Predicting scalar coupling constant using dipole moment and magnetic shielding tensor values

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Abstract

This report explains the usage of dipole moments and magnetic tensor values in the calculation of scalar coupling constant using machine learning models and outlines a brief comparative analysis between the models that use these values and those that do not.

5 1 Introduction

- 6 NMR (Nuclear Magnetic Resonance) spectroscopy is used to investigate the chemical structures of
- 7 an unknown molecule in solution. Scalar coupling constant (SCC) describes the interaction between
- 8 two magnetic nuclei in NMR spectroscopy.
- 9 The value of SCC varies with the type of coupled atoms and the number of bonds between the coupled
- 10 atoms. Thus, there is a need to accurately predict SCC values
- The current project aims to do a comparative analysis between the models that utilise the dipole
- moment and magnetic shielding tensor values to understand the importance of these values on the
- 13 accuracy of prediction of SCCs.

14 2 Dataset

- 15 The dataset for this is similar to the dataset that was used in the primary reference paper(Fang, 2021).
- 16 This dataset is from kaggle: https://www.kaggle.com/c/champs-scalar-coupling
- 17 The dataset contains molecule name, atom indices, scalar coupling constant values, dipole moments
- values, magnetic shielding tensor values, mulliken charge values and potential energy values.

9 3 Literature survey

- 20 The Fang 2021 paper mentions MPNN and transformer based architecture as some of the most
- 21 striking solutions to calculate SCC. Our project compares these models with the other regression
- 22 based models
- 23 The dipole moments and magnetic shielding tensor values are not just directly incorporated into the
- 24 training of the architecture. They need to be not only pre-processed but also need to be included in
- 25 the algorithm in specific ways. The project also outlines briefly the information regarding these ways
- 26 of incorporation.
- 27 The architectures have also used a wide variety of methods such as utilising SchNet architectures,
- dual MPNN, XgBoost, Random Forest, Regression and many more.

29 4 Baseline model(s)

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Multiple baseline models have been run for the comparative analysis. Primarily, there were three 30 types of models that were run as a baseline to understand the different kinds of architectures. The 31 first type is the models that do not utilise any chemistry information at all and purely go by black box 32 machine learning methodologies of train and test data. The second type are the models that utilise 33 some of the chemistry information given such as the structural data or dipole moments and utilise 34 standard machine learning algorithms to predict the SCCs. The theird type is the models that utilise 35 an encoder-decoder architecture with message parsing layers along with chemistry information like 36 structures to obtain a prediction of SCCs. 37

In the first type, https://www.kaggle.com/code/inversion/atomic-distance-benchmark/notebook utilises a random forest regressor model on the training data and then the model is used on the test data to obtain the predictions. The results that have been obtained by this architecture are, as expected low in accuracy due to the black box approach to the problem. The scatterplot between predicted and actual values for 1JHC is given below.

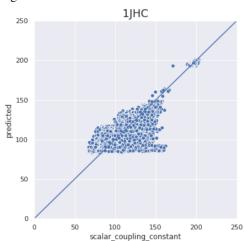
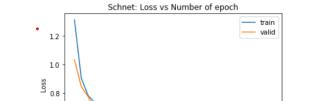


Figure 1: Predicted SCC values vs actual values

The second type of architecture includes the kind that utilises simple machine learning models by utilising some of the chemistry related data along with the test and train to get results. The example used here is a model that utilises the SchNet Architecture with train, test and structural data of the molecules to obtain the prediction of SCCs. https://www.kaggle.com/code/hau8899/schnet-1/notebook



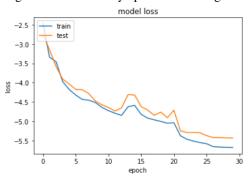
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Figure 2: Change in loss with every epoch on training and validation data

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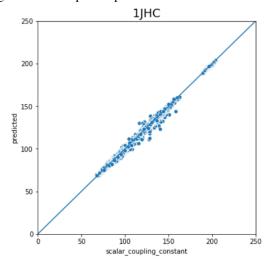
- The third type of architecture is the one that utilises message-parsing neural networks. This apprach
- provides the most accuracy when compared to the above two types of architectures. This methodology 49
- utilises edges and nodes with message parsing layers and an Adam optimiser to predict SCC values. 50
- The change in loss with every epoch is given below. 51

Figure 3: Change in loss with every epoch on training and validation data



- To get an understanding of how accurate these predictions are using MPNN architecture, a scatterplot
- of the predicted values vs actual SCC values can be plotted. This plot is for 1JGC similar to the above 53
- plot(Fig1)

Figure 4: Scatterplot of predicted vs actual values of SCC

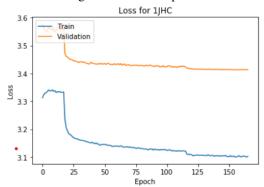


Comparitive analysis

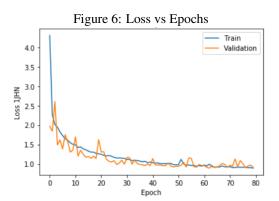
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- Another architecture utilising one hot encoding and a bidirectional RNN on dipole moment data, 56 structural data and potential energy data along with the test and train achieved a 97.7% accuracy on their validation data https://www.kaggle.com/code/manasghosal/brnn-using-additional-dataset. 58
- A Keras neural network architecture was implemented with leaky ReLU and batch nor-59 malisation utilising mulliken charges, dipole moments and magnetic shielding tensors 60 https://www.kaggle.com/code/jagannathrk/keras-neural-net-for-champs. This model was then used 61
- on the train which was split further into train and validation set for prediction. The change of loss 62 with epochs for this architecture is given below(fig5) 63
- A model developed using keras neural networks with batch normalisation and Adam optimiser with multiple dense layers and hyperparameter tuning using dipole moments, Mulliken charges, magnetic

Figure 5: Loss vs Epochs



shielding tensor values, and test and train gave a -0.80 MAE loss value. The change in loss with epochs for 1JHC is given below.(fig6)



Another similar keras NN-based model achieved a similar accuracy over 200 epochs as given below.

Mean Absolute Error

40
35
30
25
10
0 25 50 75 100 125 150 175 200

Epochs

Figure 7: MAE Loss vs Epochs

69 6 Conclusion

- The above analysis clearly indicates that the utilisation of dipole moments and magnetic shielding tensor values in the machine learning model yeilds good results. It was also observed that regardless
- of the exact model used, either a keras NN or MPNN architecture or bidirectional RNN, the accuracy
 - of these models is significantly higher than those models that are more algorithmically efficient but

- do not utilise these additional dipole moment and magnetic shielding tensor data. When it comes to
- 75 the models within those that utilise these dipole moments and tensor values, the MPNN architecture
- 76 offers the most accurate predictions with the predicted values being very close to the ground truth.

References

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