




# 1 pudu: A Python library for agnostic feature selection 2 and explainability of Machine Learning spectroscopic 3 problems.

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DOI: [10.xxxxxx/draft](https://doi.org/10.xxxxxx/draft)

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Submitted: 01 January 1970

Published: unpublished

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## 9 Statement of need

10 Spectroscopic techniques (e.g. Raman, photoluminescence, reflectance, transmittance, X-ray  
11 fluorescence) are an important and widely used resource in different fields of science, such  
12 as photovoltaics ([Fonoll-Rubio et al., 2022](#)) ([Grau-Luque et al., 2021](#)), cancer ([Bellisola &  
13 Sorio, 2012](#)), superconductors ([Fischer et al., 2007](#)), polymers ([Easton et al., 2020](#)), corrosion  
14 ([Haruna et al., 2023](#)), forensics ([P. V. Bhatt & Rawtani, 2023](#)), and environmental sciences  
15 ([Estefany et al., 2023](#)), to name just a few. This is due to the versatile, non-destructive and  
16 fast acquisition nature that provides a wide range of material properties, such as composition,  
17 morphology, molecular structure, optical and electronic properties. As such, machine learning  
18 (ML) has been used to analyze spectral data for several years, elucidating their vast complexity,  
19 and uncovering further potential on the information contained within them ([Goodacre, 2003](#))  
20 ([Luo et al., 2022](#)). Unfortunately, most of these ML analyses lack further interpretation of  
21 the derived results due to the complex nature of such algorithms. In this regard, interpreting  
22 the results of ML algorithms has become an increasingly important topic, as concerns about  
23 the lack of interpretability of these models have grown ([Burkart & Huber, 2021](#)). In natural  
24 sciences (like materials, physical, chemistry, etc.), as ML becomes more common, this concern  
25 has gained especial interest, since results obtained from ML analyses may lack scientific value  
26 if they cannot be properly interpreted, which can affect scientific consistency and strongly  
27 diminish the significance and confidence in the results, particularly when tackling scientific  
28 problems ([Roscher et al., 2020](#)).

29 Even though there are methods and libraries available for explaining different types of algorithms  
30 such as SHAP ([Lundberg et al., 2017](#)), LIME ([Ribeiro et al., 2016](#)), or GradCAM ([Selvaraju  
31 et al., 2017](#)), they can be difficult to interpret or understand even for data scientists, leading  
32 to problems such as miss-interpretation, miss-use and over-trust ([Kaur et al., n.d.](#)). Adding  
33 this to other human-related issues ([Krishnâ1 et al., 2022](#)), researchers with expertise in  
34 natural sciences with little or no data science background may face further issues when using  
35 such methodologies ([Zhong et al., 2022](#)). Furthermore, these types of libraries normally  
36 aim for problems composed of image, text, or tabular data, which cannot be associated in  
37 a straightforward way with spectroscopic data. On the other hand, time series (TS) data  
38 shares similarities with spectroscopy, and while still having specific needs and differences, TS  
39 dedicated tools can be a better approach. Unfortunately, despite the existence of several  
40 libraries that aim to explain models for TS with the potential to be applied to spectroscopic  
41 data, they are mostly designed for a specialized audience, and many are model-specific ([Rojat  
42 et al., 2021](#)). Furthermore, spectral data typically manifests as an array of peaks that are  
43 typically overlapped and can be distinguished by their shape, intensity, and position. Minor

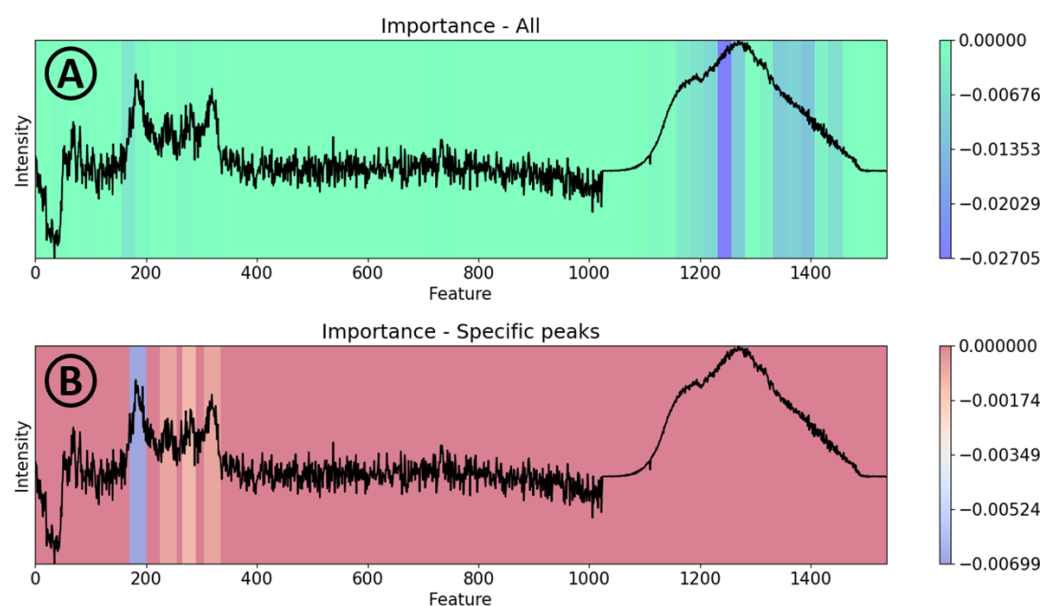
shifts in these patterns can indicate significant alterations in the fundamental properties of the subject material. Conversely, pronounced variations in the other case might only indicate negligible differences. Therefore, comprehending such alterations and their implications is paramount. This is still true with ML spectroscopic analysis where the spectral variations are still of primary concern. In this context, a tool with an easy and understandable approach that offers spectroscopy-aimed functionalities that allow to aim for specific patterns, areas, and variations, and that is beginner and non-specialist friendly is of high interest. This can help the different stakeholders to better understand the ML models that they employ and considerably increase the transparency, comprehensibility, and scientific impact of ML results (U. Bhatt et al., 2020) (Belle & Papantonis, 2021).

## Overview

**pudu** is a Python library that quantifies the effect of changes in spectral features over the predictions of ML models and their effect to the target instances. In other words, it perturbs the features in a predictable and deliberate way and evaluates the features based on how the final prediction changes. For this, four main methods are included and defined. **Importance** quantifies the relevance of the features according to the changes in the prediction. Thus, this is measured in probability or target value difference for classification or regression problems, respectively. **Speed** quantifies how fast a prediction changes according to perturbations in the features. For this, the Importance is calculated at different perturbation levels, and a line is fitted to the obtained values and the slope, or the rate of change of Importance, is extracted as the Speed. **Synergy** indicates how features complement each other in terms of prediction change after perturbations. Finally, **Re-activations** account for the number of unit activations in a Convolutional Neural Network (CNN) that after perturbation, the value goes above the original activation criteria. The latter is only applicable for CNNs, but the rest can be applied to any other ML problem, including CNNs. To read in more detail how these techniques work, please refer to the [definitions](#) in the documentation.

**pudu** is versatile as it can analyze classification and regression algorithms for both 1- and 2-dimensional problems, offering plenty of flexibility with parameters, , and the ability to provide localized explanations by selecting specific areas of interest. To illustrate this, [Figure 1](#) shows two analysis instances using the same importance method but with different parameters. Additionally, its other functionalities are shown in examples using scikit-learn (Pedregosa et al., 2011), keras (Chollet et al., 2018), and localreg (Marholm, 2022) are found in the documentation, along with XAI methods including LIME and GradCAM.

**pudu** is built in Python 3 (Van Rossum & Drake, 2009) and uses third-party packages including numpy (Harris et al., 2020), matplotlib (Caswell et al., 2021), and keras. It is available in both PyPI and conda, and comes with complete documentation, including quick start, examples, and contribution guidelines. Source code and documentation are available in <https://github.com/pudu-py/pudu>.



**Figure 1:** Two ways of using the same method *importance* by A) using a sequential change pattern over all the spectral features and B) selecting peaks of interest. In A), the impact of the peak in the range of 1200-1400 opaques the impact of the rest. In contrast, in B) only the first four main peaks are selected to be analyzed and better visualize their impact in the prediction.

## Acknowledgements

Co-funded by the European Union (GA No. 101058459 Platform-ZERO). Views and opinions expressed are however those of the author(s) only and do not necessarily reflect those of the European Union or European Health and Digital Executive Agency (HADEA). Neither the European Union nor the granting authority can be held responsible for them. This project has received funding from the European Union's Horizon 2020 research and innovation programme under Marie Skłodowska-Curie GA No. 801342 (Tecniospring INDUSTRY) and the Government of Catalonia's Agency for Business Competitiveness (ACCIÓ). Authors from IREC belong to the MNT-Solar Consolidated Research Group of the "Generalitat de Catalunya" (ref. 2021 SGR 01286) and are grateful to European Regional Development Funds (ERDF, FEDER Programa Competitivitat de Catalunya 2007-2013).

## Authors contribution with CRediT

- Enric Grau-Luque: Conceptualization, Data curation, Software, Writing – original draft
- Ignacio Becerril-Romero: Investigation, Methodology, Writing – review & edition
- Alejandro Perez-Rodriguez: Funding acquisition, Project administration, Resources, Supervision
- Maxim Guc: Formal analysis, Validation, Methodology, Writing – review & edition
- Victor Izquierdo-Roca: Funding acquisition, Project administration, Supervision

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