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Neural networks for drug discovery

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Neural networks for drug design: correlated data



Two-dimensional fragmented data

X	у
\checkmark	\checkmark
\checkmark	\checkmark
\checkmark	×
\checkmark	×
×	\checkmark

Three-dimensional fragmented data

X	У	Z
\checkmark	\checkmark	\checkmark
\checkmark	\checkmark	×
\checkmark	×	\checkmark
\checkmark	×	×
×	\checkmark	×

Three requirements for the neural network tool

X	у	Z
\checkmark	\checkmark	\checkmark
\checkmark	\checkmark	×
\checkmark	×	\checkmark
\checkmark	×	×
×	\checkmark	×

Correlated data Uncorrelated data Uncertainties Correlated fragmented data

z(x) = y(x) + c with $y = x^2$

if we know one of x, y, or z we can recover both missing quantities



Uncorrelated fragmented data

z(x) = y(x) + x with y(x) = rand(x)need to know both x and y to recover z



Uncertainty



Industrial applications of neural network tool

DFT and experimental





DFT and experimental





Quantum mechanics and experimental





Industrial applications of neural network tool

Experimental

Experimental









Structural and experimental





Proposed alloy



Cr:15.8

Ti: 3.0





Co: 20.0



Mn: 0.2

Mo: 0.5



Si: 0.2

W: 0.5



Ta: 4.9

C: 0.02



B: 0.06

Nb: 1.1



AI: 2.4

Zr: 0.18











Ni: 47.2



900°C

30 hours



Microstructure



Testing the yield stress



Testing the yield stress



Testing the yield stress



Alloys discovered

Cr-Cr₂Ta alloys Intermetallics, 48, 62



Combustor alloy GB1408536



Discovery algorithm EP14153898 US 2014/177578



Mo-Hf forging alloy EP14161255 US 2014/223465



Mo-Nb forging alloy EP14161529 US 2014/224885

RR1000 grain growth

Acta Materialia, 61, 3378



Ni disc alloy EP14157622 US 2013/0052077 A2



Database integrity



Database integrity



Found 792 erroneous points confirmed against primary sources

Protein activity database

Database contains 10,000 proteins and 2,000,000 compounds



Protein activity data

Database has protein activity for 0.1% of entries



Protein activity data

Filled in 32% of the data points with 75% accuracy

Protein



Drug

Cross-validation

Train from first half of the dataset





Test accuracy against second half of the dataset



Statistics



Introducing chemical knowledge of the drug

Exploit SMILES chemical structure to enhance predictions

CCCCN(CC)CCNC(=O)c1cc2c(nn(C)c2s1)-c1ccccc1F

to quantify chemical through 193 descriptors to capture

- Atoms present
- Functional groups
- Properties of the bonds
- Aromatic rings

Chemical knowledge in the database



Chemical knowledge of the drug improves predictions

Filled in 61% of the data points with 75% accuracy

Protein



Drug + structure

Predict activity of a new drug

Filled in 76% of the data points with 75% accuracy

Protein + structure



Drug + structure

Summary of drug discovery

Filled in the 0.1% complete protein activity database to: 32% using activity

61% using activity and drug structure

76% using activity, drug structure, and protein structure

Interface – create a network



Interface – analyze outputs





- Used artificial intelligence to discover materials and drugs
- Handle fragmented data
- Merge experiments and simulations into holistic design tool
- Worked with six different companies, formed startup intellegens