures are "equally bad"
- t-SNE: Pronounced differences in groups
- Here: Can compare e.g. rank correlations to see that t-SNE performs better
- In general: Different ranges

#### Conclusion



- Use persistent homology to analyse the behaviour of quality measures on embeddings.
- Clustering & automatic merge threshold selection.
- Judge agreement of quality measures for dimensionality reduction.