

Dimensionality Reduction Applied to Mechanical Design and Optimization

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Introduction



Cenaero

- Private research center, specialized in numerical simulations
- Two locations
 - Non-profit organization in Charleroi (BE)
 - Subsidiary in Moissy-Cramayel (FR)
- 80+ researchers
- Started in 2002 with the support of the European Regional Development Funds and the impulse of the Walloon aeronautical sector and universities
- Operational missions: collaborative research in close collaboration with industry and academia + services for companies (HPC supercomputing facilities, consultancy)







Research Themes at Cenaero





Machine Learning & Optimization

 Machine Learning (ML) activities in various scientific and industrial projects



Prediction of physical fields



Detection of defects





Time series prediction

Dimensionality Reduction

 In addition to ML activities, we also develop Minamo software for optimization and parametric studies





Dimensionality reduction in simulation-based problems

- Dimensionality reduction in the case of Cenaero's activities:
 - Need to cope with high-dimensional design spaces
 - Data coming from physics-based simulations
 - Application: engineering design and optimization (fields: aeronautics, biomedical, buildings, ...)









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- Context of this research:
 - WINGS project funded by the Walloon Region, with HPC resources from Lucia Tier 1 supercomputer (hosted by Cenaero and also funded by the Walloon Region)
 - Goal: develop dimensionality reduction methods for mechanical design examples characterized by highly flexible parametrizations

Dimensionality Reduction



Motivation for dimensionality reduction

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Text categorization of massive document resources Acoustic signal compression Data visualization of the internet network map

- Dimensionality reduction: popular in various application fields
- Black-box simulations ⇒ no analytical properties can be *a priori* assumed
- Restricted to continuous features (no integer, discrete, or categorical var.)



 $\mathbf{x} = [x_1 \dots x_N]^{\mathsf{T}}$

Input space (e.g. CAD or FFD parameters)



$$\mathbf{y} = [y_1 \dots y_M]^\mathsf{T}$$

Output space (e.g. scalar responses extracted from velocity field, stresses, ...)











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Three main paradigms:



- Generally speaking, data management issues involve:
 - Size (= number of samples)
 - Complexity (e.g., non-linearities, singularities, ...)
 - Dimensionality (= number of features/parameters/variables/inputs)
- To address these issues, several dimensionality reduction techniques have been proposed to:
 - Transform the original dataset into a new dataset representing low dimensionality...
 - ... while maintaining as much as possible the original meanings of the data [Zebari *et al.*, 2020]
 - Their performances vary in the way they can capture or (not) the inner complexity of the data



Dimensionality reduction: benefits expected

• Benefits expected:

- Reduction of the computational time and complexity of the subsequent approximation and optimization process
- Avoids overfitting and noise
- Better data visualization and interpretation

Active area of research

- Lots of papers recently published
 - In various fields (engineering, biotech, finance, etc.)
 - For several applications (image and text processing, data visualization, etc.)

Historical trend of research

- First methods focus on global projections (typically: preserve covariance between dimensions, like Principal Component Analysis or PCA)
- Now: more localized approaches (i.e. preserve pairwise distances or proximities as much as possible)



Taxonomy of dimensionality reduction techniques



Hou, C. K. J. and Behdinan, K. Dimensionality reduction in surrogate modeling: A review of combined methods. Data Science and Engineering, 7(4):402–427, 2022.



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• Visual comparison of dimensionality reduction paradigms:





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• Visual comparison of dimensionality reduction paradigms:



Focus on variable clustering:

- ⇒ The original parameter space is preserved
- ⇒ Parameters are grouped according to a given criterion, but no one is discarded



- Analysis on the state-of-the-art in variable clustering:
 - Most methods imply the resolution of a minimization problem expressed as a linear regression or matrix factorization process, coupled with additional penalty functions to deal with grouping
 - Techniques directly or remotely inspired by data clustering (k-means, ...) are frequently integrated within the search for variable groups
 - Most methods work like in a black-box mode: no visualization tool is proposed to help the user apprehending the high-dimensional space
 - In structural and multidisciplinary optimization, the methods proposed usually involve, in one way or another, a significant level of user's expertise, or imply strong physical assumptions
- Let's remind our objective: devise a generic and flexible variable clustering strategy, with visualization functionalities



 Hypothesis: variables with similar impact on the responses are grouped together





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Hypothesis: variables with similar impact on the responses are grouped together Variable clustering



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Hypothesis: variables with similar impact on the responses are grouped together Variable clustering 3



- Idea: decomposition in three distinct steps 1, 2, and 3, using the UMAP dimensionality reduction technique at step 2:
 - UMAP (Uniform Manifold Approximation and Projection, by McInnes et al., 2018) has gained popularity in recent years in various DR contexts
 - Based on fuzzy algebraic topology and Riemannian geometry, UMAP constructs a graph in high dimensions to connect the data, and then performs an optimization to find the most similar graph in lower dimensions



McInnes, L., Healy, J., Saul, N., and Großberger, L. *UMAP: Uniform Manifold Approximation and Projection*. Journal of Open Source Software, 3(29):861, 2018.



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UMAP-based variable clustering



- Various fields of applications (genetics, image processing, ...)
- Example: analyzing genotype data among 488,377 individuals in the UK, according to ethnicity [Diaz-Papkovitch et al., 2018]
- Method: dimension reduction of genomic data combining Principal Component Analysis (PCA) with UMAP to illustrate population structure in large cohorts, and capture their relationships on local and global scales
- Aims:
 - Detect correlations between genetic diseases and ethnic origins
 - Genetic anthropology



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Legend: BA, Black African; BC, Black Caribbean; BG, Bangladeshi; CHN, Chinese; IND, Indian; PK, Pakistani; WB, White British; WI, White Irish; WBC, White and Black Caribbean; WBA, White and Black African; WAA, White and Asian; AAB, Any other Asian Background; ABB, Any other Black Background; AWB, Any other White Background; AMB, Any other Mixed Background; OEG, Other ethnic group.

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- Step 0: design of experiments (dataset)
- Step 1: compute a k-neighbor connected graph in the highdimensional space



 Step 2: define a symmetric measure of similarity for each pair of data points i and j:

$$p_{ij} \coloneqq p_{j|i} + p_{i|j} - p_{j|i} \cdot p_{i|j}$$
$$p_{j|i} \coloneqq \begin{cases} \exp\left(-\frac{\left\|\mathbf{x}^{(i)} - \mathbf{x}^{(j)}\right\|_{2} - \rho_{i}}{\sigma_{i}}\right) & \text{if } \mathbf{x}^{(j)} \in \mathcal{N}_{i} \\ 0 & \text{otherwise} \end{cases}$$

• Step 3: compute a k-neighbor connected graph for the embedded points $\mathbf{y}^{(i)}$ with a similarity measure in the latent space $q_{ij} \coloneqq \frac{1}{\left(1 + a \|\mathbf{y}^{(i)} - \mathbf{y}^{(j)}\|_2^{2b}\right)}$

$$c_1 \coloneqq \sum_{i=1}^N \sum_{j=1, j \neq i}^N \left(p_{ij} \ln\left(\frac{p_{ij}}{q_{ij}}\right) + (1-p_{ij}) \ln\left(\frac{1-p_{ij}}{1-q_{ij}}\right) \right)$$

 Additional mathematical developments lead to a more tractable function to be minimized:

$$\min_{\{\mathbf{y}_i\}_{i=1}^n} c_2 \quad \coloneqq \quad \sum_{i=1}^N \sum_{j=1, j \neq i}^N \left[p_{ij} \ln \left(q_{ij} \right) + (1 - p_{ij}) \ln \left(1 - q_{ij} \right) \right]$$

- UMAP has demonstrated its effectiveness when compared to other techniques like t-SNE or LargeVis in terms of CPU time and resources
- Further modifications have been proposed in the literature (DenseMap algorithm to regularize the cost function, variant for time series, etc.)
- UMAP can be used in high dimensions (not limited to 2D/3D visualization)
- UMAP is a stochastic algorithm: randomness used both when speeding up approximation steps, and when solving hard optimization problems
- UMAP has been promoted as an efficient pre-conditioner for clustering in a few studies [Allaoui et al., 2020; Hozumi et al., 2021]



Application to design optimization



Let's go back to our original problem

- Goal: optimize highly parametrized structural and mechanical systems
- Algorithm: surrogate-based optimizer (SBO) from Minamo
- Sequential approach followed:

Design of Experiments

to evaluate the responses on a representative subset of the design space



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Dimensionality reduction to group variables by importance (using UMAP)







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Dimensionality reduction to group variables by importance (using UMAP) Surrogate-based optimization (SBO) using Minamo







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First validation: on structural test cases



• Dimensionality:

Example	Number of variables
dome	np = 696
bldg	np = 942
wing	np = 162

 Responses: outputs from a finite element analysis [FEAP program, by Taylor *et al.*, 2008]



-10

10

dome

-10 -5

0 5

10.0 7.5 5.0 2.5 0.0

-2.5

-5.0

10

 $\max(\mathbf{x})$

 $\sigma^{\max, \text{ external load}}(\mathbf{x})$

 $d^{\max, \text{ external load}}(\mathbf{x})$

 $s^{\max, \text{ external load}}(\mathbf{x})$

 $\sigma^{\rm max,\ dead\ load}({\bf x})$

 $d^{\max, \text{ dead load}}(\mathbf{x})$

Dimensionality Reduction (14/03/2025)

bldg

Is the UMAP step really useful?

- **Direct clustering vs. UMAP-preconditioned clustering**
 - Variable impacts on responses estimated by Ridge linear regression coefficients, clustering performed by DBSCAN

Average number of clusters found	Direct clustering	UMAP-based clustering
bldg	1	7
dome	2	4
wing	1	3

Illustration on the skyscraper test case (bldg):







UMAP-based: 6 clusters found

Direct: 1 cluster found **DBSCAN** without UMAP-preconditioning fails to identify meaningful clusters of variables (caveat: no hyperparameter tuning)

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- Industrial case: structural optimization of the Dutch Maritime Museum [Descamps et al., 2014 – ULB / Princeton University / Ney & Partners]
 - 1954 design variables (cross-section areas)
 - Objective: mass Constraints: on stresses, buckling, and displacements





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66 clusters detected by UMAP + DBSCAN



SBO results



Filomeno Coelho, R., Sainvitu, C., and Benamara, T. *UMAP-based Dimensionality Reduction for Variable Grouping: Application to the Design Optimization of Truss Structures*. In 13th ASMO UK / 2nd ASMO-EUROPE / ISSMO Conference on Engineering Design Optimization Product and Process Improvement, Cenaero, Belgium, July 8-9, 2024.



SBO results



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• Take aways from these first results:

- Strategy successfully tested on cases up to 1954 design variables, but on structural design cases only with symmetry / repeatibility patterns
- What about different mechanical systems like compressors?
- Should the same variable grouping be used during the whole SBO?

Multi-stage compressor (Safran Aero Boosters)

- Number of input parameters:
 - N = 144 design variables
 - Simulation: axisymmetric throughflow CFD solver, widely used at the preliminary design stage in turbomachinery [Simon, 2007]
 - Rate of successful simulations: ~55%







Cenae



Nigro, R., Baert, L., Nyssen, F., de Cazenove, J., Dominique, J., Lepot, I., Veglio, M., and Princivalle, R. *Multi-fidelity aeromechanical design framework for high flow speed multistage axial compressors*. In: Proceedings of the ASME TurboExpo 2024 conference, London, UK, June 23-28, 2024.

Multi-stage compressor (Safran Aero Boosters)

Preliminary results on a multi-stage compressor (SAB)

- Design of Experiments of 5 N = 720 geometries (LHS Latin Hypercube Sampling)
- Dimensionality reduction performed on 1175 responses
- Agglomerative clustering better adapted to this configuration



Color:

- If "flow_path": blue
- If "aero": red
- If "geom": green
- Otherwise: yellow

Marker:

- S1: v
- S2: ^
- S3: <
- ST: >
- R2: +
- R3: x

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Idea: update the clustering of variables during the SBO

• Caution: when switching from a "coarse-grained" to a "fine-grained" variable clustering, there must be a mapping between levels, in order to re-use points from previous SBO iterations:



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Multi-stage compressor (Safran Aero Boosters)

Preliminary results on a multi-stage compressor (SAB)

- SBO: 1 objective (minimize fuel burn) and 122 aerodynamical constraints
- Similar performances obtained, but with a decrease of ~50% CPU time required (faster training of the surrogates due to reduced dimensionalities)



Conclusions and future prospects



- Design of Experiments
 - Size of the design of experiments: compromise between accuracy of the results and cost of the simulations

• Variable importances:

- Ridge regression: computationally inexpensive and often acceptable [Grömping, 2009], but might be insufficient in some cases
- Currently investigated: Random forests (RF), Mutual information (MI) scores [Université de Namur]
- Grouping dependent on active constraints only?

UMAP for variable clustering

- UMAP useful to pre-condition data for clustering
- Strategy efficiently combined to surrogate-based optimization
- Ongoing developments on active learning strategies



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