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A Robust Approach to Uncertainty Quantification in Deep Learning

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Abstract

This study proposes a novel approach for quantifying the uncertainty of a deep learning model by investigating the coverage as well as the adaptivity of its prediction intervals in a Conformal Prediction context. The model investigated is designed *to impute the equivalent household income by taking both specific household group characteristics and relevant features of the main income gainer into account as it is known that there are well-known correlations in literature. The imputation of such variable is critical as outliers occur or the required information for computing it is not entirely available. Due to the relevance of income in socio-economic policy contexts, the reliability of its imputation constitutes a key aspect. The Conformalized Quantile Regression is adopted in order to evaluate the prediction intervals of the model by incorporating this approach into the same. In this study an improved assessment of the model uncertainty is achieved by separating the aleatoric component from the epistemic one. For this purpose, an appropriate selection of training data is proposed. This non random selection introduces bias which may alter model estimates causing distortions which impair the uncertainty quantification approach. As a consequence, a correction of selection bias is integrated in the uncertainty evaluation process. A real-world case study is considered to demonstrate the potential of the proposed quantification approach.*

Keywords: Uncertainty Quantification, Deep Learning, Income Imputation, Conformalized Quantile Regression, Heckman Selection

1. Introduction

In recent years the topic of income imputation has gained attention in economic research, especially when it comes to estimating the measurement of the equivalent household income [1]. This imputation is a challenging problem in general, due to the inherent uncertainty in socio economic data of which it is composed. This measure is designed for rendering the evaluation of household economic well-being more accurate by adjusting the total household income in accordance with the size of the household and relationships between its members, while keeping the economies of scale in mind as well as the different needs of the members [2]. Equivalent household income makes the comparison of different househod structures possible in that it detects various economic aspects which standard metrics might overlook [3]. Imputation techniques, such as regression-based or statistical matching methods, are not robust enough when it comes to managing incomplete or missing income data; therefore it is not insured that income distributions reflect realistic values in relation to known demographic groups [4]. Nevertheless, challenges persist, as the imputation process may introduce biases or inaccuracies that affect findings on income inequality, poverty rates, and related social policies [5]. As a result, accurate imputation remains critical for reliable socio-economic analyses and the development of equitable welfare policies. The inevitable

existence of the aleatoric uncertainty pertaining to equivalent household income components may impair the point estimation process even in the case of a robust approach. An appropriate construction of input dataset for training the model is proposed [6,7] in order to enhance the epistemic aspect of the uncertainty under investigation. This specific construction may introduce a selection bias which results in distorted income estimates. As a consequence, in this study the Heckman correction is integrated into the aforementioned model [8]. The objective is to leverage deep learning methodologies in order to propose a robust procedure for evaluating the prediction intervals of a model for income imputation by means of the Conformalized Quantile Regression (CQR) from the Conformal Prediction framework [9-12]. A case study based on the statistical register by the name of ARCHIMEDE from ISTAT is considered in order to test for the potential of the uncertainty quantification technique proposed.

2. Basic Algorithms of a Deep Learning Model

Deep learning is a specific branch of machine learning which avails of artificial neural networks for complex data patterns modelling in large-scale datasets. Their general architecture provides multiple layers of connected neurons which manage complex problem solving in various fields of research.

The essential structures of the deep learning model under investigation are described below.

2.1. Denoising Autoencoder

The Denoising Autoencoder (DAE) is a type of artificial neural network designed for robust feature learning as well as removing noise from data. The DAE encodes both numeric vectors and vectors of categorical variables which has to be transformed into

Figure 1: Denoising Autoencoder **Figure 1.** Denoising Autoencouer

of numbers included in the interval [1]. This model is trained representation, detecting essential feature to reconstruct the original input observation x from a corrupted irrelevant noise. This compressed, noisered version \tilde{x} , of the same constituting an effective algorithm for is subsequently reconstructed to obtain a data denoising and feature extraction. A pre-defined noise the data, closely matching the uncorrupt function $\eta \sim \mathcal{N}(0, 1)$ is applied to input data to disturb it, forcing training process is analogous to other po the DAE to reconstruct $x = f(x)$ as the data is being disturbed. algorithms. In this case, the loss function A sketch of a simple one-layer architecture is shown in Fig. 1. error, defined as follows: denoising and feature extraction.
A pre-definition of the constant extraction noise function \sim N (0, 1) is applied no interaction of the const dummy variables so that the input data vectors become sequences The encoder compresses each input into a

The encoder compresses each input into a smaller, more abstract representation, detecting essential features while filtering out irrelevant noise. This compressed, noisereduced rappresentation is subsequently reconstructed to obtain a noise-free version of the data, closely matching the uncorrupted original input. The training process is analogous to other popular neural network algorithms. In this case, the loss function is the *reconstruction error,* defined as follows:

$$
\mathcal{L}(\tilde{\mathbf{x}}, \mathbf{x}) = \frac{1}{N} \sum_{i=1}^{N} (\mathbf{x}_i - \hat{\mathbf{x}}_i)^2
$$
(1)

is the reconstruction error, defined as follows: the DAE subsequent to the noise application to the original observation x. ates the reconstructed input observation by multiple layers of neurons arranged in a where $\hat{\mathbf{x}}$ indicates the reconstructed input observation by multiple layers of neurons \sum_{x} observation x. $\frac{1}{2}$

2.2 Multilayer Perceptron **2.2. Multilayer Perceptron**

neural network rather similar to that described in the previous MLP model $y = f(x)$ is trained by adj problems in data science. The basic structure is composed of the error between predicted outputs The Multilayer Perceptron (MLP) is another type of artificial activation functions to model complex Section even though it is versatile in solving a great number of connections between neurons based on the regression and classification problems. Its core architecture usually includes an input

problems. Its core architecture usually includes an input Perceptron (MLP) is another type of artificial activation functions to model complex patterns in data. The multiple layers of neurons arranged in a feed-forward structure, which is highly effective for regression and classification layer x, one or more hidden layers, and an output layer. Each layer consists of interconnected neurons that apply non-linear MLP model $y = f(x)$ is trained by adjusting the weights of connections between neurons based on the backpropagation of the error between predicted outputs

Figure 2: Multilayer Perceptron

 \hat{y} and actual outputs y. This error is usually optimized by minimize it. MLP training for regression pr implementing the stochastic gradient descent in order to requires the following loss

 $y \text{ is } y$. This error is usually optimized by minimize it. MLP training for regression problems usually requires the following loss function:

$$
\mathcal{L}(\tilde{y}, y) = \frac{1}{N} \sum_{i=1}^{N} (y_i - \hat{y}_i)^2
$$
 (2)

true value of the dependent variable y and its predicted value \hat{y} B contains the second households of the same $=f(x)$ in a supervised learning perspective. question $y = f(x)$ maps the changes in the which is defined as the *Mean Square Error* (MSE) between the datasets: A which contains the first household of

Setting up training bata
The selection process of training data is one of the innovative household group characteristics subject to registers require sophisticated approaches of analysis as is the similarly distributed. The idea behind this study case when detecting similar data structures in large scale datasets. two independent uncertainty evaluation proced To be more specific, different groups of individuals related the model with the same data (similar household one to another for kinship and utility reasons, *i. e.* household contribution of the random component of the day structures, deeply influence income dynamics. The equivalent this improves the accuracy of the uncertainty eventually household income of the OECD scale should take *economies* as increasing the robustness of the same by pr of scale into account without any disturbance of its predictors into model bias. In order to separate the aleat caused by aleatoric uncertainty present within them. As a result, of the uncertainty from the epistemic one, the aleatoric uncertainty in the data affects the estimation of the on the entire dataset A for the subsequent train equivalent household income by altering the economies of scale by using a random 80% partition of the *encode* and therefore hindering a reliable evaluation of the prediction the remaining data for evaluating the model. T intervals. As a consequence, an appropriate subset of the input repeated by swapping A and B in order to comparently are detected by means of the algorithm based on a Locality 4. Selection Bias Correction aspects of this study. Complex data structures in current statistical constraint. Data from both A and B sets dataset X is selected in order to facilitate this evaluation process. investigating epistemic uncertainty only. This subset is composed of pairwise similar households which Sensitive Hashing (LSH) approach [[?], [?]]. Deep learning model training is accomplished by processing this specific subset so that the Conformalized Quantile Regression which is $\frac{1}{2}$ integrated in the model estimates the prediction intervals in two different scenarios as is described in this Section. Households pairs belonging to this subset share the same size as well as in two stages. The first stage consists or the same attributes of their members. Different pairs involve estimating the probability of an observation different sizes and different attributes of their members. The the sample; essential for understanding detection algorithm transforms all the households belonging to This stage results in the *Inverse Mills Re* the input dataset into graphs and subsequently into sequences of of the deviation of the selection process fro integers which are elaborated by the LSH approach in order to IMR is defined as follows: detect the aforementioned pairs. These pairs are split into two

which is defined as the *Mean Square Error* (MSE) between the
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economies of scale into account without any disturbance of its predictors caused **4. Selection Bias Correction**

by alleated the must correction.
In this procedure known as *Heckman model* is a statistical approach the designed to address selection bias, which arises when the dataset economies of scale and the prediction of the prediction inter-
selected for model training is not a random sample in that it is value of the internal and input determining in the internal subset of the internal vielding model estimates tion as is described in this Section. Households which may be biased. The Heckman model is implemented estimating the probability of an observation being included into in two stages. The first stage consists of a *probit* model for the sample; essential for understanding the selection process. This stage results in the *Inverse Mills Ratio* (IMR), a measure of the deviation of the selection process from being random. The IMR is defined as follows:

$$
IMR = \frac{\phi(\mathbf{x}'\beta)}{\Phi(\mathbf{x}'\beta)}
$$
(3)

where ϕ (•) denotes the probability density function of a standard intervals which are guaranteed to cover normal distribution, $\Phi(\cdot)$ rappresents its cumulative distribution pre-defined probability. The *Conformalize* function, *x* is the vector of independent variables and β is the (CQR) , is a method within this framev vector of coefficients estimated by the probit model. The IMR the quantile regression and conformal p corrects the bias introduced by the non random selection process, valid adaptive prediction intervals. CQR an adjustment factor of the MLP regression model. After this well as modelindependent, requiring no a 5 training dataset as an additional feature in order to allow the calculation, in the next (second) stage the IMR is added to the neural network to improve the learning process and correcting the distortions caused by selection bias.

5. Uncertainty Quantification of the Model

The uncertainty quantification in model predictions is a key aspect of model fitting in machine learning. Conformal Prediction is a statistical framework for evaluating prediction intervals which are guaranteed to cover the true value with a pre-defined probability. The *Conformalized Quantile Regression* (CQR), is a method within this framework which combines the quantile regression and conformal prediction to compute valid adaptive prediction intervals. CQR is distribution-free as well as modelindependent, requiring no assumptions about the underlying data distribution. Classic regression models are used to predict point-estimates around an average value of the data while quantile regression models are used to estimate different quantiles, providing a range of estimates. The conformal framework adjusts the aforementioned estimates to ensure the specified coverage probability of the prediction intervals of the model. The CQR approach provides that given a dataset of *N* observations $\{(X = x_i, Y = y_i)\}\ (i = 1, 2, \ldots, N)$ with features X ∈ ℝ*^d* and response *Y* ∈ ℝ, CQR constructs a prediction interval

 $C(X) = [L(X), U(X)]$ so that: α that: σ and σ constructs a prediction interval $\mathcal{L}(\mathcal{L})=\mathcal{L}(\mathcal{L})$ so that: constructs a prediction interval $\mathcal{L}(\mathcal{L})=\mathcal{L}(\mathcal{L})$ so that: ${(\mathbf{x}, \mathbf{y})}$ (i.e., \mathbf{y}) with features \mathbf{y} and \mathbf{y} \mathbf{y} \mathbf{z} \mathbf{y} α and α prediction interval α interval α , U(X) so that: $C(X) = [L(X), U(X)]$ so that: the model. The CQR approach provides that given a dataset of N observations ${C(X) = [L(X), U(X)]}$ so that: constructs a prediction interval C(X)=[L(X), U(X)] so that: $P(X)$ so that. constructs a prediction interval C(X)=[L(X), U(X)] so that: $\frac{1}{\sqrt{2}}$

$$
\Pr(Y \in \mathcal{C}(X)) \ge 1 - \alpha,\tag{4}
$$

where $\alpha \in (0, 1)$ is the significance level. Prediction intervals $\hat{\tau}_{1-\alpha/2}$ are estimated by using a *calibration* dat are estimated by training a lower quantity regression model $\frac{n}{2}$ containing n observations which are not used
and an upper quantile regression model $\frac{n}{2}$ separately. model. By defining the score of each observati where $\alpha \in (0, 1)$ is the significance fever. Frequency intervals α $(1-\alpha/2)$ are estimated by using a *calibration* data. belonging to the measure the quantities $\frac{\alpha}{2}$ Subsequent to the training of the models the quantiles $\hat{\tau}_{\alpha/2}$ and this dataset as follows Pr(Y ∈ C(X)) ≥ 1 − α, (4) a lower quantile regression model τα/² and an upper quantile regression model τ1−α/² where $\alpha \in (0, 1)$ is the significance level. Prediction intervals $\hat{\tau}_{1-\alpha/2}$ are estimated by bubsequent to the training of the mode Subsequent to the training of the models the quantiles $\hat{\tau}$ (e.g. and σ) observations of the score of the models the quantiles $\hat{\tau}$ (e.g. and σ) observations of the models the quantiles $\hat{\tau}$ (e.g. and σ) \mathbf{b} $\frac{1}{2}$ Subsequent to the training of the m

ne significance level. Prediction intervals $\hat{\tau}_{1-\alpha/2}$ are estimated by using a *calibration* dat
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$$
s(x, y) = \max[\hat{\tau}_{\alpha/2}(x) - y, y - \hat{\tau}_{1-\alpha/2}(x)]
$$
\n(5)

1) $(1 - \alpha)$ /n is computed in order to estimate the prediction *interval pertaining* each observ t_{max} interval pertaining each observation of Xtest α belongung to the test dataset interval pertaining each observation of Xtest α belongung to the test dataset $t_{\rm b}$ the $t_{\rm c}$ and $t_{\rm c}$ is computed in order to estimate the prediction internal participan each change $\left(1 - \alpha\right)/n$ is computed in order to estimate the prediction *interval pertaining* each step in order to estimate the prediction *interval pertaining* each observed t_{max} the quantitie $q = [(n + 1)(1 - \alpha)]/n$ is computed in order to estimate the helonging to the test dataset the quantile $\hat{q} = [(n + 1) (1 - \alpha)]/n$ is computed in order to estimate the p the quantile $\hat{q} = [(n + 1) (1 - \alpha)]/n$ is computed in order to estimate the prediction *interval pertaining* each observation of $X_{test} = x$ belonging to the test dataset the guantile $\hat{\alpha} = \vec{I}(n + 1)$ (1 = a) $\vec{I}(n)$ is computed in order to estimate the prediction intern the quantile $q = [(n + 1) (1 - \alpha)]/n$ is computed in order to estimate the prediction

$$
\mathcal{C}(x) = [\tilde{\tau}_{\alpha/2}(x) - \hat{q}, \tilde{\tau}_{1-\alpha/2}(x) + \hat{q}] \tag{6}
$$

cases while larger intervals reveal more complicated cases. CQR loss function: where the quantiles $\tilde{\tau}$ are estimeted by using the true value of approach can be adopted to evaluate the prediction The quantities γ are estimeted by using the true value of approximate $V = v$. Smaller intervals correspond to simpler any n where the such the \tilde{z} are estimated by using the true value of the test dataset \tilde{z} where the quantiles $\tilde{\tau}$ are estimeted by using the true value of approach can be adopted the test dataset $Y_{test} = y$. Smaller intervals correspond to simpler any machine learning m where the quantiles T are estimated by using the test dataset \mathcal{M} test dataset \mathcal{M}

are estimeted by using the true value of approach can be adopted to evaluate the prediction intervals of v. Smaller intervals correspond to simpler any machine learning model by incorporating in y. Smaller intervals correspond to simpler any machine learning model by incorporating in it the *pinball* where the quantiles T are estimated by using the test dataset \mathcal{M} test dataset \mathcal{M} *loss* function:

$$
L_{\tau}(y, \hat{y}) = \frac{1}{N} \sum_{i=1}^{N} \max [\tau \cdot (y_i - \hat{y}_i), (\tau - 1) \cdot (y_i - \hat{y}_i)]
$$
(7)

where *y* is the true value and \hat{y} is the predicted one. estimates the estimates the set of the

6. Proposed Approach: A Case Study combination

The proposed evaluation procedure for quantifying the study in order to evaluate its efficiency. Input data is from 6 6 uncertainty in a deep learning model for imputation purposes is presented in this Section and is applied to a real-world case the ISTAT\ statistical register by the name of *ARCHIMEDE* regarding the resident population in Italy. The information gathered into the register is stored in socio-economic variables such as demographic variables, working activity, attained level of education, and income. Each record indicates an individual. People from the same household aregrouped by the same household identification number. The model being evaluated

estimates the relationships between equivalent household where y is the predicted one.
income and relevant features of the household breadwinner in to a specific region of the Italian territory only. combination with specific household group variables pertaining

6 6 **6.1. Input Data of the Model**

An initial set *X* of 253286 households was selected by picking households with $n \geq 3$ members only. As it is reported in Sec. 3, these households are mathematically described as being fully connected undirected weighted graphs. The number of vertices is variable, so that there are graphs of different dimensions in the dataset. Every vertex is related to a sequence of *K* categorical variables (nodal attributes) $a^{(i)} = \{a^{(i)}_1, a^{(i)}_2, \dots, a^{(i)}_k\}$ called the *profile* of the vertex v_i ($i = 1, 2, ..., n$). The list of attributes in this case study is reported in Tab. 1: Subsequent to the application of

Table 1: Attributes of the Household Members in the Input Dataset

the algorithm reported in Sec. 3, a dataset of $N = 254227$ pairs of *equal* households was selected from the X dataset. Selected pairs comprise 193803 distinct households of different sizes as well as different values of the attributes of each member. Pairwise similar households share the same size as well as the same attributes of their members so that the similarity value of every pair is equal to 1. The resulting input dataset is split into two partitions: dataset *A* contains the first household of every pair, while dataset *B* contains the second household of the same. These datasets are used to constitute two independent model

training processes based on different but similarly distributed data, i.e., $A \rightarrow B$ and $A \leftarrow B$.

6.2. Correction of the Selection Bias

The dataset $(A \cup B) \subset X$ is not selected at random. As a consequence, the deep learning model may be affected by bias. This selection bias is treated by using the 2-stage Heckman procedure as reported in Sec. 4. The list of the variables used in the probit model of the selection stage *sel* ∼ probit(*predictors*) is reported in Tab. 2: The dependent variable *sel* equals 1 if the

observation belongs to the subset *A* ∪ *B*

Table 2: Predictors of the Probit Model

and 0 otherwise. Subsequent to this model fitting, the IMR is added as a feature in the predictive model of the multilayer perceptron to adjust it for bias.

6.3. Coverage and Adaptivity of the Prediction Intervals

The deep learning model in this case study is composed by a DAE which is stacked with a MLP in order to estimate the equivalent household income. The DAE transforms the input vector of the categorical variables into a smaller vector made up of numerical variables, the input of the MLP. This encoded input vector is augmented by adding the IMR variable in order to correct the model for the selection bias. The predictors of the probit model in Tab.2 are necessary for estimeting the probability of households of being selected in the training dataset. Variables used as predictors of the deep learning model (DAE and MLP)

are listed in Tab.3. The categorical variables are transformed into dummy variables before being encoded by the DAE. Subsequent to this data pre-processing, the MLP is trained by using crossvalidation. The numerical variables reported in Tab.3 are also considered as being categorical as a result of a top-coding of the aforementioned1. The uncertainty quantification of this model is carried out by integrating the loss function described in Eq.7 in the MLP in order to evaluate the prediction intervals as reported in Sec.5 in accordance with Angelopoulos and Bates [11]. The proposed approach for quantifying the epistemic component of the uncertainty provides two analoous procedures: 1) $A \rightarrow B$ and 2) $A \leftarrow B$ as is described in the diagram in Fig.3. The results of the application of the $A \rightarrow B$ and $A \leftarrow B$ are reported in Tab.4 and Tab.5, where coverage and adaptivity are respectively compared in order to investigate uncertainty. The empirical coverage

Table 3: Predictors of the Imputation model

measures the percentage of cases in which the prediction intervals contain the true value of the dependent variable. This property is also evaluated asymptotically by using an efficient caching of the non-conformity scores calculated as prescribed in

the literature. Adaptivity measures the property of the prediction intervals to adapt to the cases covered by the model. Small intervals reveal good prediction ability of the model, while larger intervals suggest greater uncertainty in the predictions. In order

Figure 3: Workflow diagram of the $A \rightarrow B$ procedure

Model	Empirical	Score caching
$\sharp 1(A \rightarrow B)$	0.9545153	0.951681
$\mid \#2(A \leftarrow B) \mid$	0.9503557	0.9511803

Table 4: Coverage of the Prediction Intervals

Stats	#1 $(A \rightarrow B)$	#2 $(A \leftarrow B)$
Min	0.000001	0.0006018
1st Qu.	0.094100	0.1309214
Median	0.110540	0.1530824
Mean	0.142123	0.1946715
3rd Qu.	0.130294	0.1782837
Max	1.000000	1.0000000

Table 5: Adaptivity of the Prediction Intervals

to compare the evaluation procedures, the adaptivity was normalized by calculating as the ratio between the length of the prediction interval and the maximum value of all interval lengths as is reported in Tab. 5. In order to further investigate

luation procedures, the adaptivity was the uncertainty of the model, the boxplots of the distributions ating as the ratio between the length of pertaining to the length of prediction intervals by the input l and the maximum value of all interval categorical variables are reported in the following.

Figure 4: Prediction Interval Lengths: *BW. Income Class*

1 2 3 4 5 1 2 3 4 5 BW.MaritalStatus Figure 8: Prediction Interval Lengths: *BW.MaritalStatus*

7. Conclusions

This study proposes a general approach for evaluating the Multilayer Perceptron stacked with the Denoising Autoencoder the deep learning regression model being uncertainty of deep learning models as is the case of the

described in this paper. This evaluation approach can be extended to other machine learning models if they cater for the integration of robust techniques for quantifying uncertainty. The structure of the deep learning regression model being

 $\mathcal{L}_{\mathcal{A}}$ Figure 9: Prediction interval lengths: BW.EduLevel

Fig. 10 Prediction interval lengths: BW.Gender Figure 10: Prediction Interval Lengths: BW.Gender

BW.PensScheme Figure 11: Prediction Interval Lengths: *BW.PensScheme*

considered combines a data pre-processing algorithm and a to ana procedures, i.e. *A* to-*B* and *B*-to-*A* which are carried out in order learning model produces valid prediction regression model. The proposed evaluation approach requires the selection of two similarly distributed datasets *A* ans *B*, focusing on the comparison of complementary evaluation

similarly distributed datasets A ans B , the underlying unknown data distribution. The integration of mparison of complementary evaluation the Conformalized Quantile Regression approach in the deep B and B -to- A which are carried out in order learning model produces valid prediction to analyze coverage and adaptivity of the model, measures for assessing the reliability of prediction intervals in rappresenting learning model produces valid prediction

Fig. 12 Prediction interval lengths: AllSameCtzn **Figure 12:** Prediction interval lengths: *AllSameCtzn*

Interval length

Fig. 13 Prediction interval lengths: BW.AgeClass Fig. 13 Prediction interval lengths: BW.AgeClass **Figure 13:** Prediction Interval Lengths: *BW.AgeClass*

Figure 14: Prediction Interval Lengths: *NRetired*

of the model. The two-sided evaluation approach proposed in this observations which involve the afore study provides consistent insights as a result of the enhancement increases as well. The reduction of ale of the epistemic uncertainty coompared to the aleatoric one. This using the DAE as a data pre-processing n study does not explicitly separate the aleatoric and the epistemic it for certain subgroups, resulting in higher components of the uncertainty. The aleatoric component is This is particularly evident in households instead reduced by training the DAE on the selected datasets and different breadwinner characteristics in the evaluation process of the allocation and while the intervals to primarily reflect epistemic component while the in the evaluation process of the uncertainty allow the prediction intervals which provide an evaluation criterion of the reliability *A* and *B*. The subsequent data encoding in both the evaluation procedures equalize the aleatoric uncertainty as a result of the data similarity. The reduction of the aleatoric component contribution equalization implies that differences between the evaluations are related to epistemic uncertainty. It is important to underline that, due to its probabilistic setup, the LSH algorithm assigns the pairs of similar households to *A* and *B* in a random order, minimizong potential biases during dataset creation. The detction algorithm resembles a split-plot design, where the primary variables act as main plot factors, and breadwinner specific variables are treated as subplot factors nested within similar households. In order to further address the bias introduced by the non random selection of training data, the Heckman correction is incorporated into the regression model. This is achieved by including the Inverse Mills Ratio as a feature to account for selection bias. The Heckman correction mitigates the concept drift resulting from a non rappresentative training sample, ensuring that the predictions generalize better to the target population. However, it assumes that the selection process is fully explained by observable variables and relies on the normality of errors, which may not hold universally. Including the IMR as a covariate also adds complexity to the interpretation of the model's coefficients. An in-depth investigation of the deep learning model under evaluation reports a higher occurrence of outliers on the lefthand side in the boxplots, corresponding to low values of the categorical variables under consideration. The probability

of there being outliers is likely to increase as the number of observations which involve the aforementioned categories increases as well. The reduction of aleatoric uncertainty by using the DAE as a data pre-processing may not fully eliminate it for certain subgroups, resulting in higher frequency of outliers. This is particularly evident in households with similar structures and different breadwinner characteristics, where variations in prediction intervals reveal epistemic uncertainty pertaining to complex underrepresented groups. Differences in the lengths of prediction intervals between the categories indicate residual epistemic uncertainty steming from uneven data rappresentation. Categories pertaining to longer intervals reflect greater uncertainty, which arises from underrepresentationor complex data patterns. Outliers in these intervals suggest that the DAE does not correct input data anomalies or does not handle rare attributes, requiring a more complex pre-processing algorithm as is the case of architectures with a higher number of hidden layers or a higher number of DAEs arranged in a stack. The results of this study demonstrate the effectiveness of the comparison betwenn two opposite evaluation procedures based on noisefree similar data for smoothing the aleatoric component out from the uncertainty quantidication process while emphasize the epistemic component being reflected in the prediction intervals of the model. A further reduction of the residual uncertainty may be achieved by rcurring to data balancing in order to reduce the occurrence of rare observations or ensemble methods for combining both the evaluation proceduresin order to improvw the robustness of the prediction intervals [13-16].

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