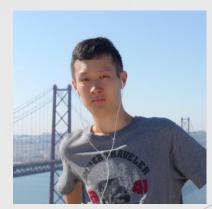


O1Who are we?





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Major in Bioinformatics, at UCPH



01 VERIFICATION

IDEA

- Using synthetic data, train a model to predict readmission.
- Check the performance of this model on synthetic data and on real data.



01 VERIFICATION

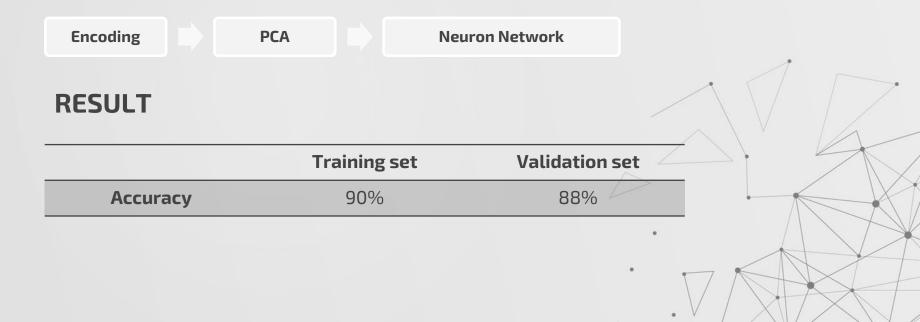
RESULT

Synthetic data (Train set)	Synthetic data (Val set)	Real data
71%	68%	53%

- There is a **significant difference** between 2 datasets.
- Real data is harder to make prediction. Using this set of synthetic data to do data analysis is risky.

02 CLASSIFICATION

MODEL



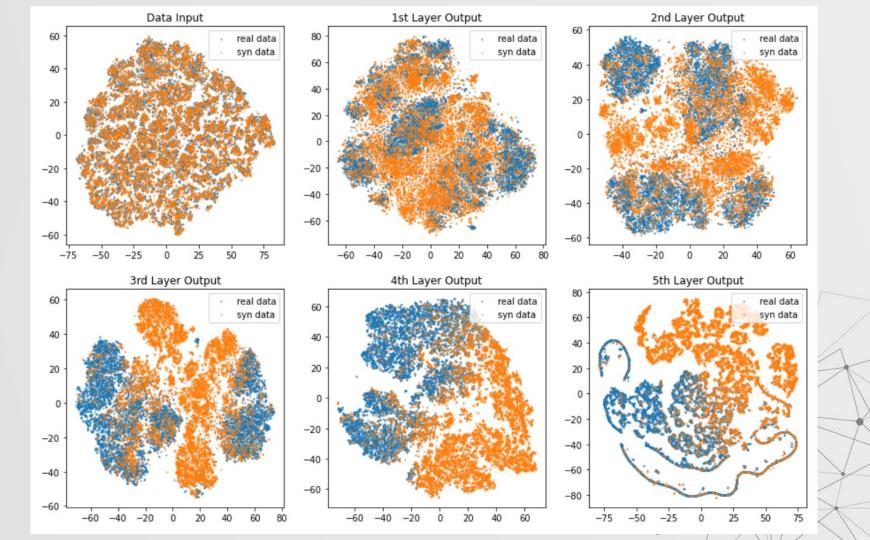
03 ANALYSIS

- Low dimensional embedding
- Check hidden-layer activation
- Re-train the model using combination of dimensions
- Analyze crucial dimensions

3.1 Low dimension embedding

- Embedding and visualization by t-SNE
- Check the decoupling of each layer in neuron network





3.2 Check hidden layer activation

- Try to find which features the first layer detects

Find the samples that mostly activate the first layer

Recover the input by taking pseudo-inverse

Compare the difference

$$A = relu(W \cdot X + b)$$
$$X \approx W^{-1} \cdot (A - b)$$

The inspiration comes from Zeiler M.D., Fergus R. (2014) Visualizing and Understanding Convolutional Networks.

3.2 Check hidden layer activation

- By comparing the result, we teased out dimensions not used in decoupling.
- Remain 22 dimensions.



3.3 Train model with combination of dimensions

- To check the most important dimensions among the 22 dimension.
- Find 3 most important dimensions:

"insulin"
"change"
"diabetesMed"



3.3 Train model with combination of dimensions

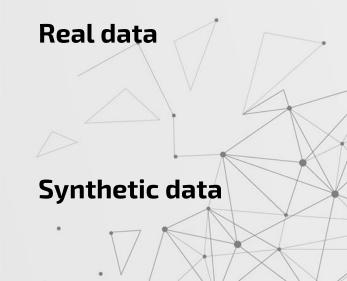
	All features	Only 3 features	All other features
Accuracy	88%	72%	74%
		7 14 11	

3.4 Understand the difference

- Compare the correlation coefficient of this 3 dimensions.

Corrs	insulin	change	diabetesMed
insulin	1.0	-0.14	0.26
change	-0.14	1.0	-0.51
diabetesMed	0.26	-0.51	1.0

Corrs	insulin	change	diabetesMed
insulin	1.0	-0.02	0.01
change	-0.02	1.0	0.02
diabetesMed	0.01	0.02	1.0



Conclusion

- The mainly difference between real data and synthetic data are dimensions "insulin", "change" and "diabetesMed";
- In real data, these three dimensions are highly related;
- while in synthetic data, these three dimensions seems to be generated **independently**.

Future Plan

 Re-sampling these 3 dimensions to generate a better synthetic data

 Use this new set of synthetic data to train a model predicting re-admission, then check if this model works well on real data

