

Use of Artificial Neural Networks to predict Aqueous Two-Phases System Optimal Conditions on Bromelain's Purification

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Bromelain is the denomination chosen to the group of endoproteases obtained from pineapple and from most of plants belonging to *Bromeliaceae* family. These enzymes have being widely studied in researches across the world due its physiological activity and biotechnological potential. While Brazil still cultivating over 60,000 hectares of pineapple, there is a optimistic trend that aim bromelain's recovery from agriculture residues (stalk and leaves) and fruit processing residues (stem and bark) leading to a fully integrated process which aggregate value to vegetal residues. Our previous studies applied Aqueous Two-Phases Systems and Fractional Precipitation to purify bromelain and achieve purification factor and yield of 11.80 and 87.36, respectively. However, such studies were designed and analysed using Design Of Experiments (DOEs), which lead to an optimal condition but cannot predict with accuracy the complex phenomena of partitioning using ATPS. This work is part of an initiative that aims establish a protocol to calculate more accurate partitioning data through the use of Artificial Neural Networks (ANNs) over a dataset that has being improved continuously. The ANN will determine the relationship between five input parameters (temperature, PEG's molar mass, concentration of PEG, concentration of ammonium sulphate and dilution factor of sample) with three output parameters (protein partition coefficient, Activity partition coefficient and purification factor). The method applied a feed-forward neural network trained with Levenberg-Marquardt algorithm and the Bayesian regularization over the normalized experimental data. The network generated proved the reliability of the method which combined datasets from different DOEs and obtained regression coefficient (~.99) and error (MSE ~0.02) satisfactory for such amount of data used so far.

1. Introduction

The group of thiol-endopeptidases known as Bromelain can be extracted from any plant belonging to the *Bromeliaceae* family (Heinicke and Gortner, 1957) and originally was used as a folk medicine by the aboriginal inhabitants of Central and South America to treats several sicknesses (Taussig and Batkin, 1988). These enzymes had proven therapeutic applications as an anti-inflammatory drug (Salas et al., 2008), in the treatment of allergic disease (Secor Jr et al., 2005), carcinopreventive agent (Harrach et al., 1994), antithrombotic and fibrinolytic activities (Maurer, 2001).

Unlike most enzymes, bromelain is stable and highly active in both acid and alkaline solutions (which expand its range of possible applications) and holds its proteolytic activity even at 60°C, when most enzymes denatures (Bhattacharyya and Bhattacharyya, 2009).

Their purification has being studied for extensively in the last decade: While Harrach et al. (1995) applied Fast Protein Liquid Chromatography as a way to Isolate and characterise the enzymes, Rabelo et al. (2004)

decided to study the use of Aqueous Two-Phase Systems, which has a higher throughput capacity than chromatography techniques, to purify the same enzymes. Over the years, researchers have investigated the use of alternatives bulk recovery techniques, such as expanded bed absorption (Silveira et al., 2009) and Fractional Precipitation (Martins et al., 2014) but the use of ATPS if by far the most employed: Reverse micelles (Umesh Hebbar et al., 2008) , High-speed counter-current chromatography (Yin et al., 2011), combining it with fractional precipitation (Coelho et al., 2012) and endless variety of salts and polymers, such as PEG/potassium phosphate (Ferreira et al., 2014) and PEO–PPO–PEO block polymers (Rabelo et al., 2004).

While the researchers seem to have tried to exhaust the possible combinations of components and modes when using ATPS, they all lack in use a fast and reliable method to determine the best operational conditions. At this moment, there is no method to determine such characteristics with no use of a time-consuming and laborious experimental method. ATPS's characterisations rely on empirical determination of purification parameters for every single modification in the systems under study. One might use statistic methods (such as Design of Experiments) to reduce experimental work but it still lacks in handle trade-off problems as a purification process.

What if we could use a cluster of randomly distributed data obtained to optimize specific parameters in a much broader purpose? That is exactly the purpose of this initiative: to combine all data generated through decades of research in a database that can be constantly improved.

2. Materials and Methods

2.1 ATPS Data acquisition

All experimental data was acquired in projects realized previously, in which we used Design of Experiments and Response Surface Methodology to optimize the parameters or determine a specific operational condition. At this study we restricted the data to those a limited number of variables and in a specific range. The chosen input variables and the correspondents ranges are presented in the Table 1. Those variables were selected from studies in which their effects in the purification were evaluated. The ones presented here showed higher impact during experiments.

Table 1: Input variables used in the neural model and their range

| Input variables | Description | Range |
|---|------------------------------------|-----------------------|
| Temperature (°C) | Operational Temperature | 5 - 25 |
| MMPEG | PEG Molecular Mass | 2,000 – 4,000 – 6,000 |
| (m/m,%) (NH ₄) ₂ SO ₄ | Concentration of Ammonium Sulphate | 7 a 20 |
| (m/m,%) PEG | PEG Concentration | 9 a 30 |
| Dilution (%) | Dilution Factor | 25 - 50 - 75 |

2.2 Mathematical definition of Output Variables

As output variables, we chose the protein partition coefficient (K_p) and the enzymatic partition coefficient (K_A), as described in the table 2. Coelho et al. (2013) describes in details the equationing for the chosen output variables.

Table 2: Output variables used in the neural model and their range.

| Output variables | Description | Minimum | Maximum |
|------------------|---------------------------------|---------|---------|
| KP | Protein Partition Coefficient | 0 | 100 |
| KA | Enzymatic partition Coefficient | 0 | 100 |
| PF | Purification Factor | 0 | 98.75 |

2.3 Results and Discussion

As no mathematical model can predict the complex nature of aqueous two-phase systems in enough accuracy, we decided to evaluate the application of Artificial Neural Networks (ANN's) in the modelling and prediction of partitioning parameters.

Basically, an artificial neural network is a system composed of hundreds of units; artificial neurons (AN) or processing elements (PE), which are connected with coefficients (weights) constituting the neural structure and is arranged in layers as can be seen in the figure 1 (Chrislb, 2005).

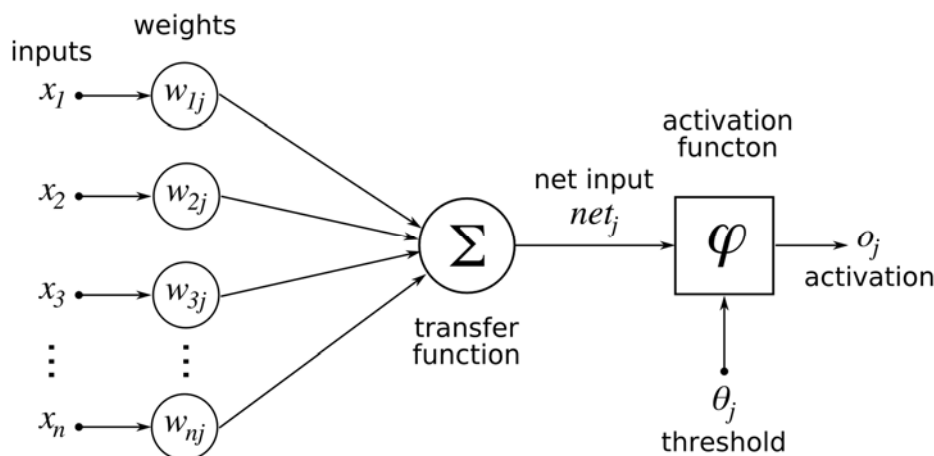


Figure 1: Diagram of an artificial neuron

When a set of input and output data is used to stimulate a “learning” network, such data is used to adjust each neuron’s “weights” through successive changes in its values so that the network implement and execute the desired functions (Brumatti, 2005) and apply the “knowledge” gained from past experiences to new problems or conditions.

This study used the Levenberg-Marquardt optimization as the training algorithm but it was used Bayesian regularization in order to improve generalization and avoid overfit. This gain is a consequence of the smaller weights calculated by the algorithm, which make the network respond smoother (Foresee and Hagan, 1997).

As mentioned, the neural model used either a backpropagation network or a feedforward network coupled with Levenberg-Marquardt and Bayesian regularization optimization algorithms, all available in the Neural Network toolbox from MATLAB © Software (The MathWorks Inc., 2013). All variables were normalized between 0 and 0.9.

As activation functions, were tested hyperbolic tangent function, sigmoid function and a linear function. The neural networks were set and trained combining the different neural models and activation functions (besides the number of neurons) in order to determine which topology converged faster. To estimate the deviation between the ANN’s results and the experimental data, we used the Mean Squared Error (MSE) and the regression coefficient (R), which are the most common parameters used on its analysis (Beale et al.). The neural network would be considered fit to the experimental data when MSE tend to zero and R to 1.

Among the results (Table 3) obtained from the topology optimization for the neural network, the best configuration is the T3, which used 30 neurons and no intermediary layers. We compared values of R, MSE and also the convergence time, being the last one the main factor that made T3 better than T7. These results were obtained during the initial step of this research project and hence used a dataset with only 120 experiments and such variance obtained is expected. Although Aqueous Two-Phases Systems has been used for decades, there is no such thing as a model that can precisely predict any property from those systems with no experimental data. This creates an even harder task to find an appropriated approach to study it.

Table 3: Neural Networks training results using KP, KA and PF as output variables

| T | (Nrn,Lyr) | R | MSE | KP | KA | PF | Inter. Function | Inter. Function 2 | Output Function |
|-----|-----------|----------------|---------|---------|---------|---------|-----------------|-------------------|-----------------|
| T1 | (10,1) | 0.8862 | 0.14369 | 0.88394 | 0.91154 | 0.67951 | Tansig | Purelin | Purelin |
| T2 | (20,1) | 0.9514 | 0.06371 | 0.95463 | 0.95836 | 0.87268 | Tansig | Purelin | Purelin |
| T3 | (30,1) | 0.9846 | 0.02045 | 0.99418 | 0.98574 | 0.92548 | Tansig | Purelin | Purelin |
| T4 | (30,2) | 0.6465 | 0.40178 | 0.63385 | 0.66119 | 0.49902 | Tansig | Purelin | Purelin |
| T5 | (30,2) | 0.9807 | 0.02573 | 0.99149 | 0.98097 | 0.91795 | Logsig | Purelin | Purelin |
| T6 | (40,2) | 0.9858 | 0.01901 | 0.99351 | 0.98713 | 0.93540 | Tansig | Purelin | Purelin |
| T7 | (50,2) | 0.9854 | 0.01953 | 0.99307 | 0.98720 | 0.93223 | Tansig | Purelin | Purelin |
| T8 | (5,2) | 0.7577 | 0.28486 | 0.75959 | 0.76642 | 0.54001 | Tansig | Purelin | Purelin |
| T9 | (10,2) | 0.8814 | 0.14927 | 0.88072 | 0.90113 | 0.70156 | Tansig | Purelin | Purelin |
| T10 | (20,2) | 0.9688 | 0.04160 | 0.98592 | 0.97862 | 0.80179 | Tansig | Purelin | Purelin |
| T11 | (30,3) | 0.9840 | 0.02118 | 0.99243 | 0.98553 | 0.92806 | Tansig | Purelin | Purelin |
| T12 | (30,4) | No Convergence | | | | | | | |

Tansig: Hyperbolic Tangent, Purelin: Linear, Logsig: Sigmoidal, Nrn: Neurons, Lyr: Layers

In this set of simulations we tried to test an even bigger number of combinations between the number of neurons, number of intermediary layers and even the activation functions but most of them couldn't even converge. Thus, the topology with 30 neurons and using a hyperbolic function as activation function provided the best results.

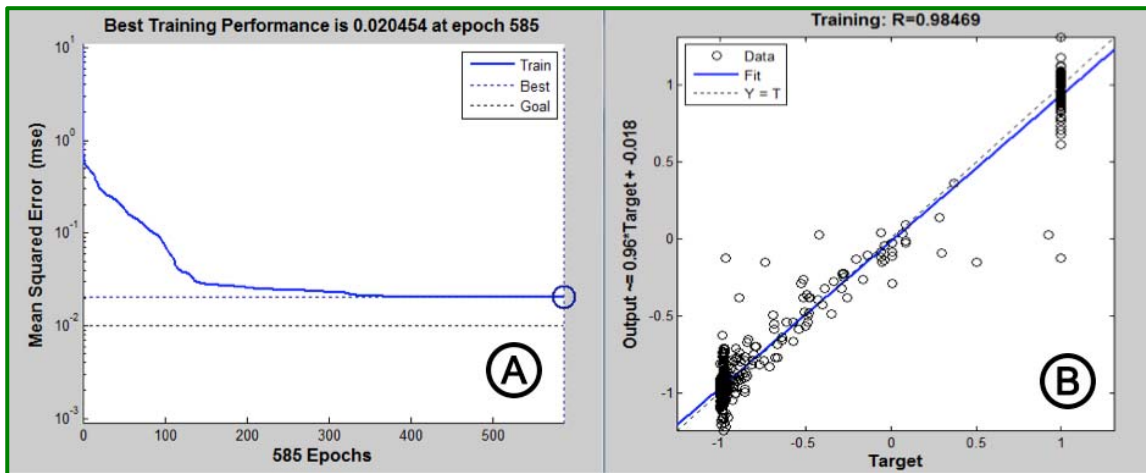


Figure 2: Convergence (A) and Regression (B) for the best topology obtained (T3, Table 3)

Figure 2 presents the fitting parameters results obtained for T3 topology, which was the one that returned the best results. Figure 3 presents the regression data using the topology T3 for the output variables (K_A , K_P and P_F). It is noticeable that data represents well the experimental data but we still expect to be able to improve the model in at least 5 %. Positive results are mainly due Bayesian regularization, which improve the generalization capability of the model even in a reasonable high operational range (Fileti et al., 2010).

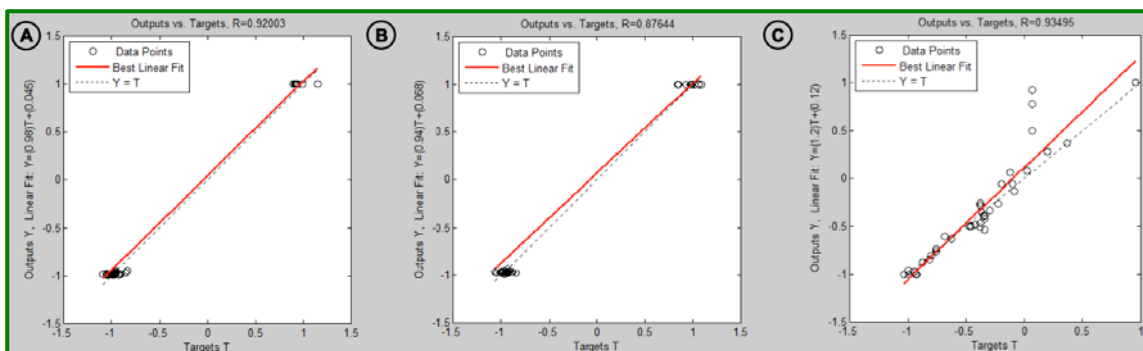


Figure 3: Output Test with T3 Topology output Neural Network for KA, KP and PF respectively

However, the study still lacks in explain why we couldn't decrease MSE and the variation observed in figure 2 and at this point the network just proved that was able to correlate data from several experiments and show we can improve the network and use it to obtain a better understanding of the process.

3. Conclusions

The network generated proved the reliability of the method by modelling combined data from different experimental designs and obtaining reasonable regression coefficient (~ 0.99) and error (MSE ~ 0.02). At this stage the neural network was able to model and predict with certain precision the data handled. However, it is necessary to improve the robustness by increasing the number of input in the database used to train the network. When complete, the neural model will be able to predict operational points, analyze influence of different factors and select conditions in which a trade-off is present.

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