

# Relative Structural Analysis on Molecular Perovskite

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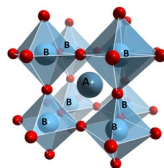
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# 1

## What are Perovskite compounds?

Compounds with  $ABX_3$ , have small cation B-site, large cation A-site and a bridging ligand anion X-site.



## How can we compare from one another?

One way is to use Descriptors: transforms a structure into a multi-dimensional constant sized vector. Smooth Overlap of Atomic Positions (SOAP) is one example from Dscribe.

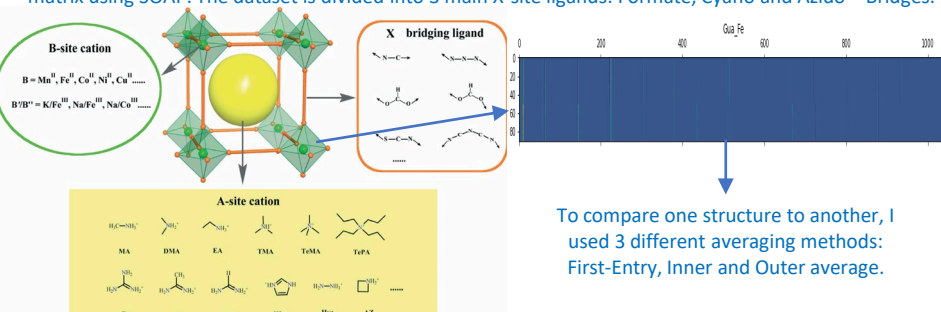
## How can you use ML to analyse perovskite compounds?

You can use unsupervised learning. Through the use of clustering algorithms, we can find hidden clusters/structure between different crystal structures of the same type.

# 2

## Dataset

Gathered various structures from a perovskite compound paper [1] and turned each structure into a matrix using SOAP. The dataset is divided into 3 main X-site ligands: Formate, Cyano and Azido – Bridges.



[1] Structural phase transitions in perovskite compounds based on diatomic or multiatomic bridges, Wei-Jian Xu, a Zi-Yi Du, b Wei-Xiong Zhang\* a and Xiao-Ming Chen, a <https://doi.org/10.1039/C6CE01485B>

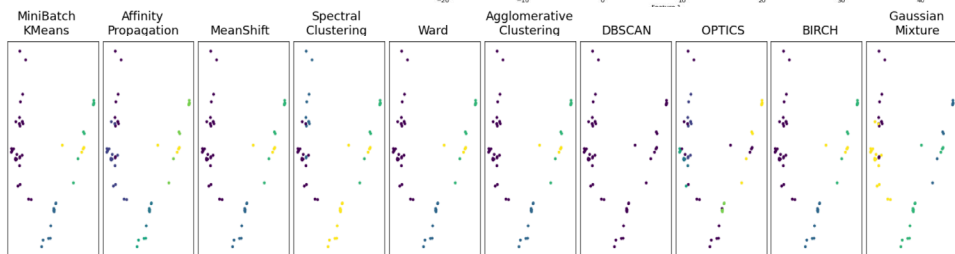
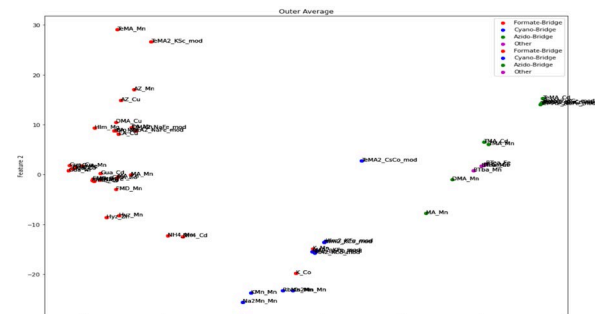
# 3

## Graphical Representation

Using PCA and Standard Scalar from scikit, I could plot my dataset.

## Clustering Algorithm

Clustering algorithm can handle multi-dimensional vector datasets.



## Optimisation

There are certain ways to measure performance of cluster models.

