

NMR Prediction, Calculation and Molecular Interpretation Complexity

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Nuclear Magnetic Resonance spectroscopy is a central tool for molecular structure elucidation. While the forward problem of predicting chemical shifts from a known molecular structure is well studied, the inverse question remains insufficiently quantified.^{1,2} To what extent does a given set of ¹H and ¹³C chemical shifts constrain the underlying molecular graph? The objective of this study is to quantify the structural information content and intrinsic ambiguity of experimental NMR data using curated entries from NMR chemical shifts databases.³ Atomic environments are represented through HOSE descriptors, enabling systematic analysis of how defined ppm intervals correspond to distinct local structural motifs. By evaluating this mapping across multiple spectral resolutions and structural radii, we assess how uniquely NMR chemical shifts specify atomic environments and how this specificity evolves with increasing structural context.

The analysis presents both methodological and theoretical challenges. Public NMR datasets contain duplicate records, inconsistent assignments, solvent-dependent variations and outliers that can artificially inflate structural diversity and bias ambiguity estimates. A controlled curation and validation workflow was therefore implemented, complemented by density functional theory (DFT) and density functional tight binding (DFTB) calculations to assess assignment consistency and evaluate theoretical chemical shift variability. Structurally, substantial overlap exists among chemical shift regions, rendering the inverse mapping inherently non-unique. The poster will therefore present quantitative mappings between ppm intervals and distinct HOSE-based environments, entropy-based measures of ambiguity across descriptor radii, sensitivity analyses evaluating the impact of data curation and spectral binning, and molecular-level complexity estimates derived from branching formulations and co-occurrence statistics.

Bibliography :

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