

Motivation

- In modern world ability to rapidly iterate through the scientific discovery process is important
- The biggest bottleneck in material science is the time-consuming process of creating and combining new substances
- We need to strategically prioritize candidate molecules for synthesis

Objectives

- Simplify the entire discovery process, spanning from proposing molecules to ranking them effectively.
- Integrate established heuristics utilized by scientists into the prediction framework.
- Create a user-friendly web application tailored for material scientists, designed to boost research efficiency.

Solution

- Build comprehensive end-to-end solution that generates new molecules and predicts the target capacity of CTFs based solely on the encoded representation of the molecular substrates in Simplified Molecular Input Line Entry System (SMILES)

Results

- No signal in the data we received from our experts --> random ranking
- With LightGBM variant and the combined dataset (our data + data from literature with already extracted features), **74% of pairwise comparisons resulted in a correct ranking**

Future Work

- Obtain more ground truth data containing SMILES
- Create more flexible molecule generation method using GNNs

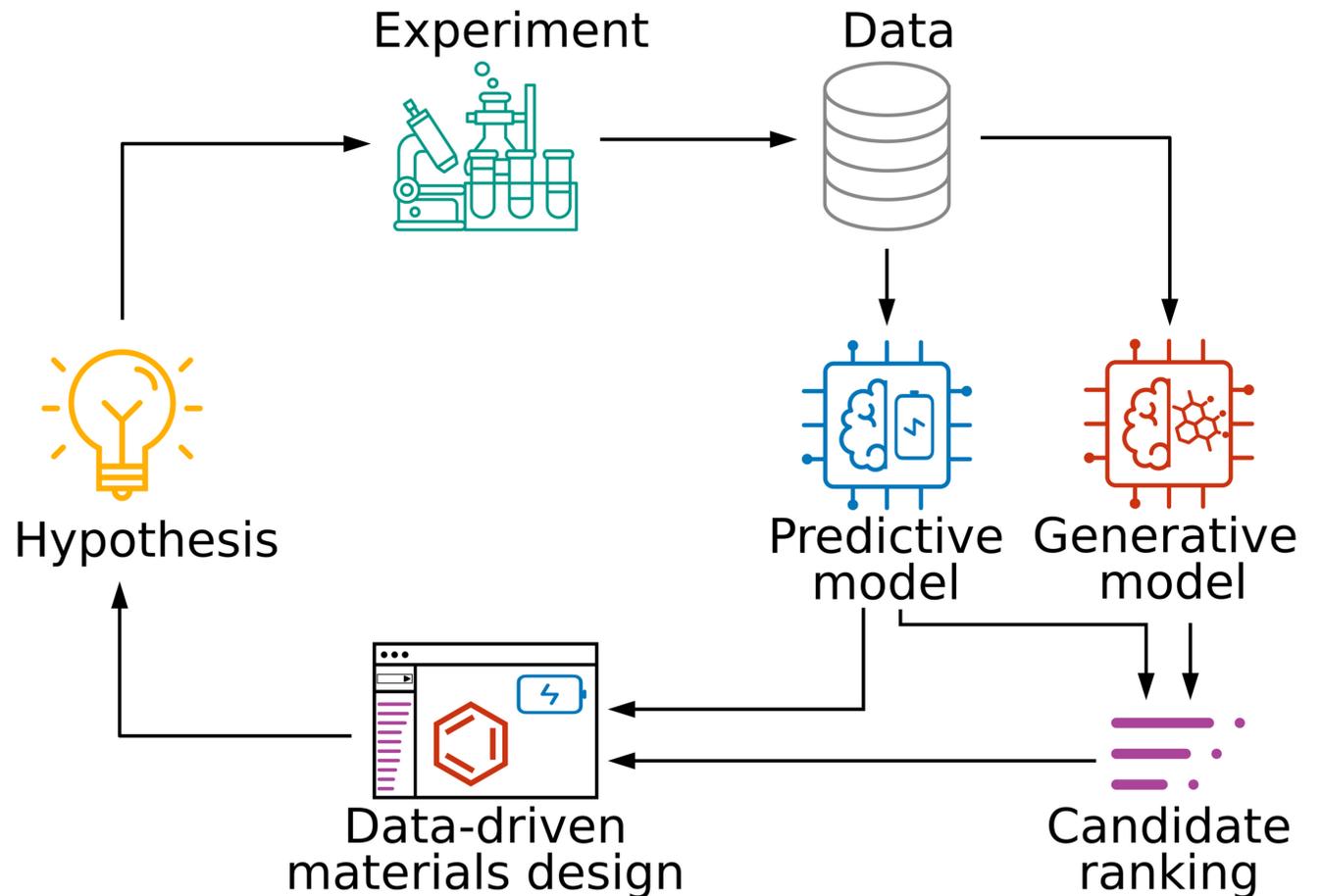
Acknowledgments

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References

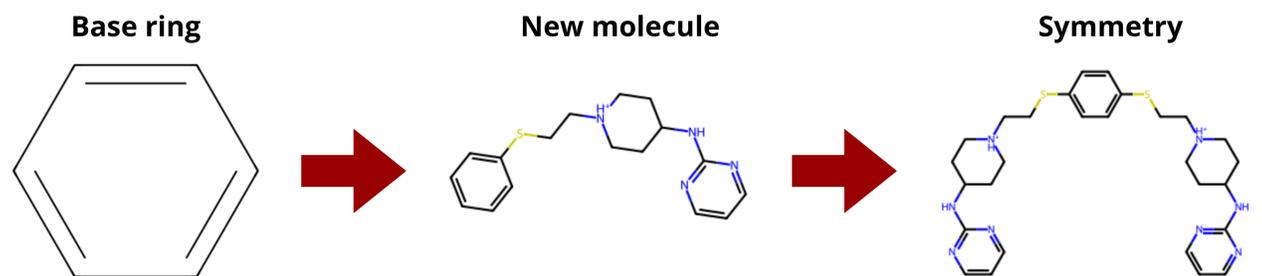
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2. Ishida et al., "Chemtsv2: Functional molecular design using de novo molecule generator"
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Framework

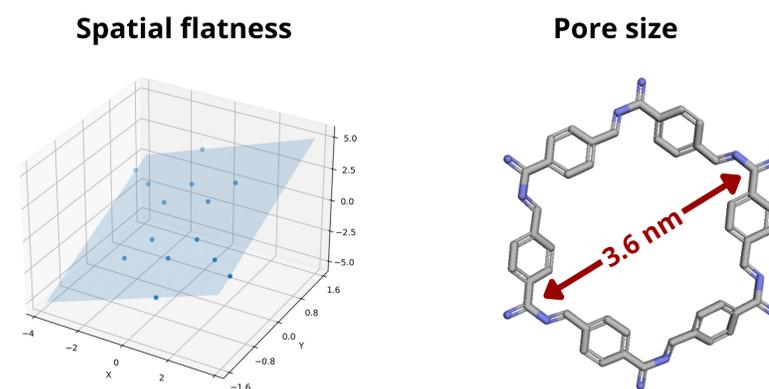


Methods

Generation



Feature engineering



CTF capacity prediction

