

Article

Prediction of Protein Content in Pea (*Pisum sativum* L.) Seeds Using Artificial Neural Networks

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Abstract: Pea (*Pisum sativum* L.) is a legume valued mainly for its high seed protein content. The protein content of pea is characterized by a high lysine content and low allergenicity. This has made consumers appreciate peas increasingly in recent years, not only for their taste, but also for their nutritional value. An important element of pea cultivation is the ability to predict protein content, even before harvest. The aim of this research was to develop a linear and a non-linear model for predicting the percentage of protein content in pea seeds and to perform a comparative analysis of the effectiveness of these models. The analysis also focused on identifying the variables with the greatest impact on protein content. The research included the method of machine learning (artificial neural networks) and multiple linear regression (MLR). The input parameters of the models were weather, agronomic and phytopenological data from 2016–2020. The predictive properties of the models were verified using six ex-post forecast measures. The neural model (N1) outperformed the multiple regression (RS) model. The N1 model had an RMS error magnitude of 0.838, while the RS model obtained an average error value of 2.696. The MAPE error for the N1 and RS models was 2.721 and 8.852, respectively. The sensitivity analysis performed for the best neural network showed that the independent variables most influencing the protein content of pea seeds were the soil abundance of magnesium, potassium and phosphorus. The results presented in this work can be useful for the study of pea crop management. In addition, they can help preserve the country's protein security.

Keywords: artificial neural networks; multiple linear regression; protein prediction; pea; sensitivity analysis; weather conditions



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1. Introduction

In terms of cultivation, legumes are the world's second largest crop after cereals. They constitute about 30% of the world's plant production [1]. The crop residues of these plants are characterized by a positive organic matter (MO) balance, which makes them a very good forecrop for many crops such as cereals, potatoes and beets. In the years with an uneven distribution of precipitation (temperate climates) or a shortage of precipitation (southern European climatic conditions), when there is a poor uptake of mineral nitrogen, a particularly favorable after-effect of legumes is observed [2,3]. The decomposition of *Fabaceae* crop residues in the soil provides available forms of nitrogen to both successor plants and soil microorganisms.

This process contributes to the intensification of biological nitrogen sorption. The interaction between the legume residues and MO mineralization determines the amount of available N for the next plant [4]. The participation of legumes in crop rotation also contributes to the reduction of weed, pest and disease populations [5]. When grown in rotation with cereals, they counteract soil erosion and improve soil fertility [6] by changing physical and chemical properties. A properly developed root system of legumes is formed

through loosening the soil, which increases soil aeration and becomes a source of a large amount of organic matter rich in nitrogen and other minerals [7]. A significant property of this group of plants is the ability to fix atmospheric nitrogen as a result of symbiosis with *Rhizobium* spp. bacteria. This trait plays an important role in argoecosystems and sustainable crop production, which seeks to reduce the use of mineral fertilizers [8,9]. The symbiosis of legumes with papillary bacteria also reduces inputs and resources by reducing the need for nitrogen fertilizers [10]. The importance of these plants is particularly important in an era of high and volatile mineral fertilizer prices. In many European countries, these fertilizers are periodically becoming a scarce commodity, so legumes are an important element in countering the fertilizer crisis and fitting in with the ideas of sustainable agriculture.

One of the most important plants in the *Fabaceae* family is the pea. More than half of its world production is in Canada, Russia, the United States of America and India [11]. In Poland, peas are the most widely grown legume right after yellow lupin. In 2022, the area of pea cultivation in Poland was more than 105 thousand hectares. For example, the cultivation of soybeans and beans in the same year amounted to 48.20 and 30.70 thousand hectares, respectively. In 2022, peas accounted for 0.68% of the total area in the structure of national sowings [12].

Peas are most valued for their high seed protein content, which can be as high as 31% [13,14]. Pea protein has high nutritional value due to its relatively high content of lysine, an amino acid that limits the nutritional value of cereals. In addition, it is characterized by low allergenicity [15,16]. Despite its significant and obvious advantages, the area under cultivation for this crop is relatively low due to poor profitability as a result of biotic-abiotic factors [17]. However, there is growing interest in plant-based proteins as a substitute for animal-based proteins [18]. The reason for this phenomenon is greater awareness of nutrition, environmental concerns and ethical issues [11]. Therefore, it can be assumed that peas will become increasingly popular among farmers over the next few years, which will be reflected in the increasing area under cultivation.

The demand for protein will continue to grow in the coming years due to the world's expanding human population [1]. This makes it likely that interest in legumes, including peas as a valuable source of protein, will be greater than before. In addition, peas are a very good component in feed production. As a protein-raw material with a satisfactory amino acid composition, it is used in the feeding of slaughter, dairy and laying animals without adversely affecting production and fattening performance [19]. For many years, efforts have been made in European countries, including Poland, to increase the production and use of domestic protein raw materials to replace, or at least supplement, expensive imported post-extraction soybean meal. These measures are also aimed at preserving the country's protein security, as the feed market is largely dependent on imported protein raw materials. Therefore, there is a risk (for now only theoretical) of a shortage of protein feed for animals, and consequently a shortage of food for the population [20]. Therefore, the ability to predict the protein content of pea seeds is very important for the possibility of ongoing decision support, management of national protein resources and risk management [21]. However, prediction of crop quality traits is a very difficult task. During the growing season, plants are exposed to a number of factors that limit both yield and quality, and predicting many of these factors is often an impossible task [22]. The non-uniform course of weather conditions, soil variability or pest pressure causes the growth and development of agricultural crops to proceed differently in each growing season [23]. In addition, the non-linear interaction between environmental factors and plant growth can result in a low-precision predictive model with a large prediction error [24,25]. The large number of factors influencing the quality of yield poses a significant difficulty in the selection of independent variables. Therefore, the construction of predictive models should be supported by very good knowledge of the research object [22]. This knowledge will allow the selection of those input variables that significantly affect yield quality. Among the most commonly used independent variables in the prediction of yield and its chemical and biochemical

characteristics are weather data [26–29]. Average air temperature, total precipitation or total sunshine provide valuable information on plant development conditions. Data on soil mineral abundance, fertilizer application rates, and the course of plant phenological traits are commonly used to build predictive models [30–32].

One effective method for yield quality prediction is machine learning, among which artificial neural networks (ANNs) are of great interest [33–37]. The prototypes of ANNs are the nerve cells that build the human brain, so the operation of artificial neural networks is similar to that of the human brain [38,39]. Each neural network is made up of many simultaneously working and jointly processing elements called neurons. Neurons, due to their function, can be divided into three basic groups: input neurons (they are responsible for inputting the signal into the network), information processing neurons and output neurons (“producing” the results of the network to the outside world) [38,40]. Each of these groups of neurons forms a separate layer, the function of which is the same as the function of the elements from which it is built. Thus, the first is the input layer, which contains a number of neurons equal to the number of independent variables. Its task is to separate the input data into a number of neurons contained in the hidden layer. The hidden layer is built from the n -th number of neurons, the number of which depends on the complexity of the problem being solved by the network. In the structure of a neural network, there can be a different number of hidden layers. The decision on how many hidden layers to use is made by the network developer and is generally an arbitrary decision. The last third layer usually contains only one neuron, responsible for transmitting the result. A neural network, as a layered structure, works by connecting adjacent layers on an “each to each” basis [36,41].

This paper presents the possibility of using ANN to predict the protein content of pea seeds. An extensive analysis of the literature revealed a lack of scientific work of a similar nature. There are no reports on the possibility of predicting the protein content of peas under Polish weather and habitat conditions. In this study, three hypotheses are put forward for verification: (i) artificial neural networks are an effective tool for predicting the protein content of *pea* seeds 20 days before harvest; (ii) it is possible to create a model predicting the protein content of pea seeds based on five-year field trials; and (iii) the ANN model predicts the protein content of pea seeds with greater accuracy compared to the MLR model.

2. Materials and Methods

Experimental data were obtained from a 5-year cycle of field experiments with peas which were conducted in Poland. The results of the experiments were obtained from the field books of the system of the Research Center for Cultivar Testing (COBORU) [42]. Among other things, this institution is engaged in research on distinctiveness, uniformity and durability (DUS) of crop varieties. It is also within the scope of COBORU to conduct field trials for cultivation and use value (VCU). Obtaining positive results from these experiments allows a given variety to be included in the National Variety List. In addition, COBORU supervises the legal protection of varieties entered in the National Register [43]. The field books were created based on data from the official results of experiments under the Program of Registered Varietal Testing (PRVT; in Polish, PDO). PRVT is a system of permanent or periodic testing on the economic value of crop species listed in the National Register or included in the Community Catalogs of Agricultural/Vegetable Varieties (CCA/CCV). PRVT covers both varietal and varietal-agronomic experiments [44].

Field experiments were conducted at the Stations and Experimental Plants for Variety Testing of COBORU located in: Bezek (51°12′6.722″ N 23°16′7.656″ E), Głębokie (52°38′33.18″ N 18°26′16.26″ E), Kawęczyn (52°10′15.157″ N 20°20′49.328″ E), Krzyżewo (53°1′33.535″ N 22°45′28.438″ E), Pawłowice (50°27′14.049″ N 18°29′28.912″ E), Radostowo (53°59′20.566″ N 18°44′41.429″ E) and Sulejów (51°21′8.03″ N 19°52′7.517″ E) (Figure 1). These localities were chosen for their optimal conditions for pea cultivation. These locations are dominated by clay soils of class II-IIIb (polish classification). The experiments

were conducted in accordance with COBORU methodology, which includes a number of agrotechnical recommendations. All studies on selected pea varieties were conducted on plots of 13.86 m². The experiments were conducted in an arrangement with variety groups for species in which different morphological types are studied (e.g., traditional and self-terminating varieties, tall, medium-high and low varieties, etc.). In a system with groups of varieties, first the place in repetitions is drawn or determined, and then the order of varieties in groups. The number of repetitions for each variety in each year of the study was 3. A model based on artificial neural networks (N1) and multiple linear regression (RS) was developed for 11 general-purpose pea varieties: Arwena, Astronaute, Batuta, Mecenasa, Medyk, Mentor, Olympus, Spot, Starski, Tarchalska and Tytus. These varieties are widely recommended for cultivation in Poland due to their relatively high yield levels and relatively high resistance to biotic factors.

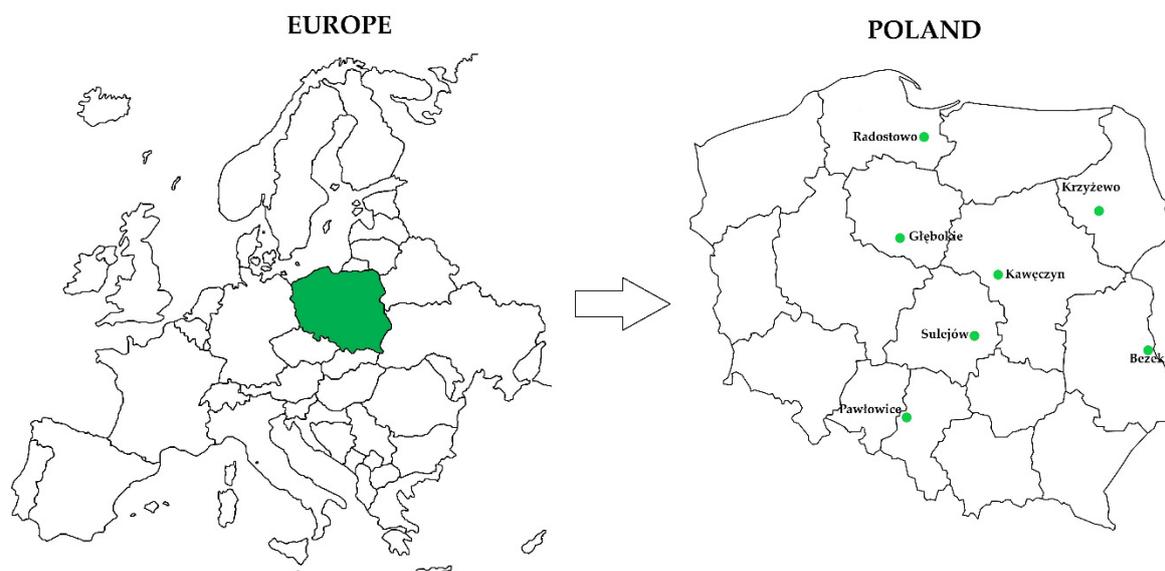


Figure 1. Location of field experiments conducted.

2.1. Data for Model Construction

Two categories of variables were used during the creation of the N1 and RS model. The first group was agronomic data, phytophenological data and results of protein content of pea seeds, all of which came from COBORU field books. The second category of variables was meteorological data, which came from a dataset of meteorological phenomena and observations recorded at each COBORU Variety Testing Station and Department. Missing data, such as sunshine totals, were supplemented using historical data from meteorological stations of the Institute of Meteorology and Water Management–National Research Institute. Measurements from meteorological stations that were located closest to the experimental facilities were used [25]. This information was obtained from a public archival database, available electronically [45].

2.2. Construction of the Database

Nineteen independent variables were used to construct the N1 and RS models, as shown in Table 1. The dependent variable was the percentage protein content of pea seeds. Data from a total of 1155 plots were used to construct and verify the N1 and RS model. Each of the analyzed plots constituted one separate case for model construction. The information was grouped into two sets A and B. Set A contained data from 1040 plots and was used to build a neural and regression model. Set B, on the other hand, contained cases from 115 plots and was used to validate the models. Therefore, this set was not used to build the neural network and regression model. It should be noted that the data of set B was selected randomly. However, the determinant representing each case was the

variety. The percentage of protein in the seeds was the dependent variable for the predictive models created.

Table 1. Variables used to build the N1 and RS model.

Symbol	Unit of Measure	Variable Description	Data Range
Independent Variables			
RAIN	mm	Total rainfall from sowing date to 14 July	96.9–312.4
SUN	h	Total sunshine from sowing date to 14 July	630.5–1051.5
TEMP	°C	Average air temperature from sowing date to 14 July	11.0–17.5
N_F	kg·ha ⁻¹	Total nitrogen from mineral fertilizers	10–90
P2O5_F	kg·ha ⁻¹	Total phosphorus from mineral fertilizers	0–80
K2O_F	kg·ha ⁻¹	Total potassium from mineral fertilizers	0–119
SOWI	days	Number of days from 1 January to sowing date	83–102
P_EMER	days	Number of days from 1 January to the beginning of plant emergence	96–133
HAR	days	Number of days from 1 January to the date of harvesting	184–221
FLOWE	days	Number of days from 1 January to the beginning of flowering	126–169
INI_MA	days	Number of days from 1 January to onset of maturity	167–211
TECH_M	days	Number of days from 1 January to technical maturity	171–216
P_HIG	cm	Plant height	43–156
WEGW	days	Number of plant growing days	87–137
PH	-	Soil pH	5.5–7.5
P2O5_C	Scale from 0 to 4 *	P ₂ O ₅ content in the soil	0–4
K2O_C	Scale from 0 to 4 *	K ₂ O content in the soil	0–4
MGO_C	Scale from 0 to 4 *	MgO content in the soil	0–4
GEN	feature coded 101 to 111	General variety of peas	-
Dependent variable			
PROT	%	Percentage of protein in pea seeds	18.56–29.22

* The scale from 0 to 4 refers to the abundance of macronutrients in the soil and is determined as follows: 0—very low, 1—low, 2—medium, 3—high, 4—very high.

The construction of the ANN model was performed on the basis of the predicted date for the calendar year, i.e., 14 July. Based on the analysis of data from all five years of the study (2016–2020), it was shown that pea harvesting was most often performed on 3 August, and the latest on 10 August. The date of 14 July, which is the date of prediction, is the dominant beginning of maturity of the analyzed varieties. This approach to predicting the protein content of pea seeds makes it possible to make predictions 20 days before harvest (based on the dominant harvest date) in the same calendar year.

2.3. Determination of Protein Content in Pea Seeds

Protein determination using the Kjeldahl method is a standard method for determining proteins in plant raw materials [46]. It involves the conversion of protein nitrogen into ammonium sulfate with concentrated sulfuric acid. In the first step, the ground and dried sample is mineralized with sulfuric acid (VI) in the presence of K₂SO₄ and CuSO₄ catalysts. To the mineralized and cooled sample, 75 mL of distilled water and 2 of receiving solution are added. The solution thus prepared is distilled for about 4 min. In the final step, the resulting distillate is titrated with standard hydrochloric acid (0.1 mol·L⁻¹) until a gray-green color appears. The amount of hydrochloric acid used for titration is the basis for calculating the total protein content of the sample [47].

2.4. ANN Model Development

The construction of the N1 model was performed on the basis of the predicted date for the calendar year, i.e., 14 July. Based on the analysis of data from all five years of the study (2016–2020), it was shown that pea harvesting was most often performed on 3 August, and the latest on 10 August. The date of 14 July, which is the date of prediction, is the dominant beginning of maturity of the analyzed varieties. This approach to predicting the protein content of pea seeds makes it possible to make predictions 20 days before harvest (based on the dominant harvest date) in the same calendar year.

The choice of network type was made by repeatedly building neural networks using an automatic network designer, as well as by reviewing the available literature [21,48–51]. A total of 10,000 networks were tested, which allowed the selection of a multilayer perceptron type network with the following architecture: MLP 19:19-32-1:1 (Figure 2). The selection of the network architecture was guided using the size of the error of the validation, test and learning sets, as well as key network quality parameters, which are shown in Table 2. Achieving the smallest error values by these sets leads to an ANN with high prediction accuracy [52]. An important element in the construction of the prediction model was the division of set A (1040 fields) into three subsets: learning, test and validation. These subsets consisted of 520, 260 and 260 cases (50%–25%–25%), respectively. The entire procedure was performed in the Statistica v7.1 software (TIBCO Software Inc., Palo Alto, CA, USA).

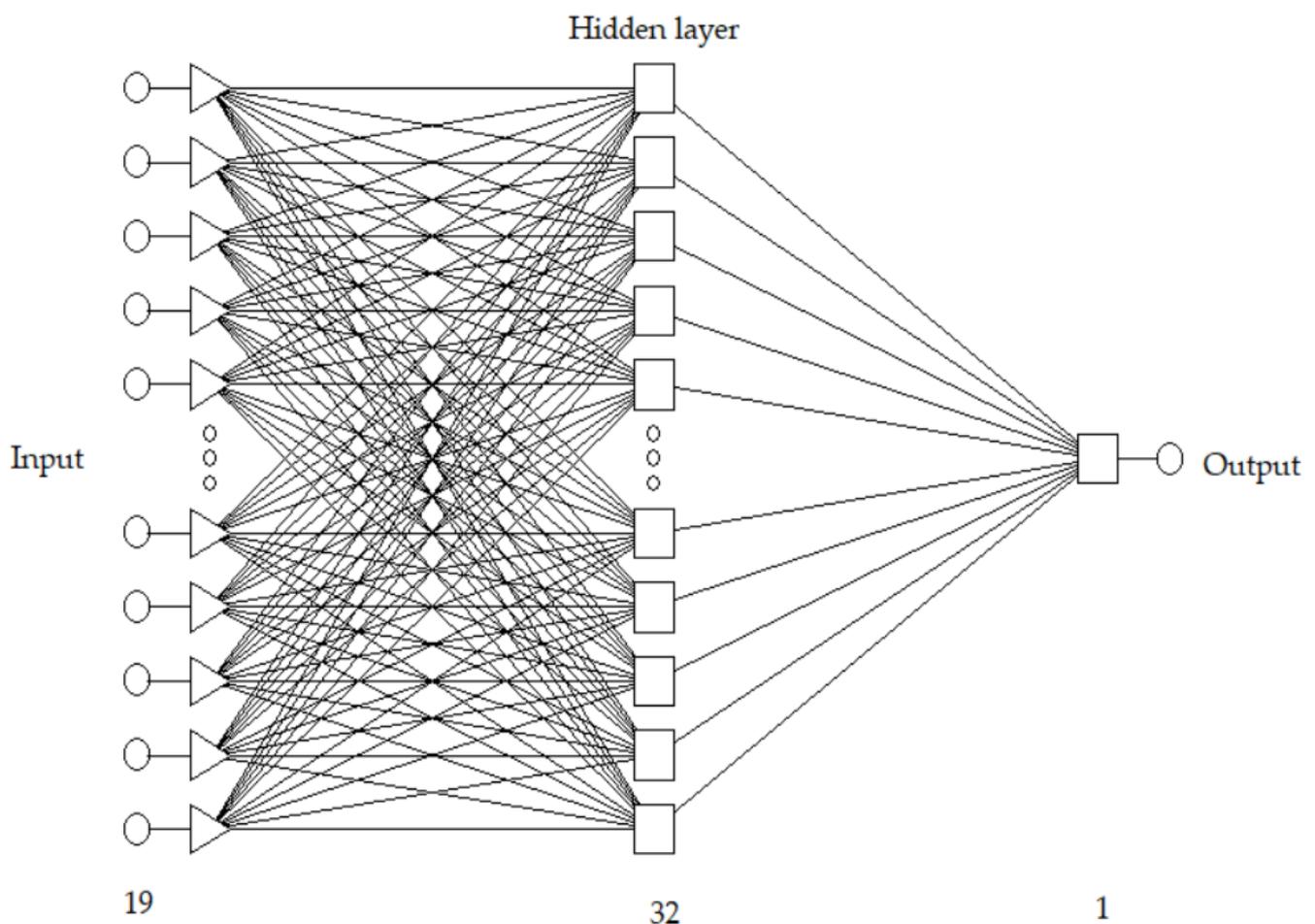


Figure 2. Architecture of neural network with MLP topology.

Table 2. Assessment of the abundance of soils in Poland.

Resources	Phosphorus, mg P ₂ O ₅ ·kg ⁻¹ Soil	Potassium, mg K ₂ O·kg ⁻¹ Soil			Magnesium, mg MgO·kg ⁻¹ Soil		
		Soil Agronomic Category			Soil Agronomic Category		
		Light	Medium	Heavy	Light	Medium	Heavy
very low	up to 50	up to 100	up to 105	up to 170	up to 80	up to 105	up to 120
low	51–80	101–160	106–170	171–260	81–135	106–160	121–220
average	81–115	161–275	171–310	261–350	136–200	161–265	221–330
high	116–185	276–380	311–420	351–510	201–285	266–330	331–460
very high	>185	>380	>420	>510	>285	>330	>460

Due to the coding of the variable P2O5_C, K2O_C and MGO_C, Table 2 presents an assessment of the abundance of Polish soils in phosphorus, potassium and magnesium. Very low abundance of soils in these elements was not recorded in the localities where field experiments were conducted.

2.5. MLR Model Development

Multiple linear regression (MLR) is a statistical tool for detecting interdependencies between independent characteristics and the dependent variable. It makes it possible to determine the strength and type of the detected dependence and to build a functional model that makes it possible to forecast the direction of change in one characteristic on the basis of others [53]. In addition, this method allows you to identify significant and non-significant variables at a given level of probability. This knowledge is particularly important in the further stages of modeling and allows you to identify those invariant variables that do not affect the dependent variable [54]. The general regression formula is shown in Equation (1).

$$Y = b_0 + b_1 \cdot X_1 + b_2 \cdot X_2 + \dots + b_p \cdot X_p + \epsilon, \quad (1)$$

where: Y-dependent variable (explained variable), X₁, X₂ ... X_p-independent variables (explanatory variable), b₀, b₁, b₂ ... b_p-equation parameters, ε-random component (rest of the model).

In this study, a model developed using multiple linear regression (stepwise progressive) was used to predict the percentage protein content of pea seeds. For its construction, the independent variables shown in Table 1 were used. The model was built to compare the effectiveness of the prediction of protein content in peas and was contrasted with a nonlinear model (ANN).

The RS model, like N1, was created based on 1040 cases. The analysis continued through 17 steps, and of the 19 explanatory variables, two of them (number of growing days and number of days from the beginning of the year to emergence) were removed using the model. 115 random observations (set B) were used to predict protein content. Multivariate linear regression analysis was performed using Statistica v7.1 software, and the results are presented graphically and in tabular form.

2.6. Verification of the N1 and RS Models

The obtained N1 and RS models were verified on the basis of measures of predictive properties (Equations (2)–(7)). To calculate them, a set B (115 fields) was used on the basis of which the difference between actual and predicted values was determined. This is example 2 of an equation [22,55]:

$$RAE = \sqrt{\frac{\sum_{i=1}^n (y_i - y'_i)^2}{\sum_{i=1}^n (y_i)^2}}, \quad (2)$$

$$\text{RMS} = \sqrt{\frac{\sum_{i=1}^n (y_i - y'_i)^2}{n}} \quad (3)$$

$$\text{MAE} = \frac{1}{n} \sum_{i=1}^n |y_i - y'_i| \quad (4)$$

$$\text{MAPE} = \frac{1}{n} \sum_{i=1}^n \left| \frac{y_i - y'_i}{y_i} \right| \cdot 100\% \quad (5)$$

$$\text{MAX} = \max_i |y_i - y'_i| \quad (6)$$

$$\text{MAXP} = \max_i \left| \frac{y_i - y'_i}{y_i} \right| \cdot 100\% \quad (7)$$

where: n -number of observations, y_i -actual values, y'_i -predictive values, obtained with the model.

2.7. Sensitivity Analysis of the Neural Network

The final step in creating the N1 model was to identify the independent variables that most influenced the variable explained by the model. To do this, a sensitivity analysis of the network was performed, which allows the rank of each feature to be quantified. The ranks are determined by the size of the deviation quotient, which is the ratio of the error to the error received by all independent variables. The importance of a given feature is greater the higher the deviation quotient achieved by it.

3. Results

3.1. Neural Network Learning and Quality Assessment of Models Predicting Protein Content in Pea Seeds

The obtained multilayer perceptron-type neural network was learned using two methods. The first method of learning-backward error propagation took 100 epochs. The best learning result was obtained by continuing this process with the coupled gradients method and the best result was achieved at 55 epochs. This approach is common in creating predictive models using ANNs [25,28,56–58]. The error for the network did not exceed 0.6 for each of the sets-i.e., learning, validation and test. The results of the N1 predictive model and its basic features are shown in Tables 3 and 4.

Table 3. Subset error size and number of learning epochs of neural networks.

Subsets	Teaching	Validation	Testing
Size of error	0.0551	0.0535	0.0595
Quality	0.3642	0.4145	0.0551
Epochs of learning			
Back propagation method of error		100	
Coupled gradients method		55b *	

* b (best)-the best result in the indicated learning epoch.

Table 4. Quality parameters of the N1 model.

Quality Measures	Value
Average	22.857
Standard deviation	1.895
Average error	0.008
Error deviation	0.744
Average absolute error	0.574
Deviation quotient	0.393
Correlation coefficient r	0.920

The N1 model had a high correlation coefficient ($r = 0.920$). Satisfactory values were also obtained for the mean error and mean absolute error, which were 0.008 and 0.574, respectively. Low error values and a high correlation coefficient were among the many parameters that determined the selection of the N1 model for further analysis.

An analysis using multiple linear regression showed that the explanatory variables statistically insignificant at the $\alpha = 0.05$ level were the number of days from 1 January to the beginning of flowering (FLOWE), variety (GEN), soil potassium and phosphorus abundance (K2O_C and P2O5_C, respectively) and soil pH (PH).

Based on the results shown in Table 5, the multiple linear regression equation is of the form:

$$\text{PROT} = 40.445 + 0.576 \cdot \text{MGO_C} - 0.005 \cdot \text{RAIN} - 0.021 \cdot \text{K2O_F} - 0.028 \cdot \text{N_F} - 0.065 \cdot \text{HAR} + 0.027 \cdot \text{P_HIGH} - 0.214 \cdot \text{TECH_M} + 0.152 \cdot \text{INI_A} + 0.020 \cdot \text{P2O5_F} - 0.416 \cdot \text{TEMP} + 0.061 \cdot \text{SOWI} + 0.001 \cdot \text{SUN} \quad (8)$$

Table 5. MLR analysis results.

Factor	MLR: $r = 0.6949$ $R^2 = 0.4829$ Standard Error of Estimate = 1.374					
	Beta	Standard Error Beta	b	Standard Error b	p	Significance
Free Term	-	-	40.445	0.859	0.000000	+
MGO_C	0.323	0.035	0.576	0.062	0.000000	+
RAIN	-0.154	0.034	-0.005	0.001	0.000007	+
K2O_F	-0.298	0.041	-0.021	0.003	0.000000	+
N_F	-0.187	0.031	-0.028	0.005	0.000000	+
HAR	-0.316	0.041	-0.065	0.009	0.000000	+
P_HIG	0.234	0.033	0.027	0.004	0.000000	+
FLOWE	0.062	0.042	0.021	0.015	0.146017	-
TECH_M	-1.063	0.127	-0.214	0.026	0.000000	+
INI_A	0.745	0.138	0.152	0.028	0.000000	+
P2O5_F	0.195	0.043	0.020	0.004	0.000006	+
TEMP	-0.340	0.086	-0.416	0.104	0.000077	+
SOWI	0.182	0.071	0.061	0.024	0.009821	+
GEN	0.0439	0.023	0.026	0.014	0.056315	-
K2O_C	-0.055	0.031	-0.102	0.063	0.083382	-
SUN	0.067	0.032	0.001	0.0006	0.036298	+
P2O5_C	-0.036	0.029	-0.074	0.060	0.211476	-
PH	0.040	0.032	0.158	0.128	0.219669	-

Determination of the level of statistical significance: - non-significant. + significant for $\alpha = 0.05$.

3.2. Sensitivity Analysis of Neural Networks

Verification of the predictive model based on artificial neural networks was carried out using 115 cases (plots). The N1 model with a structure of 19:19-32-1:1 was prepared based on 19 independent variables. The dependent variable was the percentage protein content of pea seeds. The sensitivity analysis performed on crop A showed that the factor with the greatest effect on the protein content of pea seeds was soil magnesium abundance (Table 6). This trait received a rank of one, and removing this variable from the N1 model would result in an increase in the cumulative error value by 2.366 times. The independent variables that received a rank of two and three were the potassium and phosphorus content in the soil. Not including these variables in the model would have increased the error

by 1546 and 1413 times, respectively. Average daily air temperature received a rank of four and, of all the weather variables, had the greatest effect on the protein content of the eleven pea seed varieties.

Table 6. Quality parameters of the N1 model.

Variable	Quotient	Rank
GEN	1.082	18
RAIN	1.158	12
SUN	1.110	16
TEMP	1.396	4
N_F	1.110	15
P2O5_F	1.095	17
K2O_F	1.194	8
SOWI	1.364	5
P_EMER	1.160	11
WEGE	1.257	7
HAR	1.179	9
FLOWE	1.049	19
INI_MA	1.265	6
TECH_M	1.131	13
P_HIG	1.136	13
PH	1.175	10
P2O5_C	1.413	3
K2O_C	1.546	2
MGO_C	2.366	1

The protein content of pea seeds predicted using the N1 model was compared with actual values (Figure 3). A coefficient of determination was obtained at a relatively high level ($R^2 = 0.7979$), which means that the model's response is very close to the observed values, and that the network has the ability to correctly represent the relationships that are characteristic of the issue being modeled. This procedure was also performed for the RS model (Figure 4). The obtained coefficient of determination of the studied characteristics was 0.3357, so the model has much weaker predictive properties compared to the N1 model.

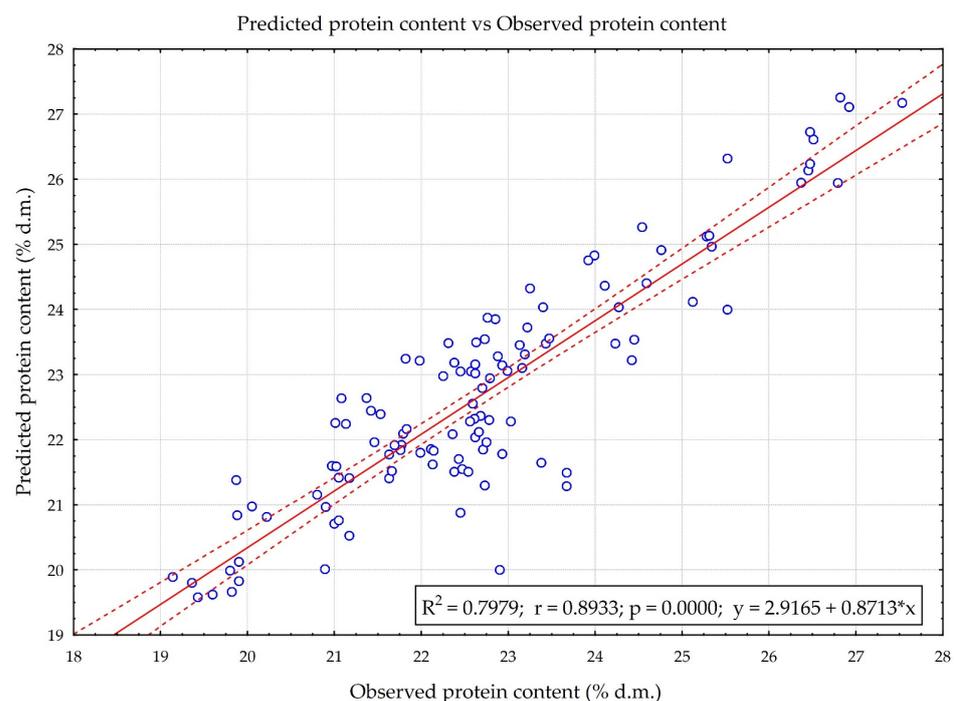


Figure 3. Scatter plot of observed and predicted values for the N1 model.

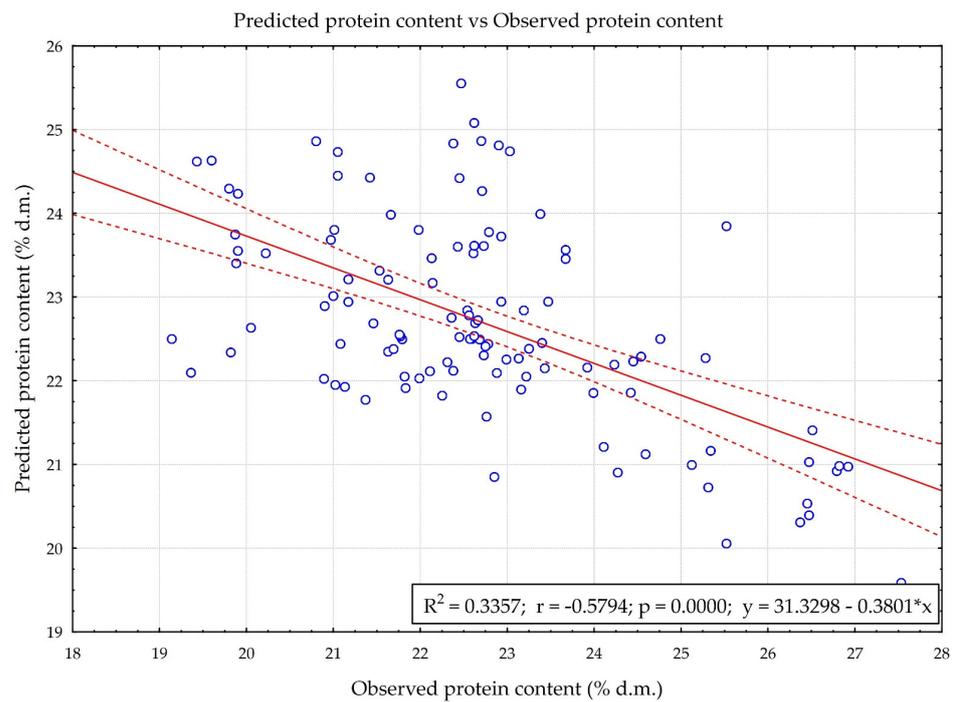


Figure 4. Scatter plot of observed and predicted values for the RS model.

From Figure 5, it can be observed that the protein content of pea seeds changed with the increasing soil magnesium and potassium abundance. An increase in the concentration of magnesium in the soil causes an increase in the percentage of protein content in the analyzed plant. A similar trend is observed for potassium. However, this increase is not as high as in the case of magnesium.

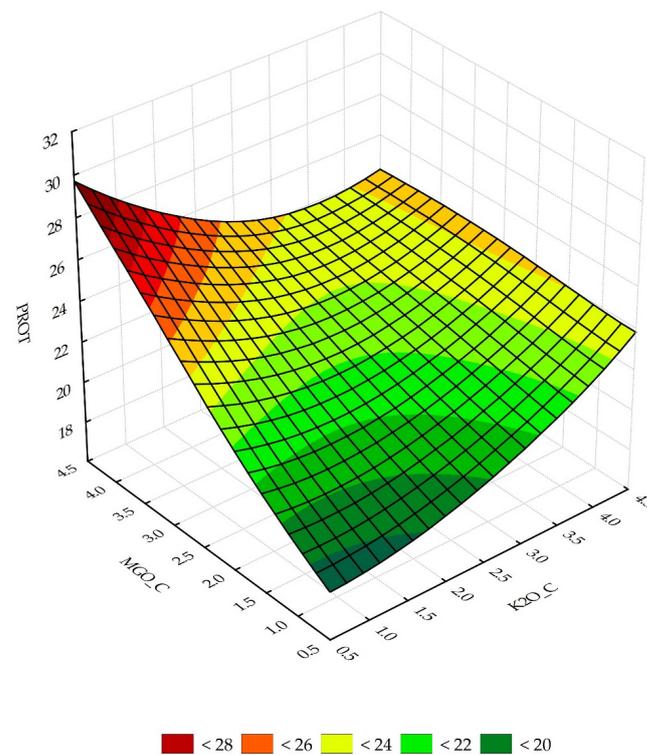


Figure 5. Response surface of the N1 model for percent protein and two dependent variables MGO_C and K2O_C.

Figure 6 shows the relationship of the independent variables (TEMP and P2O5_C) from the sensitivity analysis of the artificial neural network in relation to the dependent variable. From it, it can be observed that an increase in average daily temperatures during the pea growing season promotes the accumulation of protein in its seeds. This relationship also applies to the amount of phosphorus available in the soil. However, as the average daily temperature decreases, the high abundance of phosphorus in the soil does not result in the accumulation of protein in pea seeds.

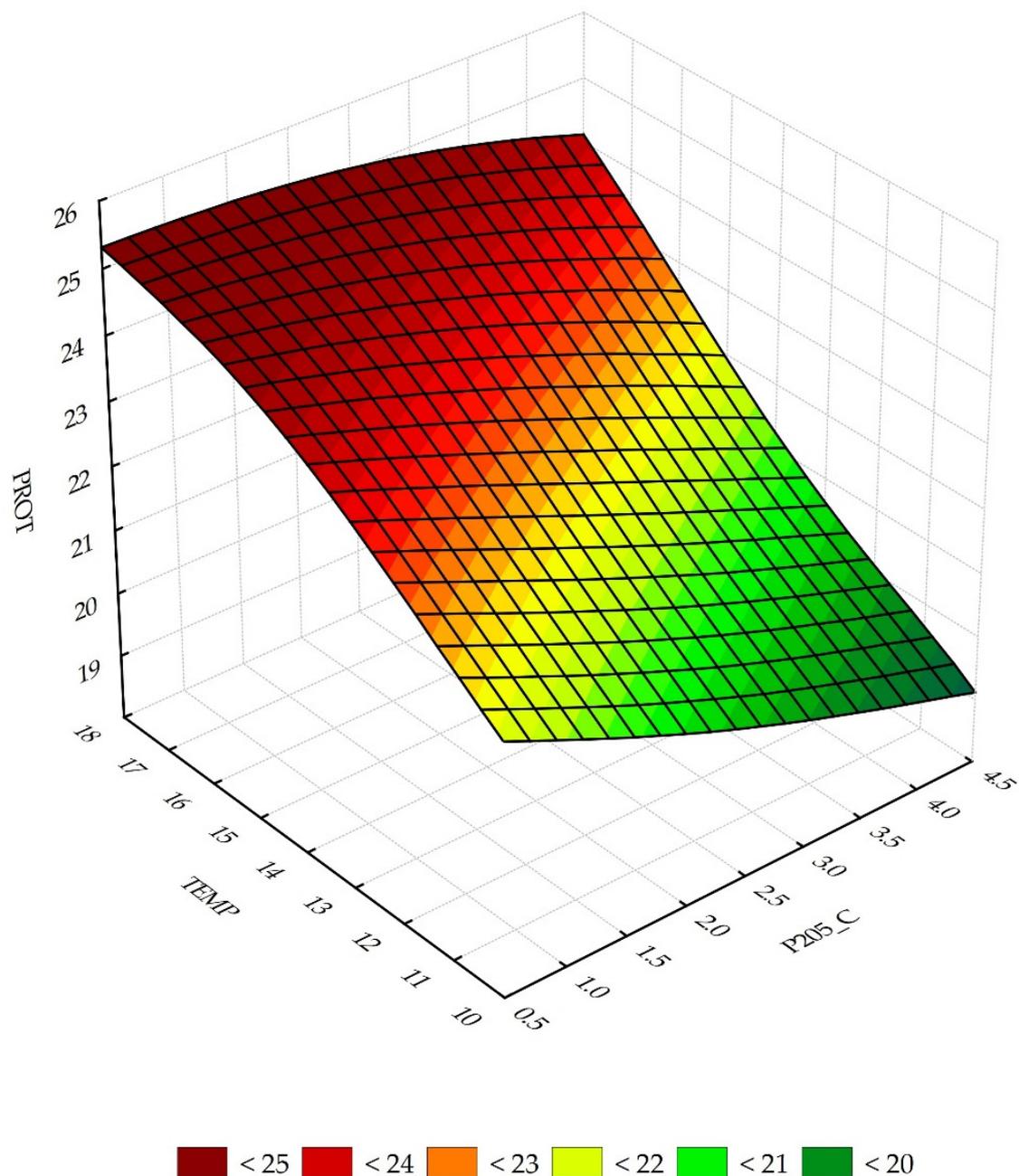


Figure 6. Response surface of model N1 for percent protein and two dependent variables TEMP and P2O5_C.

Changes in protein content as influenced by changes in average daily air temperature (TEMP) and soil magnesium abundance (MGO_C) are shown in Figure 7. An increase in the value of the independent variable TEMP at a low soil magnesium concentration caused the protein content of pea seeds to be less than 21%. A high MgO content in the soil

promotes protein synthesis in seeds only when there are sufficiently high average daily air temperatures during the pea growing season.

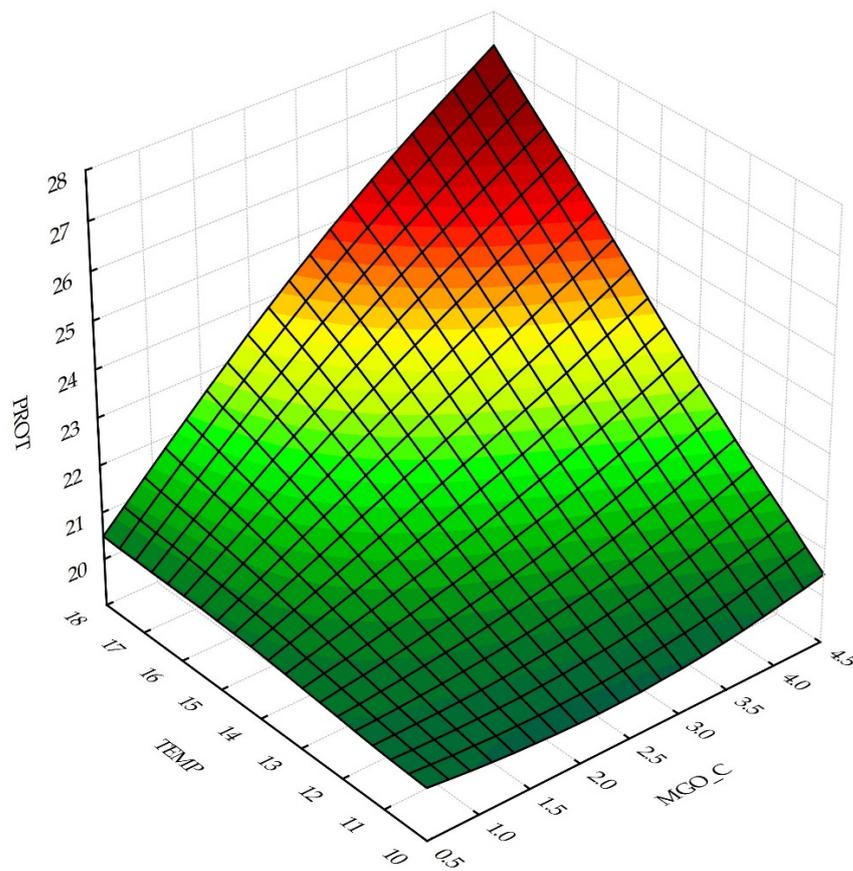


Figure 7. Response surface of the N1 model for protein percentage and two dependent variables TEMP and MGO_C.

3.3. Predictive Properties of the N1 and RS Model

Verification of the correctness of the N1 model and RS model was carried out based on six quality measures: RAE (global relative error of model approximation), RMS (root mean square error), MAE (mean absolute error), MAPE (mean absolute percentage error), MAX (maximum error determined for the whole model) and MAXP (maximum percentage error). Ex-post analysis (Table 7) showed that the N1 model had smaller error values compared to the RS model. The RS model achieved an RMS error of more than three times that of the N1 model. There were similarly large disparities in MAE and MAPE error. The N1 model was also characterized by a more-than-two-times smaller maximum percentage error.

Table 7. Quality assessment of the N1 and RS model.

Error Type	N1 Model	RS Model
RAE	0.037	0.118
RMS	0.838	2.696
MAE	0.617	2.032
MAPE	2.721	8.852
MAX	2.977	7.943
MAXP	13.001	28.853

4. Discussion

An important advantage of artificial neural networks is their ability to model the complex non-linear relationships that occur in agricultural crops [59]. A network with MLP (multi-layer perceptron) topology was used to predict cotton yield. The network was built from four categories of variables: weather data, drought indices, crop vegetation indices and yield. The resulting model had a MAPE error of 1.35%, and the R^2 value was 0.88 [60]. In contrast, Abrougui et al. [32] predicted the yield of potato grown in Chott Meriem (Tunisia). The analysis showed that the best measures of prediction quality (MSE = 0.006, % error = 1.116%) were characterized by a model with a topology of two hidden layers, which consisted of eight neurons in each layer. ANNs have also been successfully used to predict the quality characteristics of plant products such as essential oil content [61], ferulic acid concentration including mycotoxins in wheat grains [62], changes in protein, gluten and water content of stored wheat grains [63] and free radical content of sunflower, palm and rapeseed oil [64]. Niedbała [65] predicted the yield of winter rapeseed (*Brassica napus* L.) using ANN as of 30 June. The study covered fields located in Poland, in the southern part of the Opole Province. The neural network with a MLP topology predicted the explained variable with a MAPE error of 9.43%. The obtained N1 model in the in-house work was also characterized by low prediction errors. For example, the MAPE error was 2.721, which, according to Peng et al. [66] testifies to the model's excellent degree of fit, as the error does not exceed 10%. When MAPE is in the range of 10–20%, the degree of model fit is good. A forecasting model that achieves a MAPE error of more than 30% should be rejected due to poor mapping of predicted values with actual values. A low MAPE error (7.203%) was also obtained in a study by Piekutowska et al. [25], where a neural network with a MLP topology was built to predict the yield of very early potato varieties 40 days before harvest. Prediction of yield and its quality traits before harvesting gains importance in the era of changing climatic conditions. The unstable course of weather during the growing season of crops makes the quality of the crop variable each year. Obtaining models to accurately forecast food production is crucial for policy making and managing national food security plans [67]. Artificial neural networks were also successfully used in wheat yield prediction. The MLP model obtained had an RMS error of 0.4237 [68].

Niazian et al. [30] predicted the essential oil content of ajowan (*Carum copticum* L.) using ANN and MLR. Field studies were conducted from 2014–2015 in central Tehran. Four phenological traits were used as input data. The selection of independent variables was preceded by simple correlation analysis. The study showed that the ANN model with two latent layers predicted essential oil content with a mean squared error of 0.23% and a mean absolute error of 0.14%. In addition, the authors' research showed higher performance of the ANN model compared to the MLR, which had an RMS error of 0.26% and an MAE error of 0.18%. The artificial neural network model also had a higher coefficient of determination value ($R^2 = 0.88$) compared to the multiple regression model ($R^2 = 0.74$). The results of our own study also showed the superiority of neural networks over MLR in predicting the protein content of pea seeds. These results are consistent with another study [36], where the performance of MLP and stepwise regression networks was compared in predicting the essential oil content of fennel (*Foeniculum vulgare* Mill.). A total of 11 independent variables were used to build the neural network. The resulting MLP model with a topology of 11:11-9-7-1:1 was characterized by a coefficient of determination of 0.953 and 0.929 for the training and test set, respectively. The stepwise regression model, on the other hand, was characterized by an R^2 magnitude of 0.553. The neural network was additionally characterized by lower prediction errors compared to the MLR. The RMS and MAE error for the ANN were 0.544 and 0.385, respectively, while for the stepwise regression, the magnitudes of these errors were obtained at the level of 0.819 and 0.624. As can be seen from the data in Table 5, the N1 model predicted the protein content of peas with an RMS error of 0.838 and an MAE error of 0.617. The values of these errors are smaller than those obtained using the RS model, demonstrating the greater effectiveness of artificial neural networks over multiple regression in predicting the issue under analysis.

The advantage of artificial neural networks over classical regression modeling is due to the ability of ANNs to approximate non-linear functions [33]. In agricultural crops, many relationships between the analyzed variables have a complex and non-linear course, and MLR models are capable of predicting linear phenomena. Therefore, MLR cannot explain complex nonlinear relationships between independent variables and the dependent variable [69]. In addition to artificial neural networks, random forest regression (RFR) has also been used in agricultural science. Machine learning tools were built to predict the yield of winter rapeseed. The results obtained showed better predictive ability of the RFR model compared to the ANN model [33].

The sensitivity analysis performed for the ANN model showed that the protein content of peas was most affected by soil minerals (Mg, K, P). These variables received rankings one, two and three, respectively (Table 6). Interestingly, the response of pea plants, in terms of protein content, was greater for soil richness in these components than for mineral fertilization. Soil micronutrient and macronutrient abundance critically affects plant growth and development, as well as the quality of the yield obtained [70]. Plants respond less to current mineral fertilization than to high soil nutrient abundance [71]. Yano and Kume [72] noted, in pot experiments, an increased growth of corn root length on soil with a locally elevated phosphorus content. Low root mass increased the efficiency of phosphorus uptake per unit root weight with a low consumption of photosynthetic products.

Magnesium (Mg) received a rank of one in the neural network sensitivity analysis conducted. This means that this element had the greatest impact on the protein content of pea seeds. Mg is one of nine key plant nutrients. It is used in large quantities by plants for proper growth, development and reproduction [73]. Mg has many important physiological functions. It is an essential component of chlorophyll [74], and is involved in CO₂ assimilation reactions in the chloroplast [75]. Magnesium, like potassium (K), is essential for protein biosynthesis and significantly affects the absorption, utilization and metabolism of nitrogen (N) in plant roots [76]. For example, in soybean (*Glycine max* (L.) Merr.), nitrate uptake by the root system was influenced by Mg²⁺ and K⁺ ions through the regulation of the NRT2 transporters [77]. In a study conducted by Geng et al. [78], it was shown that rapeseed fertilized with magnesium exhibited increased nitrogen uptake at all fertilization levels (from 0 to 45 kg Mg·ha⁻¹). Much of the Mg contained in leaves appears to be directly or indirectly related to protein synthesis, due to its role in nitrogen metabolism and in the structure and function of ribosomes [76,79,80]. Ribosomes are macromolecular structures that are responsible for protein biosynthesis [81].

A four-year field study conducted in Croatia on six soybean varieties showed that foliar application of magnesium resulted in an increase in protein content in addition to an increase in seed yield. These differences, relative to the control, were statistically significant at the significance level of $\alpha = 0.05$ [82]. In contrast, a study by Sawan et al. [83] investigated the effect of potassium fertilization on cotton protein yield. The study was conducted in Giza, Egypt and covered two growing seasons. The results showed that protein yield significantly increased in plots where potassium was applied (47.4 kg·ha⁻¹) compared to the control, where plants were not fertilized with this element. Similar results were obtained in the present study. A higher Mg content in the soil caused an increase in the protein concentration in pea seeds (Figure 3). A similar relationship was observed for potassium (this variable received a rank of two in the network sensitivity analysis). However, the increase in protein content as a result of higher soil abundance of this element was not as great as in the case of magnesium.

Soil phosphorus content and average daily air temperature were the variables that received ranks three and four, respectively, in the sensitivity analysis (Table 6). Interestingly, TEMP received the highest rank among the weather variables. However, in order for this development not to be disturbed, air temperatures should remain at optimal levels throughout the growing season. Each species has an optimal temperature range for its development, while an excessive decrease or increase can contribute to plant damage [84]. Research conducted by Walter et al. [85] from 2016–2018 show that air temperature was

positively correlated with the protein content of pea and bean (*Vicia faba* L.) seeds grown in Germany. Higher temperatures were associated with higher protein concentrations in both pea and broad bean seeds. These results correlate with the data presented in this paper (Figure 4), where higher protein concentrations were recorded in peas as the average daily temperature increased. The increase in seed protein content was also associated with a higher soil phosphorus content. However, this effect was smaller the lower the daily temperatures were. Low temperatures result in reduced P uptake by plants. When the air temperature is less than 12 °C, the uptake of this element by the root system is largely blocked [86].

Phosphorus is an integral part of cell membranes and nucleic acids and is directly involved in protein synthesis [87]. Therefore, a close relationship between plant P nutrition and various physiological and biochemical traits is frequently and widely reported in the literature [88–91]. The proper nutrition of plants with this component leads to an increase in protein nitrogen and nitrogen of essential amino acids. However, the expected effect can be variable: positive (increase in protein content), neutral or negative (decrease in protein content following the so-called dilution effect). The negative effect results from an increase in usable yield, but in quantitative terms (e.g., for protein yield) it is positive [92].

The results concerning the significance of independent traits generated by using the models do not always coincide. The network sensitivity analysis showed that nitrogen fertilization of pea plants had a negligible effect on seed protein content. However, stepwise regression showed that this variable had a statistically significant effect on the dependent variable under study. Nitrogen is a macronutrient that significantly affects protein synthesis by plants [93]. However, according to a study conducted by Faligowska et al. [17], nitrogen in pea seeds grown in Brody (Poland) was accumulated from three main sources: soil, atmosphere and fertilizers. The largest amount of accumulated nitrogen came from the soil (57.9%), followed by the atmosphere (35.2%) and fertilizers (6.8%). The analysis of the multiple regression results confirmed the well-known state of knowledge about the relationship between nitrogen fertilization and protein content. However, the resulting neural network highlighted less well-known and much more interesting aspects of certain relationships. Relationships between given factors in agricultural production are determined mainly by weather conditions prevailing in a given growing season, habitat conditions and genotypes of crops grown.

5. Conclusions

The developed MLP-type artificial neural network model successfully predicted the protein content of pea seeds 20 days before the harvest date. The model had higher prediction accuracy and lower ex-post error values with respect to the stepwise regression model. The accuracy of MLP networks is mainly dependent on the accuracy of the data at one's disposal and the amount of information fed into the model. In the case of artificial neural networks, an important task is to maintain a balance between the model's ability to approximate and generalize. The analysis conducted showed that the network accurately predicted the dependent variable based on a five-year field study. The results obtained did not allow the rejection of the null hypotheses, thus confirming the validity of the assumptions made at the outset. The N1 model, due to its lower error values and higher R^2 value, more accurately predicted the protein content of peas. For this reason, this model seems to be better in practical application.

This work can serve as a strand for future research on the prediction of pea seed quality such as the content of fat, nitrogen-free compounds, the largest part of which in pea seeds is starch and anti-nutritional compounds. A continuation of the present research will be the optimization of NPK fertilization taking into account the liming needs of general-purpose pea varieties.

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Abbreviations

ANN-artificial neural networks; CCA-Community Catalogs of Agricultural; CCV-Community Catalogs of Varieties of Vegetable; COBORU-Research Center for Cultivar Testing; DUS-distinctiveness, uniformity and durability; FLOWE-number of days from 1 January to the beginning of flowering; GEN-general variety of peas; HAR-number of days from 1 January to the date of harvesting; INI_MA-number of days from 1 January to onset of maturity; K2O_C-K2O content in the soil; K2O-potassium oxide; K2O_F-Total potassium from mineral fertilizers; kg-kilogram; MAE-mean absolute error; MAPE-mean absolute percentage error; MAX-maximum error determined for the whole model; MAXP-maximum percentage error; mg-miligram; MgO-magnesium oxide; MGO_C-MgO content in the soil; MLP-multilayer perceptron; MLR-multiple linear regression; MO-organic matter; n-number of observations; N_F-total nitrogen from mineral fertilizers; N1-built its own neural network model; P_EMER-number of days from 1 January to the beginning of plant emergence; P_HIG-plant height; P2O5-phosphorus(V) oxide; P2O5_C-P2O5 content in the soil; P2O5_F-total phosphorus from mineral fertilizers; PH-Soil pH; PROT-Percentage of protein in pea seeds; PRVT-Program of Registered Varietal Experimentation; RAE-global relative error of model approximation; RAIN-total rainfall from sowing date to 14 July; RMS-root mean square error; RS-built its own linear regression model; SOWI-number of days from 1 January to sowing date; SUN-total sunshine from sowing date to 14 July; TECH_M-number of days from 1 January to technical maturity; TEMP-average air temperature from sowing date to 14 July; VCU-trials for cultivation and use value; WEGW-number of plant growing days; y^i -predictive values, obtained with the model; y_i -actual values.

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