



# 6<sup>TH</sup> EUROCC VILNIUS WORKSHOP ON USING HPC



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# HIDRA – a Hierarchical DFT accuracy transformer model to Reconstruct Atomic geometry

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Predicting molecular properties from string representations enables high-throughput screening without expensive 3D structure generation. We present a systematic comparison of SMILES [1] and SELFIES [2] representations using transformer encoders [3] trained on 3.9 million DFT-optimized molecules from the PubChemQC database, predicting seven structural and symmetry properties.

This work addresses three questions: which string representation better encodes molecular structure for property prediction, how transformer depth should scale given limited quantum chemistry data, and whether multi-task learning improves performance through shared representations. We evaluate three architectural variants shown in Fig. 1: single-task models with one property per encoder, parallel multi-task with all prediction heads at the final layer, and hierarchical multi-task with heads at different encoder depths and cross-attention feedback between properties.

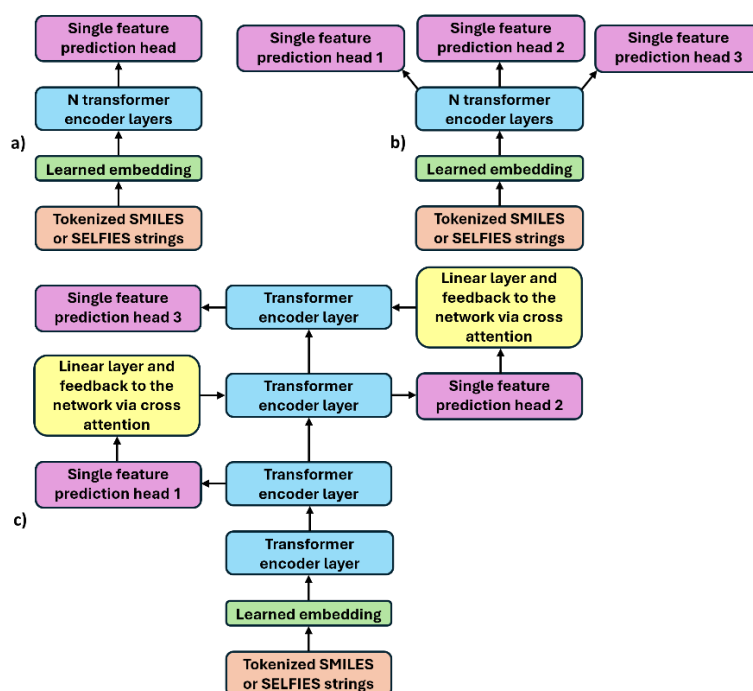


Fig. 1. (a) Single-property, (b) parallel multi-property, and (c) hierarchical multi-property transformer encoder networks.

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