Associative Neural Network

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Supervised regression methods

- Memoryless: multiple linear regression analysis, neural networks, polynomial neural networks, usually these are global models
- Memory-based: k-nearest neighbours (KNN), Parzen-window regression, memory-based reasoning, usually these are local models

Associative Neural Network (ASNN)

A prediction of case *i*:
$$[\mathbf{x}_i] \bullet [\mathbf{ANNE}]_M = [\mathbf{z}_i] = \begin{bmatrix} \mathbf{M} \\ z_k^i \\ \mathbf{M} \\ z_M^i \end{bmatrix}$$

$$\overline{z}_i = \frac{1}{M} \sum_{k=1,M} z_k^i$$

Pearson's (Spearman) correlation coefficient $r_{ij} = R(z_i, z_j) > 0$

$$\overline{z}'_{i} = \overline{z}_{i} + \frac{1}{k} \sum_{j \in N_{k}(\mathbf{x}_{i})} \left(y_{j} - \overline{z}_{j} \right) \quad <<= \text{ASNN bias correction}$$

 $\begin{bmatrix} z_1^i \end{bmatrix}$

The correction of neural network ensemble value is performed using errors (biases) calculated for the neighbor cases of analyzed case \mathbf{x}_i detected in space of models (neural network associations of the given model)

Illustrative example

0.1

0.3



Interpolation of $y=sin(x=x_1+x_2)$

Simple ensemble average

ASNN (one hidden neuron)



Gray (black) line corresponds to neural networks with one (two) hidden neurons. The bias problem (underfitting) is more prominent for one-hidden neuron networks. ASNN dramatically decrease bias of the network prediction.

Similarities in input/output space



 $Y=Gauss(x_1+x_2)$

Similarities of symmetric & non-symmetric functions



Nearest neighbors of case $(x_1, x_2)=(0,0)$ are shown as black circles. Nearest neighbors of case $(x_1, x_2)=(1,0)$ are shown as gray circles.

Gauss function interpolation with fresh data

Features:

fast, no weights retraining; correction is not limited by the range of values in the training set.



N.B! KNN in the output space works better, since it takes into account invariance $x=x_1+x_2!$

ALOGPS - program to predict lipophilicity (logP) and aqueous solubility (logS) of chemicals

LogP: 75 input variables corresponding to electronic and topological properties of atoms (E-state indices), 12908 molecules in the database, 64 neural networks in the ensemble. Calculated results RMSE=0.35, MAE=0.26, n=76 outliers (>1.5 log units)

LogS: 33 input E-state indices, 1291 molecules in the database, 64 neural networks in the ensemble. Calculated results RMSE=0.49, MAE=0.35, n=18 outliers (>1.5 log units)

Tetko et al, JCICS, 2001, 41, 1488-1493 & 1407-1421

Percentage of molecules within indicated error range for lipophilicity prediction

logP _{pred} - logP _{exp}	LOO for the training set	BASF, 6100 "as is"	BASF, 6100 LOO ¹
00.3	63%	30%	60%
00.5	81	49	80
01.0	96	82	96
02.0	99	98	99

¹Tetko, 2002, JCICS, 42, 717-728.

What are the Roots of ASNN? Efficient Partition Algorithm!



selection of cases (feedback loop) Tetko & Villa, ICANN'95, and Neural Networks, 1997



Tetko, I.V.; Villa, A. E. P. Neural Networks 1997, 10, 1361-1374.

ASNN & logP

- More theoretical articles:
- Tetko, I.V. Neural Network Studies. 4. Introduction to Associative Neural Networks, *J. Chem. Inf. Comput. Sci.*, 2002, 42, 717-728.
- Tetko, I.V. Associative Neural Network, Neural Processing Letters 2002, 16, 187-199.
- Tetko, I.V.; Villa, A. E. P. Efficient Partition of Learning Datasets for Neural Network Training, *Neural Networks* 1997, 10, 1361-1374.
- More applied one:
- Tetko, I.V.; Tanchuk, V. Yu. Application of Associative Neural Networks for Prediction of Lipophilicity in ALOGPS 2.1 program, *J. Chem. Inf. Comput. Sci.*, 2002, in press.

 These articles + posters are available at <u>http://vcclab.org/lab/pdf</u>

ASNN is available at http://vcclab.org/lab

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