

Structure-to-Shift Prediction for ^{19}F NMR: Comparison of Graph Neural Networks, HOSE Code, and Large Language Models

Nathan O'Neill¹, Adam P Jones¹, Katharina T Huber¹, E Kate Kemsley^{1,3}, Carlos Cobas^{2,3}, David Williamson³, Gary Sharman³, John Hollerton³

1.University of East Anglia (UEA), Norwich Research Park, NR4 7TJ, Norwich, UK

2.Centro de Investigación Mestrelab (CIM), Av. de Barcelona, 7, 15706, Santiago de Compostela, Spain

3.SciY-Mestrelab, Av. de Barcelona, 7, 15706, Santiago de Compostela, Spain

Motivation and Methods

Fluorine (^{19}F) NMR is useful for structural verification of fluorinated compounds, important in the pharmaceutical and agrichemical industries. This study compared different approaches for predicting ^{19}F chemical shifts directly from molecular structure:

Graph Convolutional Neural Networks (GCNNs)

Trained on a proprietary database of approximately 15,000 structures annotated with groundtruth ^{19}F chemical shifts; uses a 'message-passing' network architecture.

Mnova's ModGraph-NMRPredict (SciY-Mestrelab)

Commercial tool that uses the same underlying database as the GCNN approach but applies empirical HOSE-code prediction.

Large Language Models (LLMs) from Anthropic and OpenAI

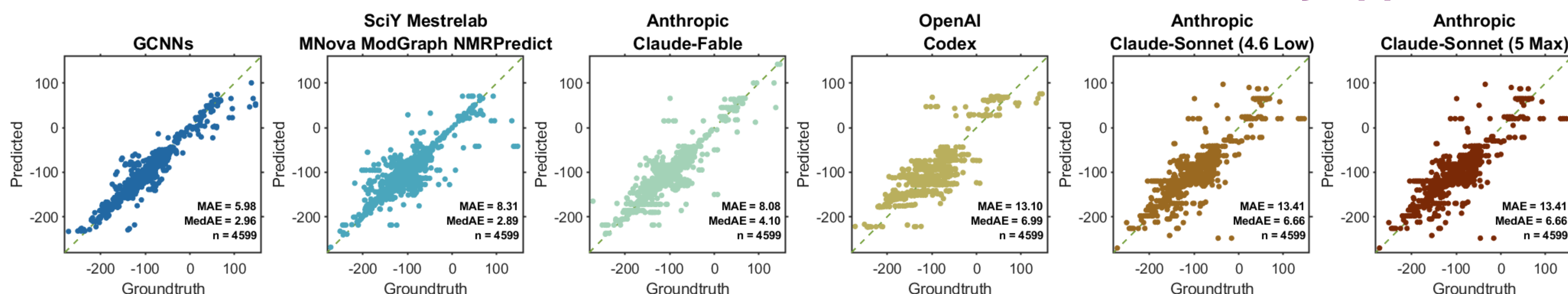
Prompted to design, develop and execute their own zero-shot prediction workflows, based on molecular environment analysis and chemical reasoning.

Test Dataset

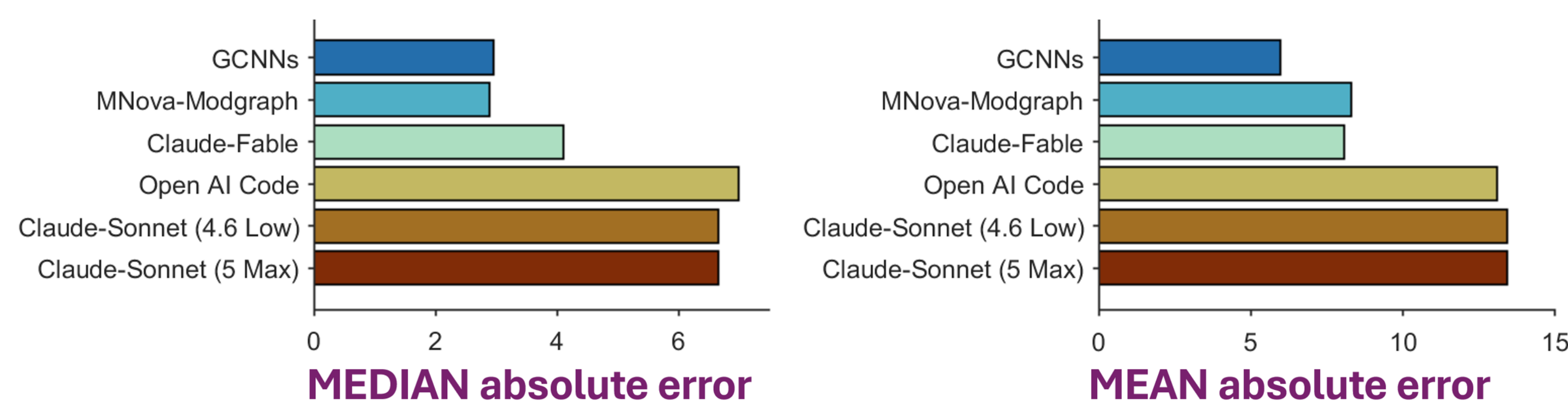
A test collection was sourced from the public *nmrshiftdb2* database, comprising 795 diverse fluorinated structures annotated with ^{19}F chemical shifts.

None of these were present in the GCNN/ModGraph training set. The LLMs were specifically instructed to refrain from search and retrieval. Thus, all methods were benchmarked exclusively on unseen compounds.

Predicted versus Groundtruth ^{19}F chemical shifts for Test set, by approach

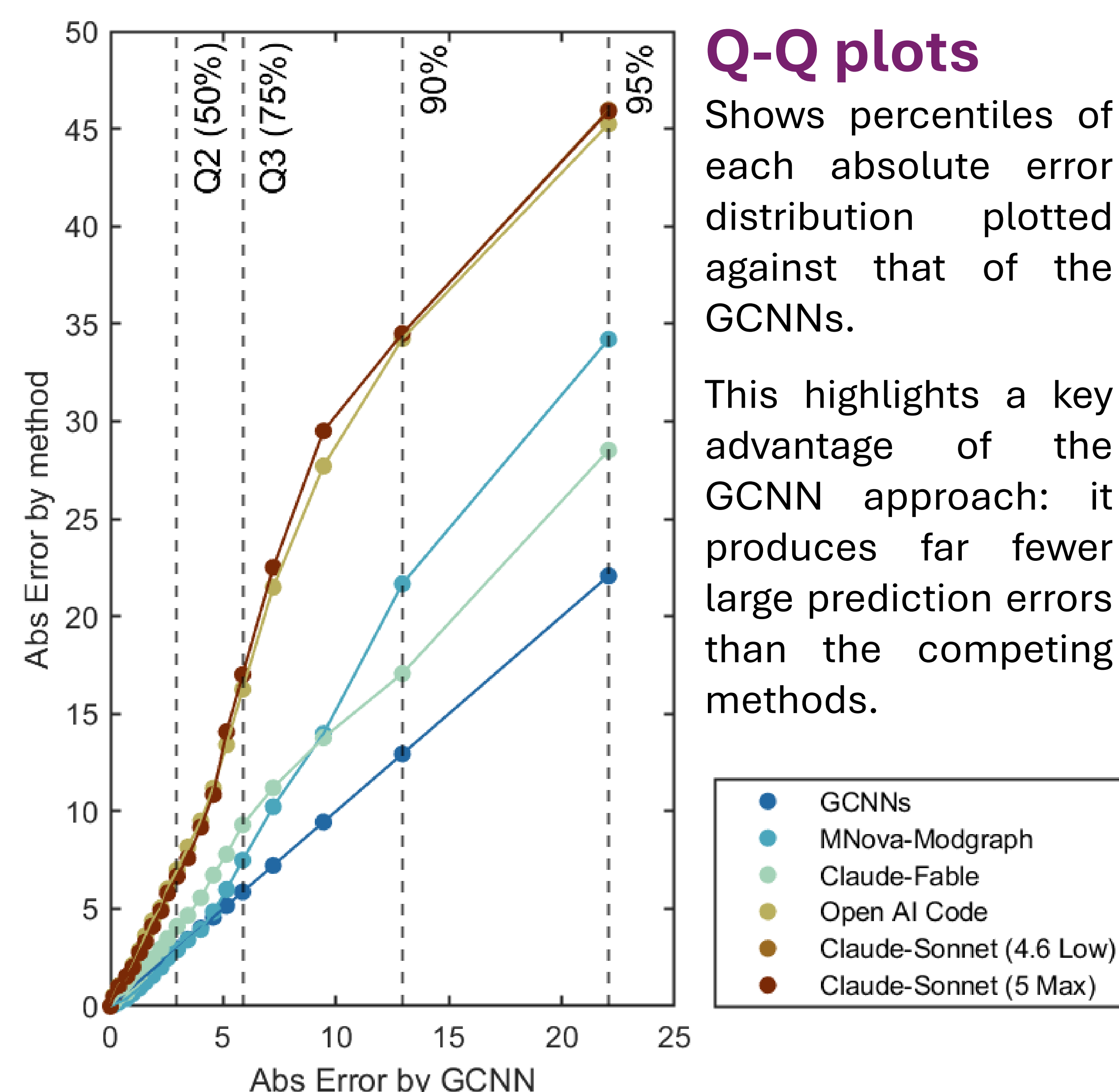


- GCNNs and ModGraph achieved similar **MEDIAN Absolute Errors** (2.96 and 2.89 ppm respectively), both outperforming the LLMs.
- GCNNs provided the best overall accuracy and generalisation to unseen compounds and achieved the lowest **MEAN Absolute Error** (5.98 ppm), >25% lower than any other approach.



Conclusion

Although modern LLMs demonstrate impressive chemical reasoning capabilities, for quantitative ^{19}F chemical shift prediction they currently do not match the performance of dedicated NMR models trained on high-quality experimental data.



Acknowledgments



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