

Predicting DNA Binding Proteins with Machine Learning and Deep Learning
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1. Ensure 'Dataset' folder is on the same layer as 'Code' folder so that the path will be correct when reading in the dataset.
2. 'Code' folder contains 4 python notebooks (1 for each model) and `feature_extraction.py`
3. `Feature_extraction.py` contains the common functions used for each model.
4. Run each of the python notebooks by clicking 'Run all'.