Predicting Nutrient Density in Foods Using Machine Learning Models: A Comparative Study

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Abstract: Fine-tuning and thus accurately estimating nutrition density in foods is useful in optimizing diets and

improving health standards. Several challenges have been observed with the traditional methods for nutrient evaluation. Most of these challenges can be trimmed down by adopting the use of Machine Learning (ML) models, which possess better capabilities of giving efficient and accurate assessments. To this end, different regression models were applied to estimate nutrient density, namely Linear Regression, Ridge Regression, Decision Trees, and Random Forests. The used set had 2397 food items for which 33 nutrients had been deemed relevant. The missing values in the chosen dataset were addressed before model training through imputation and normalization for better data quality. The models were trained and evaluated using separate training and test sets, with performance indicators such as Mean Absolute Error (MAE) and R-squared (R²) used to measure their accuracy. Results showed that linear models, such as Linear Regression and Ridge Regression, achieved the best accuracy, with an R² of 0.999, while tree-based models exhibited overfitting tendencies, resulting in lower predictive performance on unseen data. These findings demonstrate the effectiveness of machine learning in predicting nutrition density, significantly improving the precision of

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1 INTRODUCTION

Dietary habits are now considered to be a vital part of health promotion and maintenance, especially in the etiology and prevention of chronic communicable diseases such as obesity, type-2 diabetes, hypertension, and atherosclerosis associated with unhealthy diets (Drewnowski, 2009). An example is the laboratory analysis and nutrition databases that are employed in the traditional approach to evaluate nutritional quality. In response to these problems, there is increasing concern about artificial intelligence for improved assessments of nutrition density in foods due to their superior performance (Drewnowski et al., 2019). Machine learning (ML), especially, shows great potential advancements in delivering helpful recommendations to individuals, thus helping them make the right decisions regarding their choice of foods to take (Drewnowski, 2019).

The learning capability, intrinsic to machine learning algorithms, has been illustrated with several

works on many tasks, and the same applies to nutritional science. Conventional approaches seem to have their limitations in this area, so it only makes sense to attempt to use the relatively more advanced machine learning technique. With big data, many factors that are not easily observable or even recognizable may be modeled and analyzed to predict Nutrition Density more efficiently than in systems that employ traditional analytics (Drewnowski, 2005). Explaining how nutrition density scores balance overcrowded food product health claims with Henley's health-supporting mission, Drewnowski best encapsulates the beneficial relationship between health and food. Subsequent studies have established that those machine learning algorithms can make probable an undertaking of nutrition density by evaluating several nutritional parameters including fat content, sugar content, and vitamin and mineral contents (Shen et al., 2020). Armand et al. (Armand, 2024) described an inherent capability of ML within nutrition science to transfigure the field through the measurements of nutrition density and the provision of personalized advice.

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Furthermore, due to their capacity to work with big and intricate data sets, the ML models are especially useful for nutritional analysis, where more conventional approaches might fail to capture the associations of the different nutrients. For instance, Lucas Prado Osco et al. (Osco, 2020), were able to utilize machine learning to determine nutrition value content in agricultural produce and this is a further example of the application of these techniques. Also, Timsina et al. (Timsina, 2021) presented how ML has been useful in enhancing nutrition management in the agriculture field; therefore, valid in the food and health industries.

For an effective and precise estimation of nutrition density, Linear Regression, Regression, Ridge Regression, Decision trees, Random Forest, and AdaBoost models are chosen in this study. These models have been selected to illustrate that they are distinct from each other, from linear models through different types of ensemble techniques. These linear models were particularly effective when it came to the modeling of the relations between features and nutrition density. In assessing the performance of each model, specific evaluation metrics were employed including Mean Absolute Error (MAE), Mean Squared Error (MSE), and R-squared (R2) to get a full analysis of the models' prediction potentials.

2 METHOD

2.1 Dataset Preparation

The dataset (Dey, 2024) used in this study was meticulously designed to capture the detailed nutritional profile of various food items, providing a comprehensive foundation for predicting nutrition density. The dataset includes 36 columns, each representing a specific nutrient or related metric essential for understanding the nutritional content of the foods. These columns cover a wide range of essential macronutrients, micronutrients, vitamins, and minerals, which are critical for evaluating the overall nutritional value of each food item.

The dataset consists of 2,397 rows, each corresponding to a unique food item. To ensure robust model training and evaluation, the dataset was divided into two subsets: a training set and a test set. The training set, containing 1,674 rows, was used to train the machine learning models. The test set, comprising 723 rows, was reserved for evaluating the performance of the trained models. This separation of data allows for an unbiased assessment of the model's

predictive capabilities, ensuring that the models generalize well to new, unseen data.

The dataset includes the following key columns:

- Caloric value: This column captures the total energy provided by the food, measured in kilocalories (kcal) per 100 grams. It serves as a baseline for comparing nutrition density relative to energy content.
- Carbohydrates: This includes both total carbohydrates and sugars, highlighting the presence of simple sugars, which can significantly impact health outcomes.
- Vitamins and minerals: The dataset includes detailed information on a wide range of vitamins (A, B1, B11, B12, B2, B3, B5, B6, C, D, E, K) and minerals (Calcium, Copper, Iron, Magnesium, Manganese, Phosphorus, Potassium, Selenium, Zinc), each measured in milligrams per 100 grams. These nutrients are vital for various bodily functions, from bone health to immune support.

Last but not least, the Nutrition Density column is the last column that serves as the nutrition richness of the food per calorie. This column is significant for the analysis as it goes to the heart of the study's purpose which is to predict nutrition density.

Again, before engaging in the analysis, the dataset went through some basic exploratory data analysis steps to clean the data. For the handling of missing values, the proper imputation methods were used to fill in the missing values and normalize the data set was used to transform all the features into one convenient scale. Such a preprocessing enables the models to learn from alterations in the data without being influenced by differences in feature scales, or lack of missing values.

During the model training, the 'Food', 'Identifier', and 'Nutrition Density' feature columns were omitted from the features because they are not themselves predictors of nutrition density and are categorical variables for food identification and target variables respectively. This led to the use of 33 features, in total for developing the model used for training the classifier. The rest of the columns given the data offered the inputs to the machine learning models while the Nutrition Density served as the output.

Cutting the dataset into the training and testing sets was done to enable a conclusive assessment of the model's accuracy. This means, the study was able to keep a portion of the data for testing and could be able to determine whether the model was overfitting on the training data or it could perform well on foods that were not imported into a model. This is common

in machine learning-oriented practices and it is very important when it comes to creating models that generalize to other situations.

2.2 Machine Learning Models-based prediction

2.2.1 Linear Regression Models with Variants

Linear Regression: Linear Regression is used as a baseline model and notwithstanding its eccentricity in finding linearity between the features and the target variable it is simple. Although it might be less accurate than other models of higher complexity, it is valuable for its interpretability of the contributions of the features. It is stated by Yao et al. (Yao, 2013) that there are other methods such as using modal regression; however, using Linear Regression is inevitably simpler and more transparent while obtaining shorter prediction intervals, even in cases where the distribution is skewed.

Lasso Regression: Lasso Regression, uses L1 regularization hence solving the problem of data overfitting, and does feature selection by setting coefficients to zero. This method is especially useful in big data, especially in a situation where there are many independent variables to consider since it reduces the possibility of making large prediction errors and provides a measure of checking the model's complexity. According to Ranstam et al. (Ranstam, 2020), even as it imposes potential bias in estimating individual parameters, Lasso lends itself to be an effective technique for obtaining high overall accuracy in the prediction which is more desirable when working with large numbers of predictors.

Ridge Regression: Ridge Regression makes use of the L2 regularization to handle the issue of multicollinearity between the features to ensure more accurate computation of coefficients that leads to better predictions. They include genetic data analysis where the number of predictors surpasses the observations, thus making it suitable for high dimensions. By removing mean-square error for the correlated predictors, as noted by Cule (Cule, 2013), Ridge Regression can be used in cases where the basic regression can be non-applicable while still offering good predictive quality and stability.

For these regression models, the primary hyperparameters to tune were the regularization strength (denoted by alpha). Grid Search Cross-Validation was employed to explore a range of alpha values to determine the optimal regularization parameter. This process involved systematically testing multiple values of alpha and selecting the one that minimized the validation error.

Grid Search: The Grid Search method exhaustively searches over a specified parameter grid for each model. For instance, in Ridge and Lasso Regression, various alpha values were tested to find the one that best balanced model complexity and prediction accuracy.

Support Vector Machine (SVM) Regression: SVM Regression examines possible non-linearity of relationships between dependent and independent variables by transforming the input data set by various kernels. It is an advantage to provide high accuracy, and at the same time, generalization was pursued for data with many features for the number of cases. However, as stated by Pisner (Pisner, 2020), overfitting is common with SVM but it is still very useful in many regression problems that pose both linear and non-linear curvatures.

The SVM regression model required tuning of the C parameter (regularization), the kernel type (e.g., linear, polynomial), and the gamma parameter (for non-linear kernels). Again, Grid Search Cross-Validation was used to explore combinations of these parameters.

Grid Search: The grid search for SVM involved testing different kernel functions and their respective parameters (e.g., C and gamma). The combination that provided the best cross-validation performance was selected as the optimal model configuration.

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2.2.2 Tree-based Model

Decision Tree: Decision Tree models work for data with non-linear relations as segmentation of variables forms a tree, thus, classifying the data. This is a non-parametric approach technically capable of handling large-sized data and missing values without requiring strenuous assumptions. According to Song (Song, 2015), Some of the common uses of Decision Trees include variable selection, testing of variable importance, and making of forecasts which are reasons why this method is prevalent, especially for

medical research that requires simple models that are easy to understand.

For the Decision Tree model, the hyperparameters tuned included the maximum depth of the tree, the minimum samples required to split a node, and the minimum samples required at a leaf node. These parameters control the complexity of the tree and prevent overfitting.

Grid Search: Grid Search was applied to identify the best combination of these parameters. By varying the maximum depth, minimum samples split, and minimum samples leaf, the grid search determined the configuration that yielded the best validation performance.

2.2.3 Ensemble Models

Random Forest: The decision tree is an individual model; Random Forest is an advanced method that combines many decision trees through an average that minimizes the risk of over-fitting. It is a complex calculation and not easy to decipher, but when it comes to large and interaction-riddled data as well as non-parametric analyses, it stands out. While Random Forest may be known as the "black box", according to Rigatti (Rigatti, 2017), the abilities to model non-linear effects make it important for many predictive tasks.

The Random Forest model required tuning the number of trees, the maximum depth of each tree, the minimum number of samples required to split a node, and the number of features to consider when looking for the best split.

Randomized Search: Instead of an exhaustive grid search, a Randomized Search Cross-Validation was employed for Random Forest due to the larger parameter space. This method samples a fixed number of parameter settings from the specified distributions and tests them, making it more efficient for models with numerous hyperparameters.

AdaBoost: AdaBoost is an ensemble model functioning as an enhanced weak learner through weight assignment based on misclassifications of instances. It transforms weak predictors into strong models and it is among the best algorithms in data mining since it has an impact on other learning algorithms, according to Cao et al. (Cao, 2013). This reconstruction based on the scores has made AdaBoost to be a popular and successful tool in machine learning.

For AdaBoost, the key hyperparameters tuned were the number of estimators (i.e., the number of weak learners to combine) and the learning rate (which controls the contribution of each weak learner). These parameters were optimized to improve the ensemble's accuracy and reduce overfitting.

Grid Search: Like other models, Grid Search Cross-Validation was used for AdaBoost. The method tested different values for the number of estimators and the learning rate to find the combination that resulted in the best model performance.

2.3 Evaluation Metrics

To ensure the accuracy and reliability of the models in predicting nutrition density, four key validation metrics were employed. These metrics assess different aspects of prediction accuracy and error distribution, providing a comprehensive evaluation of each model's performance. The validation was performed on the test set, which contains data not seen by the models during training. The metrics used are as follows:

$$MAE = \frac{1}{n} \sum_{i=1}^{n} |y_i - \widehat{y}_i| \tag{1}$$

$$MSE = \frac{1}{n} \sum_{i=1}^{n} (y_i - \hat{y}_i)^2$$
 (2)

$$RMSE = \sqrt{\frac{1}{n} \sum_{i=1}^{n} (y_i - \hat{y}_i)^2}$$
 (3)

$$R^{2} = 1 - \frac{\sum_{i=1}^{n} (y_{i} - \hat{y}_{i})^{2}}{\sum_{i=1}^{n} (\hat{y}_{i} - y_{i})^{2}}$$
(4)

3 RESULT AND DISCUSSION

The Linear Regression, Ridge Regression, and SVR models demonstrated excellent performance across all metrics shown in Table 1. These models achieved an R² value of 0.999, indicating nearly perfect predictions on the test set. The MSE and RMSE values for these models were extremely low (MSE: 0.002, RMSE: 0.040 for both Linear and Ridge Regression), further confirming their accuracy. These results suggest that linear models and SVR are highly effective in capturing the relationship between the features and nutrition density, making them reliable choices for this predictive task.

	Train MSE	Test MSE	Train R ²	Test R ²	Train MAE	Test MAE	Train RMSE	Test RMSE
Linear regression	0.001	0.001	0.999	0.999	0.018	0.022	0.033	0.039
Lasso Regression	133.890	26.416	0.996	0.997	5.432	3.163	11.571	5.139
Ridge Regression	0.001	0.001	0.999	0.999	0.018	0.022	0.033	0.039
SVM	543.093	774.441	0.985	0.939	9.855	12.859	0.033	27.828
Decision Tree	1396.349	424.893	0.961	0.966	4.918	7.515	23.304	20.612
Random Forest	3288.969	2766	0.910	0.783	43.877	41.752	57.349	52.597
Adaboost	0.006	0.004	0.999	0.999	0.053	0.051	0.079	0.069

Table 1: Model Performance Metrics Summary

On the other hand, the Decision Tree and Random Forest models, while performing well on the training data, exhibited signs of overfitting. The Decision Tree model, for example, had a significantly higher test MSE of 774.441 and an R² of 0.939, which, although still relatively high, indicates a drop in performance compared to the linear models. Similarly, the Random Forest model had a test MSE of 424.893 and an R² of 0.967. These figures suggest that while tree-based models can capture complex interactions within the data, they may struggle with generalization, particularly when not properly tuned.

The AdaBoost model struggled significantly in this application. It had the highest test MSE of 2766.445 and the lowest R² value of 0.783 among all models tested. The MAE and RMSE were also much higher compared to other models, indicating that AdaBoost was not well-suited for predicting nutrition density in this dataset. This poor performance could be due to AdaBoost's sensitivity to noisy data or its tendency to overfit, especially in the absence of strong individual learners.

The success of machine learning in predicting nutrition density in fruits is supported by similar studies, such as the prediction of fiber content in Australian packaged foods using the k-nearest neighbors (KNN) algorithm. This study found that KNN significantly outperformed manual prediction methods with an R² value of 0.84, highlighting the potential of machine learning to predict important nutritional metrics efficiently, according to Davies et al. (Davies, 2021). In research, models like Linear Regression and Ridge Regression achieved even higher accuracy, demonstrating that machine learning can be a robust tool for various nutritional predictions.

Interestingly, this work reveals that simple models: Linear Regression, Ridge Regression, and

SVR have higher accuracy rates than complex models: Decision Tree, Random Forest, and AdaBoost for nutrition density. This may have caused a higher performance of these models because a linear flow of data makes it easy for models not to overfit while training by establishing the relationship that exists between features and the target variable. It is quite evident that these models have great generalization ability, especially by looking at how well they have performed on both training and test sets.

On the other hand, the Tree Models, much as they are capable of handling nonlinearity and interaction, had the vice of overfitting the data. This is usually the case when models learn noise within the training data and not the repeated patterns thus affecting the generalization of new data. The AdaBoost model has shown poor results, which can be attributed to the relative sensitivity to noise and the difficulty of combining several weak learners.

Therefore, this study has shown that different models' selection should be determined and guided by the characteristics of the given dataset. Compared to more complex models, the simpler ones might perform better, especially when the linear relationship prevails between the features and the target variable. In more complex models there can be the problem of overfitting, which reduces the ability of these models to be effective. One could take up future research on the hybrid approaches or the incorporation of more features to develop better the performance of nonlinear models in similar predictive tasks.

4 CONCLUSIONS

This research was able to use machine learning models to accurately predict nutrition density in food, and thus show that ML could be of great value in boosting the analysis of diets. Regression analysis, especially linear regression, as well as ridge regression, decision trees, Random forests, and other models, were used to compare and contrast various factors to nutrition density. Based on the obtained experimental results, it was found that Linear Regression models including Linear Regression and Ridge Regression exhibit the highest accuracy with the value of R² of a nearly perfect degree of 0.999. It also shows that machine learning is more useful in dealing with a massive dataset and is generally reliable in enhancing the forecasted probability as compared to the simple nutritional analysis which took a lot of time and resources.

Therefore, the research outcomes show the applicability of machine learning algorithms for determining nutrition density so that better and evidence-based dietary advice could be given. However, it also emerged that Decision Trees and Random Forest others may encounter problems such as overfitting which infers that further tuning or perhaps the use of a hybrid model could be considered. Subsequent studies may also look at extending the existence of other features or the raw employ of superior models, to enhance the exactness of predictive operations as well as the reductions in nonlinear data sets.

REFERENCES

- Armand, T., Kintoh Allen Nfor, Kim, J.-I., & Kim, H.-C. 2024. Applications of Artificial Intelligence, Machine Learning, and Deep Learning in Nutrition: A Systematic Review. Nutrients, 16(7), 1073–1073.
- Cao, Y., Miao, Q.-G., Liu, J.-C., & Gao, L. 2013. Advance and Prospects of AdaBoost Algorithm. Acta Automatica Sinica, 39(6), 745–758.
- Cule, E., & De Iorio, M. 2013. Ridge Regression in Prediction Problems: Automatic Choice of the Ridge Parameter. Genetic Epidemiology, 37(7), 704–714.
- Davies, T., Louie, J. C. Y., Scapin, T., Pettigrew, S., Wu, J.
 H., Marklund, M., & Coyle, D. H. 2021. An Innovative Machine Learning Approach to Predict the Dietary Fiber Content of Packaged Foods. Nutrients, 13(9), 3195
- Dey, U. 2024. Food Nutrition Dataset. Kaggle.com. https://www.kaggle.com/datasets/utsavdey1410/food-nutrition-dataset/data
- Drewnowski, A. 2005. Concept of a nutritious food: toward a nutrient density score. The American Journal of Clinical Nutrition, 82(4), 721–732.
- Drewnowski, A. 2009. Defining Nutrient Density: Development and Validation of the Nutrient Rich Foods Index. Journal of the American College of Nutrition, 28(4), 421S426S.

- Drewnowski, A. 2019. Impact of nutrition interventions and dietary nutrient density on productivity in the workplace. Nutrition Reviews, 78(3), 215–224.
- Drewnowski, A., Dwyer, J., King, J. C., & Weaver, C. M. 2019. A proposed nutrient density score that includes food groups and nutrients to better align with dietary guidance. Nutrition Reviews, 77(6), 404–416.
- Lucas Prado Osco, Paula, A., Pinheiro, F., Saito, A., Nilton Nobuhiro Imai, Nayara Vasconcelos Estrabis, Ianczyk, F., Fernando, Veraldo Liesenberg, Jorge, Li, J., Ma, L., Wesley Nunes Gonçalves, José Marcato, & José Eduardo Creste. 2020. A Machine Learning Framework to Predict Nutrient Content in Valencia-Orange Leaf Hyperspectral Measurements. Remote Sensing, 12(6), 906–906.
- Pisner, D. A., & Schnyer, D. M. 2020. Chapter 6 Support vector machine (A. Mechelli & S. Vieira, Eds.). ScienceDirect; Academic Press.
- Ranstam, J., & Cook, J. A. 2016. Overfitting. British Journal of Surgery, 103(13), 1814–1814.
- Rigatti, S. J. 2017. Random Forest. Journal of Insurance Medicine, 47(1), 31–39.
- Shen, Z., Shehzad, A., Chen, S., Sun, H., & Liu, J. 2020.

 Machine Learning Based Approach on Food Recognition and Nutrition Estimation. Procedia Computer Science, 174, 448–453.
- Song, Y.-Y., & Lu, Y. 2015. Decision tree methods: applications for classification and prediction. Shanghai Archives of Psychiatry, 27(2), 130–135.
- Timsina, J., Dutta, S., Devkota, K. P., Chakraborty, S., Neupane, R. K., Bishta, S., Amgain, L. P., Singh, V. K., Islam, S., & Majumdar, K. 2021. Improved nutrient management in cereals using Nutrient Expert and machine learning tools: Productivity, profitability, and nutrient use efficiency. Agricultural Systems, 192, 103181.
- Yao, W., & Li, L. 2013. A New Regression Model: Modal Linear Regression. Scandinavian Journal of Statistics, 41(3), 656–671.