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## Deep Learning – the Death of Physics-Based Kernel Representations?



Physics-based representations for machine learning properties of chemical reactions<sup>[1]</sup>

- Reaction representations are constructed using conventional QML representations, accounting for the nuclear charge and coordinates of the species involved in the reaction.
- A new reaction representation (B<sup>2</sup>R<sup>2</sup>) is created by investigating the traditional representation ingredients relevant to high-performing reactions.

Method:

• Reactant-only, product-only, or combined (addition or difference) representations of both can be used to represent a chemical reaction.

$$X_{reactants(products)} = \sum_{i(j)=1}^{N(M)} X^{i(j)} \quad X_{difference} \approx X_{TS} = \sum_{j=1}^{M} X^{products} - \sum_{i=1}^{N} X^{reactants}$$

- Analysis of diverse datasets showed that combined representations perform better.
- The analysis of the key factors that contribute to the robustness of SLATM<sub>d</sub>, namely its meaningful difference, emphasis on two-body interactions, and separation of relevant two-body feature bags, is used to engineer the B<sup>2</sup>R<sup>2</sup> representation.
- B<sup>2</sup>R<sup>2</sup> is based on the notion of differences in pairwise interactions between reactants and products;
- Three bagging strategies are used by B<sup>2</sup>R<sup>2</sup>: canonical (pairwise bags), linear (elemental bags), and constant-sized (no bags).

Acceptable accuracy is achieved when the B<sup>2</sup>R<sup>2</sup> representation is used as input for KRR models when predicting reaction properties.

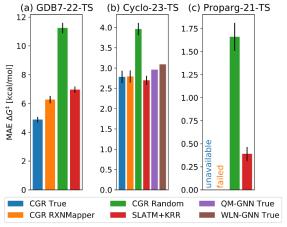
- <u>Model performance</u>: For quantitative activation energy analysis, a model with high test MAE (*ca.* 10 kcal/mol) is useless. Using model baselines to compare performance is essential.
- <u>Model input</u>: For reaction energies, the output is computed at no extra cost because the model uses DFT-optimised geometries. Methods cheaper than DFT must be used to incorporate 3D information (only if really needed).
- <u>Data splitting</u>: Training models on random splitting can deduce irrelevant patterns, hindering generalization. Using different splits (including chemically relevant ones) can help determine applicability and compare this model to others.

Splitting	SLATM <sub>d</sub> <sup>(2)</sup>	DMPNN
Random	10.01±0.16	4.11±0.07
Scaffold	13.32±0.73	19.08±0.94

MAEs (kcal/mol) are reported for 10,165 training points of GDB7-20-TS dataset.

## Reply to critical comments<sup>[3]</sup>

- Model input and performance: Deep NN models use atom-mapped reaction
  SMILES, and the mapping method (handmade or algorithmically evaluated) greatly affects model performance. Both "model types" have advantages and disadvantages, and different datasets present different challenges.
- <u>Data splitting</u>: All dataset reactions appear in the test set in one of CV folds. Community consensus on best practices is lacking.



[2] Mach. Learn.: Sci. Technol. 2023, 4, 048001.

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[1] Mach. Learn.: Sci. Technol. 2022, 3, 045005.