

# Chemical Space Exploration via Semi-Supervised Learning with Molecular Graphs

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Computational Materials Science & Chemistry

### **Problem Characterization**



- Chemical Space exploration is a recurrent activity in MS
- Generative ML methods to fulfill this purpose are emerging, such as VAEs
- VAEs' latent spaces' dimensions do not follow properties
- Current graph-based VAEs have big limitations and could be much better





- Graph-based VAE with navigable latent space, emphasizing properties
- Find a cost function for the VAE that corroborates with the objective
- Include disentanglement procedures to make navigation even better
- Project an elastic model for the iterative ampliation of the latent space
- Test the developed technology using chemical datasets of interest



## Methods - GNN and Message Passing

- Graphs encode molecules well and store more information than typical string representations.
- GNN receives graphs as input and encodes the vertices and edges as feature vectors.
- MP algorithm is applied so vertices are informed about their neighbors



Figure 1: Message Passing

representation <sup>4</sup>

### Methods - Variational Autoencoder



Figure 2: Variational Autoencoder

- Autoencoders, by default, work like the identity function
- It is possible to sample data from the latent space of a VAE
- Disentanglement can be applied to a VAE to force different neurons to learn different properties

#### Methods - Semi-Supervised Autoencoder





Figure 4: GVAE vs SGVAE (Oliveira, A. F., Da Silva, J. L., Quiles, M. G., 2022)

Figure 3: Semi-Supervised

Autoencoder topology

 It is possible to combine a predictor and a VAE to further organize the latent space 6



# Methods - Grammar AE vs Graph AE

Grammar Autoencoder encoding Alanine represented by SMILES.



Graph Autoencoder encoding Alanine represented by 3D Graph.

## **Expected Results**



- Semi-supervised learning system capable of finding useful chemical compounds through chemical space exploration
- Reduction on materials screening time (one of the most time-consuming tasks), improving research productivity





- Encoding large molecules
- Directing properties
- Configuring hyper-parameters
- Obtaining useful molecules

#### Acknowledgments

