Using deep neural networks with heterogeneous chemical data to support phenotypic assay campaigns

Antonio de la Vega de León





Deep neural networks: challenges and opportunities

Antonio de la Vega de León



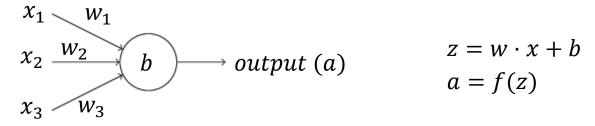


Hyperparameter selection

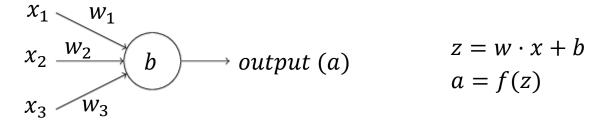
Effect of missing data

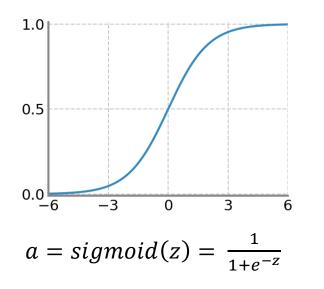






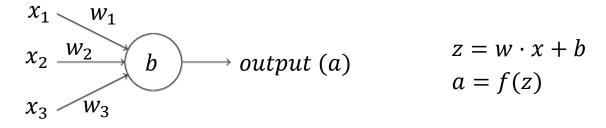


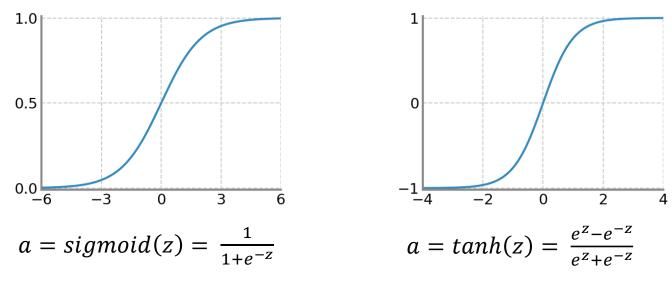






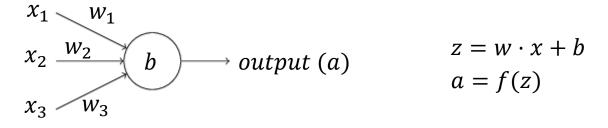


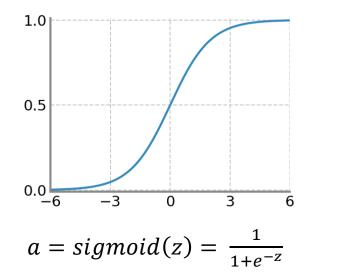


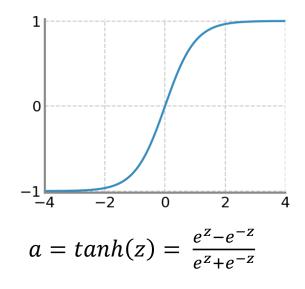


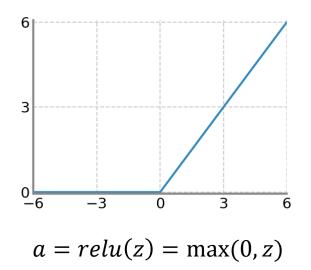










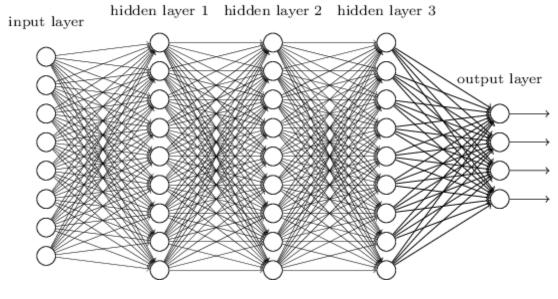






Neurons are organized into layers, where neurons in one layer are connected to all neurons in the previous and the next layer

The input layer represents the data descriptors while the output layer has a neuron per prediction task





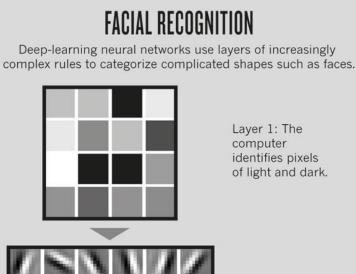


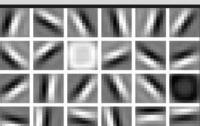
Deep learning

The idea behind deep learning is that of a hierarchical learning process

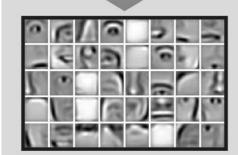
Early layers identify simple patterns that latter layers use to learn complex patterns

This reduces the need for feature selection or complex feature engineering

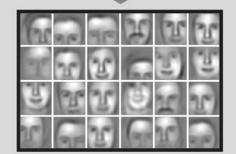




Layer 2: The computer learns to identify edges and simple shapes.



Layer 3: The computer learns to identify more complex shapes and objects.



Layer 4: The computer learns which shapes and objects can be used to define a human face.





Hyperparameter selection

Effect of missing data





Hyperparameter selection

DNNs usually require a large number of hyperparameters to be set in advance:

- Optimizer functions
- Learning rate
- Activation function
- Number of neurons
- Number of hidden layers

- Dropout amount
- Batch size
- Number of training steps
- Weight initialization

How best to find the combination that gives highest performance?

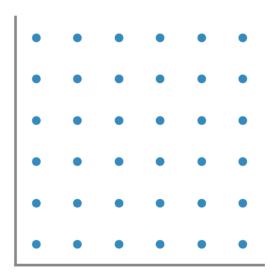




Search strategies

Grid search

- For each parameter, a set of values is fixed and all possible parameter combinations are tested
- Impractical for large number of parameters





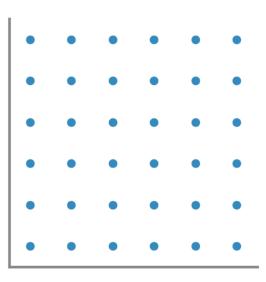
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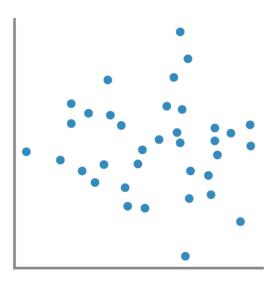
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Random search

- For each parameter, a value range is fixed and at each iterations random numbers are used
- Can require large number of iterations to cover parameter space adequately









Search strategies

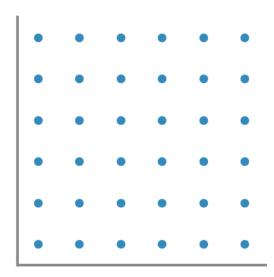
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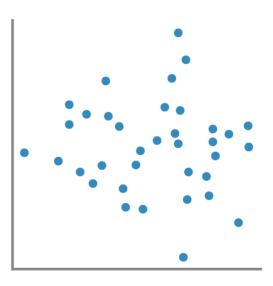
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Bayesian optimization







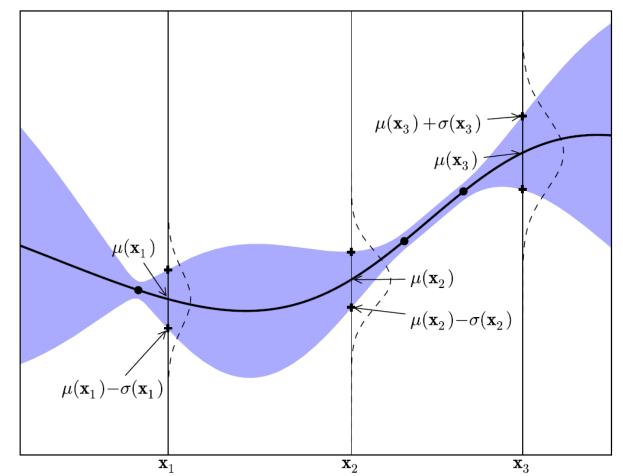


Bayesian optimization

Methodology to find the optimum of a black-box function over a range of parameter values

Based on previous evaluations, it builds a probabilistic representation of the function (Gaussian Process or GP)

Based on the GP, a utility function determines the best point in parameter space to test in the next iteration





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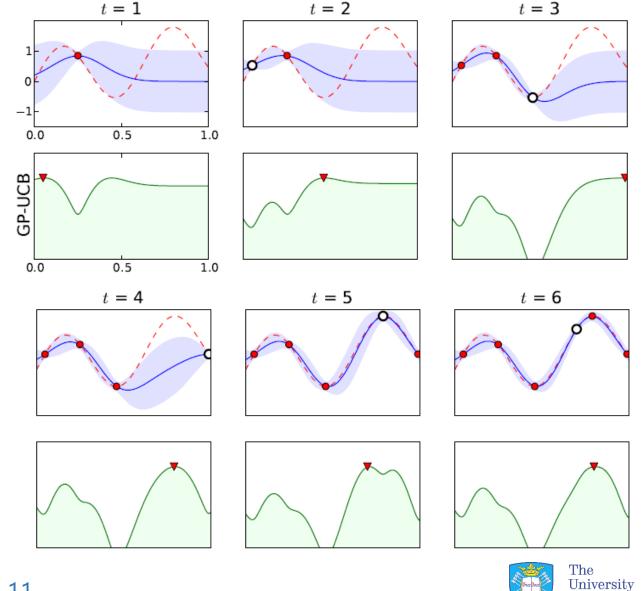
Bayesian optimization

Utility functions guide the search for the optimum:

- Probability of Improvement
- Expected Improvement
- GP-Upper Confidence Bounds

They all have a parameter that balances exploration/exploitation

At each step, the GP becomes better at representing the real formula



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Bayesian optimization test

Tox21 data: collection of 12 toxicity assays with active/inactive labels

Random parameter selection vs Bayesian optimization with the 3 functions (POI, EI, and UCB) at values: 10⁻⁷, 10⁻⁵, 10⁻³, 10⁻¹, or 10¹

200 DNN models per run were trained where the parameters varied:

- Optimizer algorithm: Adagrad, Adam, Ftrl, RMSProp, or SGD
- Learning rate: 10⁻⁷, 10⁻⁵, 10⁻⁴, 10⁻³, 10⁻², 10⁻¹, 1, 10¹, or 10³

Performance is measured as area under the ROC curve (AUC)



https://tripod.nih.gov/tox21/challenge/

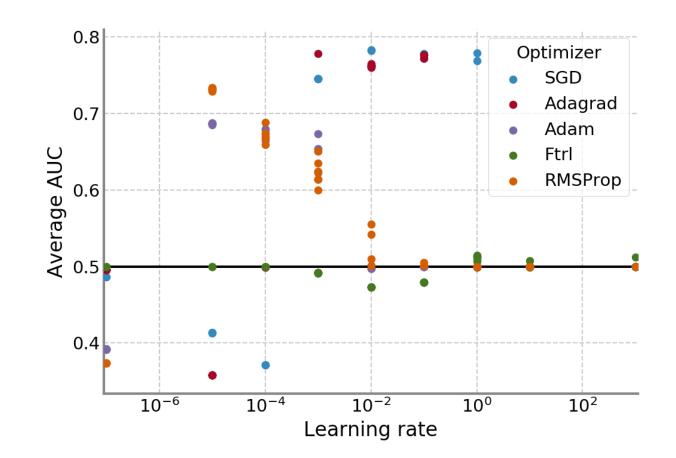


Bayesian optimization test

The effect of both of these parameters are heavily related

Many combinations of optimizer and learning rate lead to models with random or worse performance

The winner of the Tox21 data challenge achieved an average AUC value of 0.84







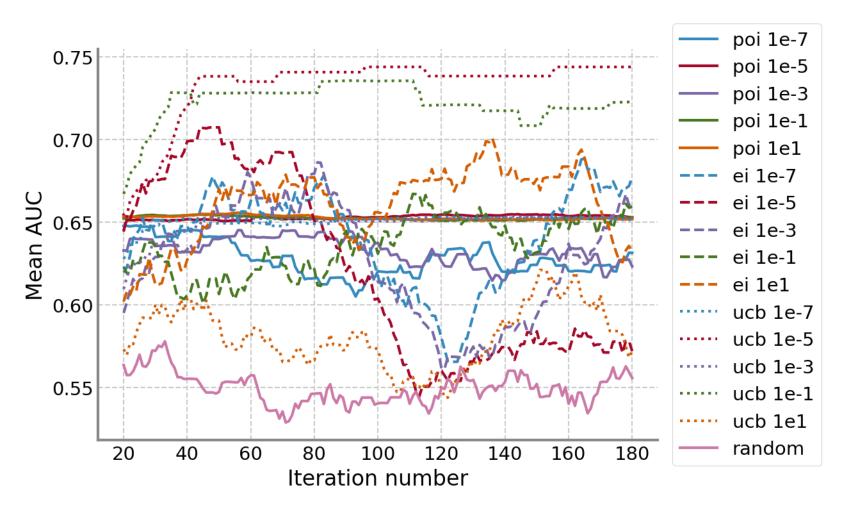
Bayesian optimization test

Random search has very low AUC values throughout

Results of Bayesian optimization are really mixed

Some runs have very stable performance, but far from the optimum

Most runs find good values but do not hold there





Hyperparameter selection

Effect of missing data







When combining activity records from several targets/assays in a pathway, the resulting activity matrix is usually incomplete

It has been stated that even small data sets can help in preventing bias and generalizing the networks internal representation

We were interested to test how performance deteriorates with increasing sparseness of activity labels







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Image: Sector sector

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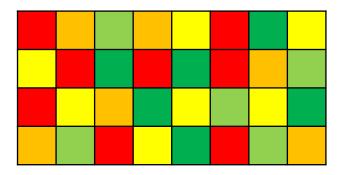
Ramsundar B et al. arXiv 1502.02072

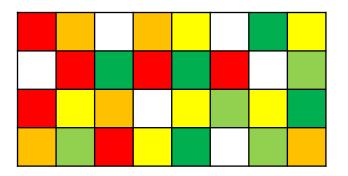


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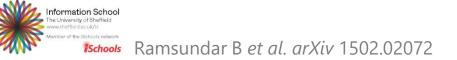
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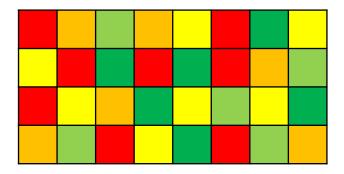


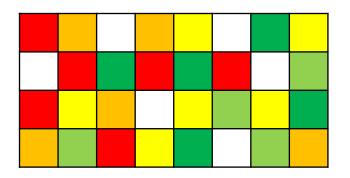


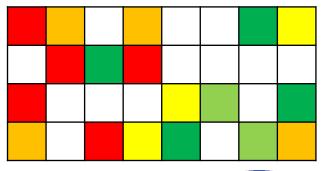
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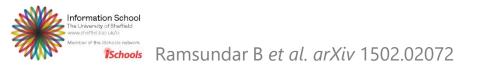
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PKIS (GSK Published Kinase Inhibitor Set) data: percent inhibition of ~ 370 compounds in 454 kinase assays (for most kinases there are values at 1 and 0.1μ M)

¹/₄ of the compounds are used as test set and the rest is used for training

10 sets of DNN hyperparameter values were set, and for each set 100 DNN models were trained with different amounts of training labels removed

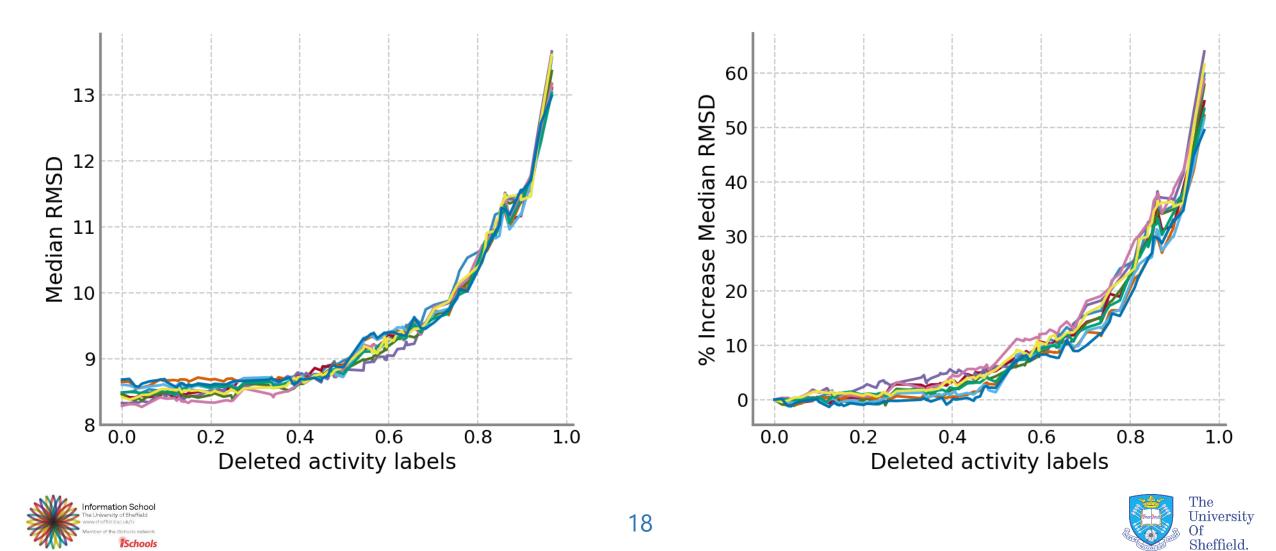
Performance was measured as RMSD RMSD

$$D = \sqrt{\frac{\sum(y_i - \widehat{y_i})^2}{n}}$$



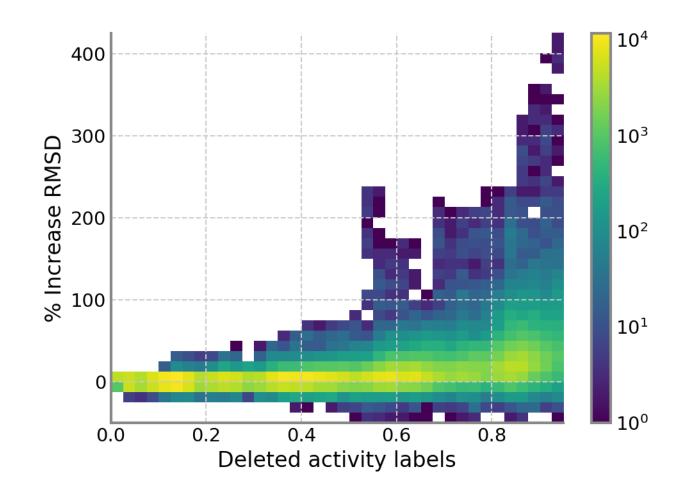


Median performance holds well even if half of the training data is removed



Median performance masks large differences between individual prediction tasks

Most tasks see small differences even with large decreases of training data



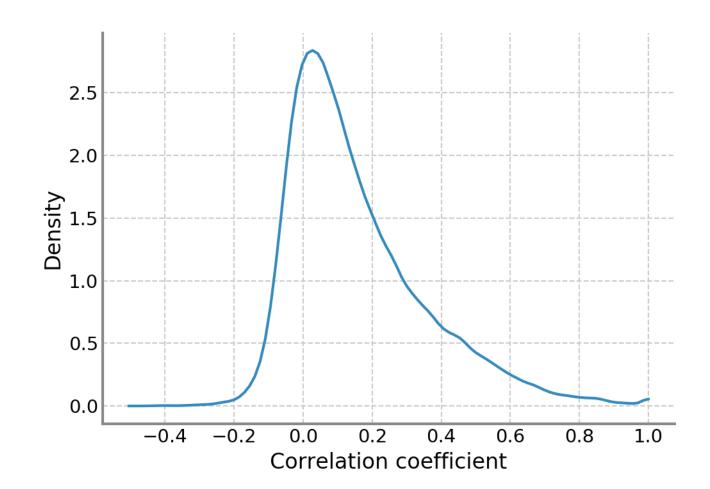




Median performance masks large differences between individual prediction tasks

Most tasks see small differences even with large decreases of training data

Correlation between tasks is not large enough to explain the performance consistency







Conclusions

Deep neural networks are becoming increasingly popular in chemoinformatics

Bayesian optimization provided mixed results and might not be worth the computing cost compared to random parameter selection

Data sparseness is a frequent issue in multi-target data sets such as those that model biological pathways

The performance of deep neural networks is resilient in low to middle data sparseness scenarios





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