

Assignment 4 - DL

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Preprocess step on both datasets:

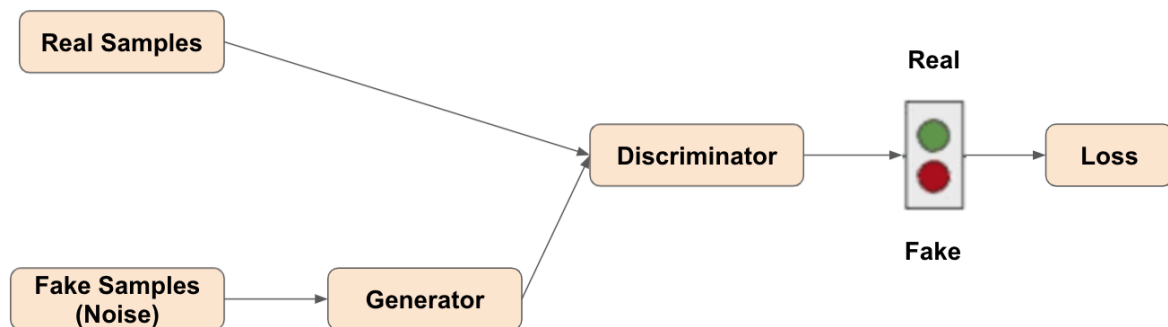
Categorical features - we encode categorical features as to integer values. We encode the features to numerical values and are not using one hot encoding.

Continuous features - we kept the same.

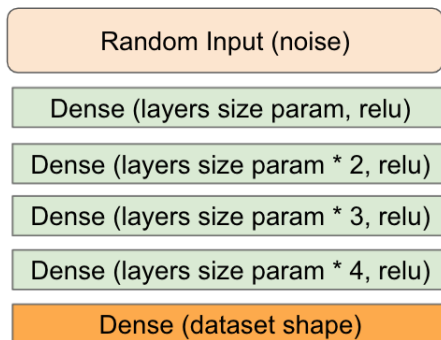
We used PowerTransformer scaler (from Sklearn) on all the features to get a Gaussian distribution for the data.

Part 1 – Generative Adversarial Networks

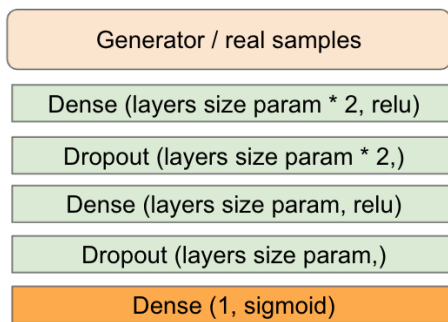
The architecture of the GAN:



The architecture of the Generator:



The architecture of the Discriminator:



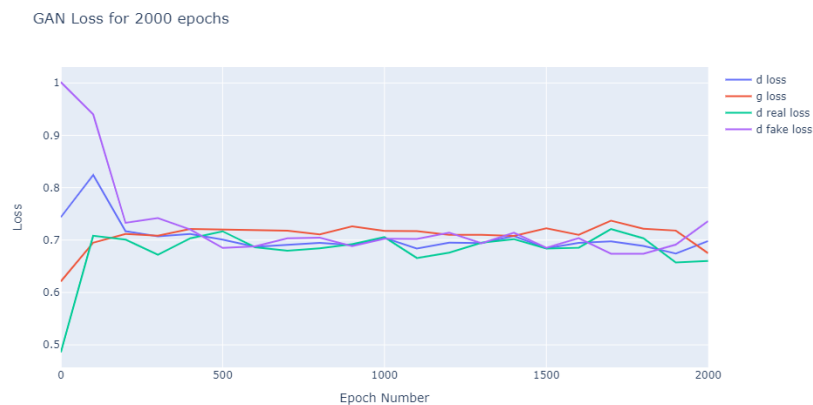
Results on Datasets:

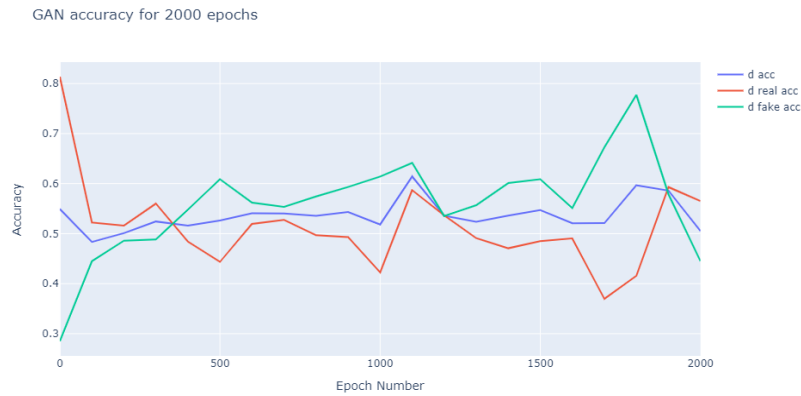
1. Diabetes Dataset:

Hyper parameters:

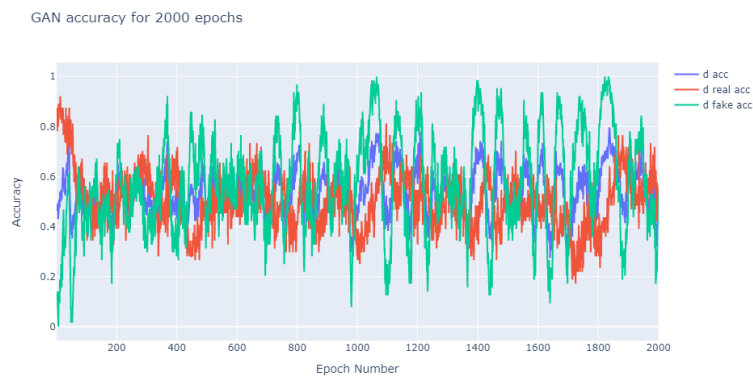
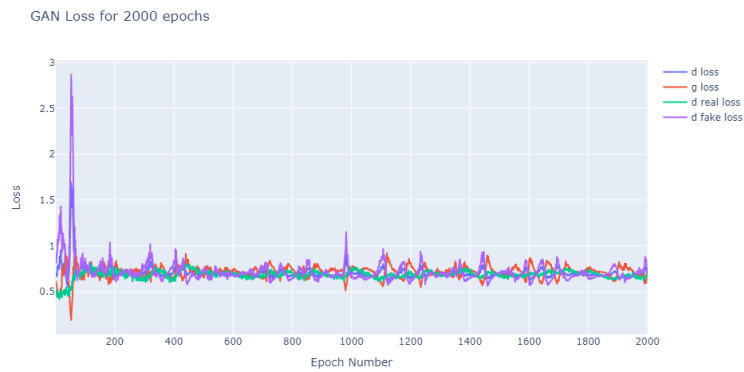
Parameter description	Parameter value
batch size	64
Number of epochs	2000
Learning rate	5e-4
Noise dimension size	32
Number of units in layers factor	30

Average statistics per 100 epochs:





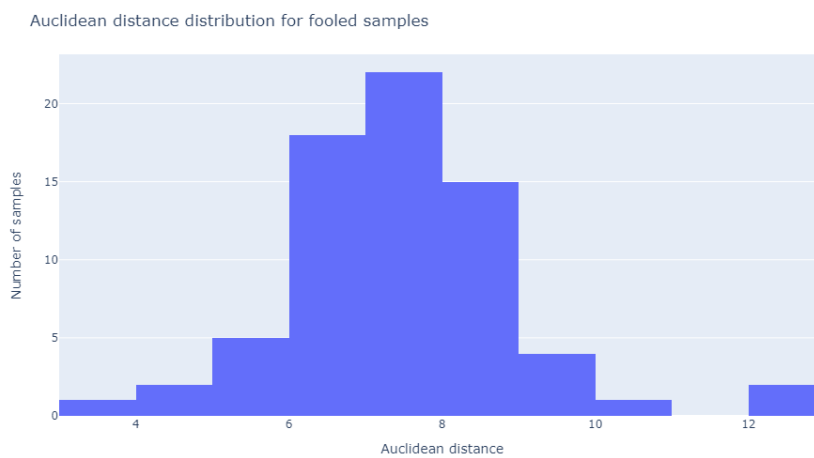
Raw statistics:



The GAN converged at epoch 2000.

Samples that fooled the Discriminator:

Out of 100 generated samples, 70 fooled the Discriminator, while 30 did not.



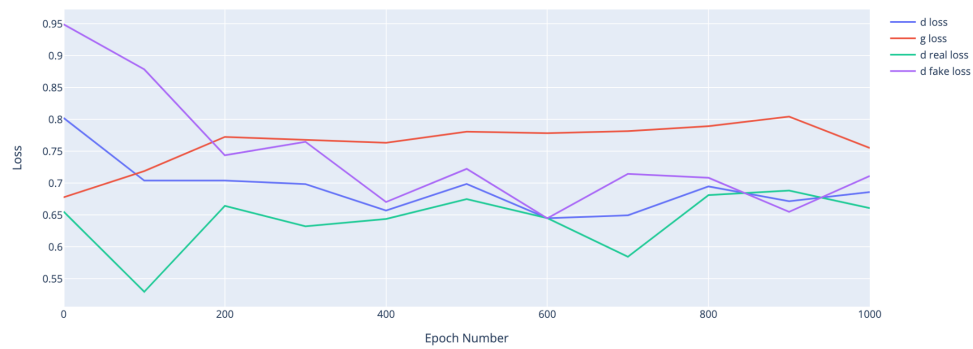
2. German credit Dataset:

Hyper parameters:

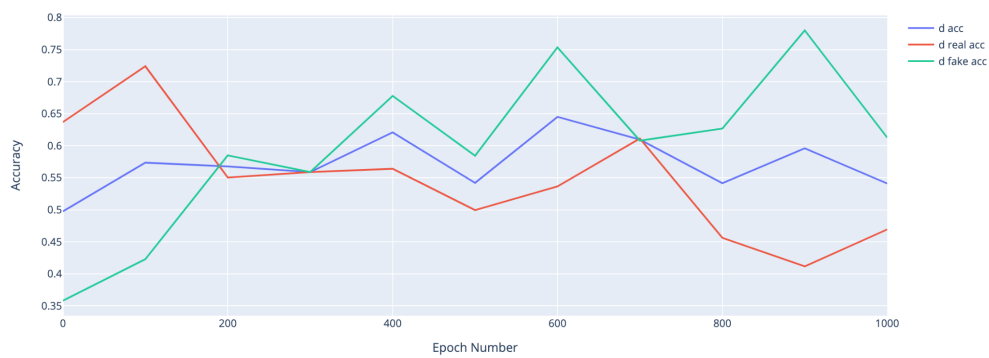
Parameter description	Parameter value
batch size	64
Number of epochs	1000
Learning rate	1e-3
Noise dimension size	64
Number of units in layers factor	25

Average per 100 epoch

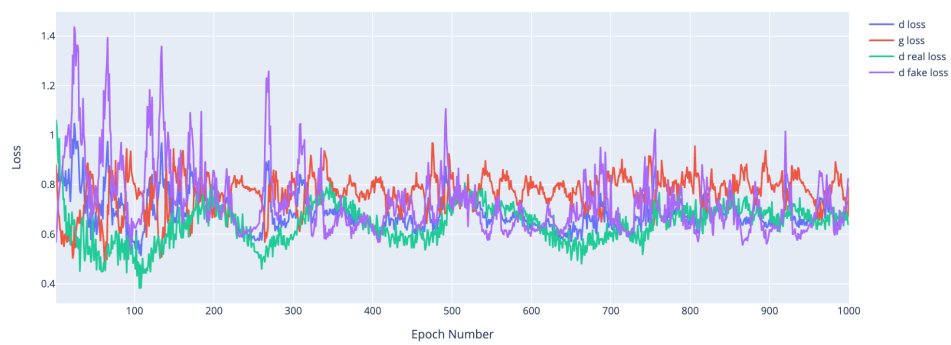
GAN Loss for 1000 epochs



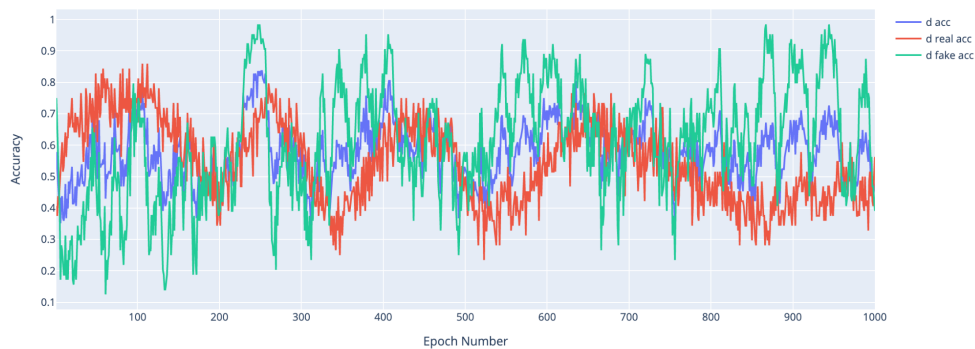
GAN accuracy for 1000 epochs



Raw statistics:



GAN accuracy for 1000 epochs

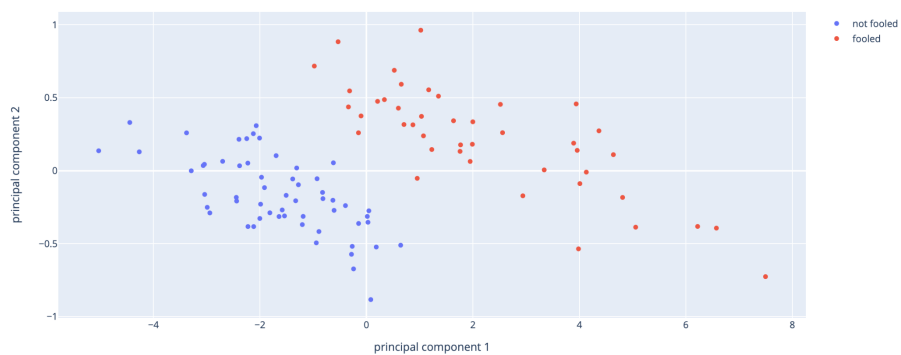


The GAN converged at epoch 1000.

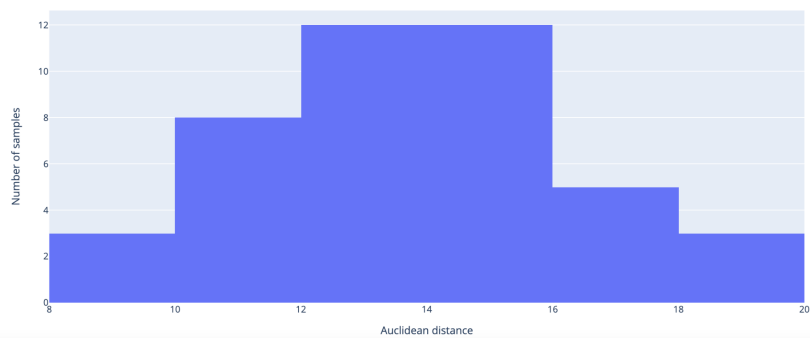
Samples that fooled the Discriminator:

Out of 100 generated samples, 43 fooled the Discriminator, while 57 did not.

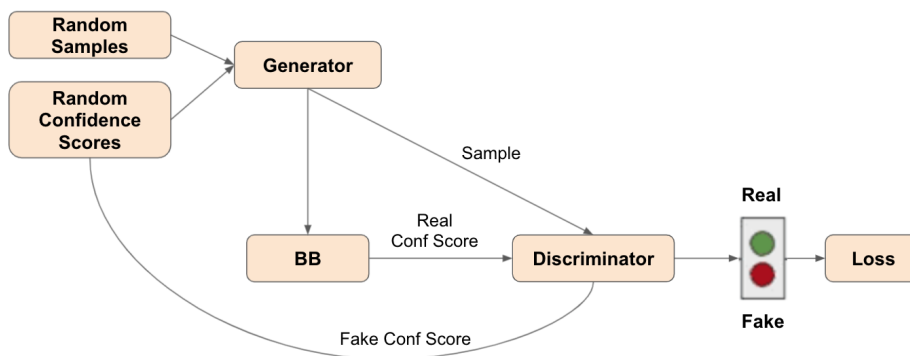
Fooled samples visualization with PCA



Auclidean distance distribution for fooled samples

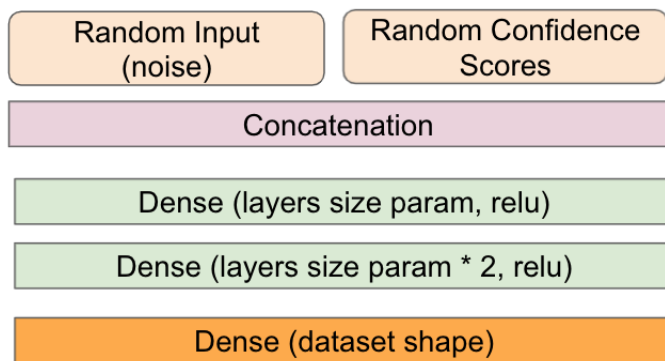


Part 2 – Generative model for sample generation

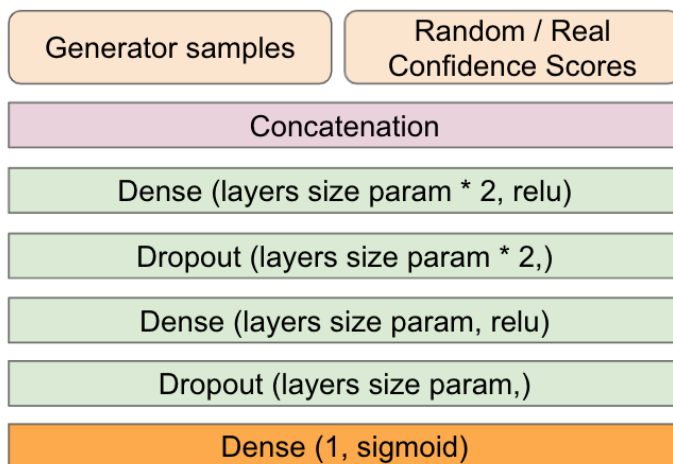


The architecture of the GAN:

The architecture of the Generator:



The architecture of the Discriminator:



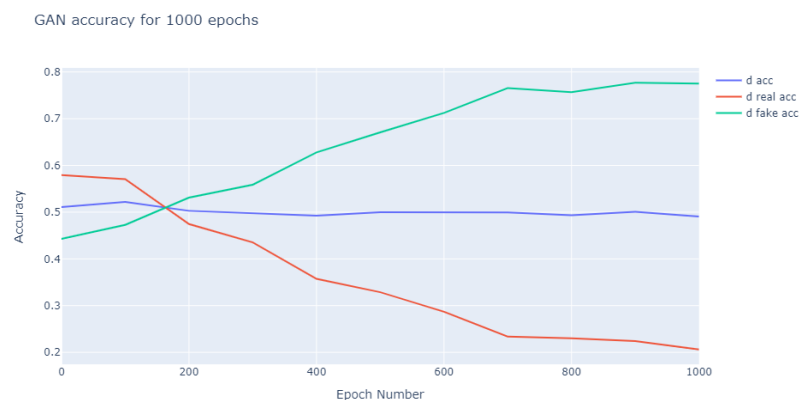
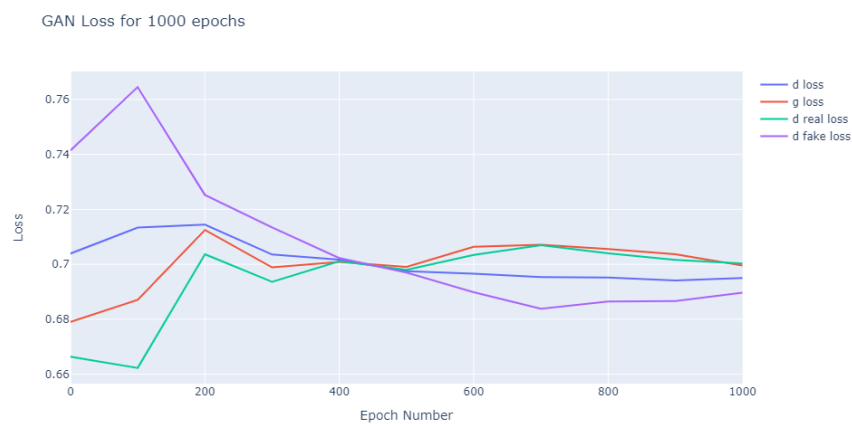
Results on Datasets - part 2:

3. Diabetes Dataset:

Hyper parameters:

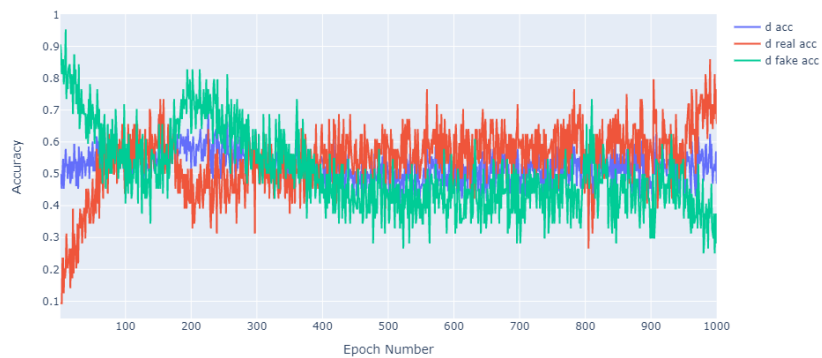
Parameter description	Parameter value
batch size	64
Number of epochs	1000
Learning rate	5e-4
Noise dimension size	8
Number of units in layers factor	16

Average statistics per 100 epochs:

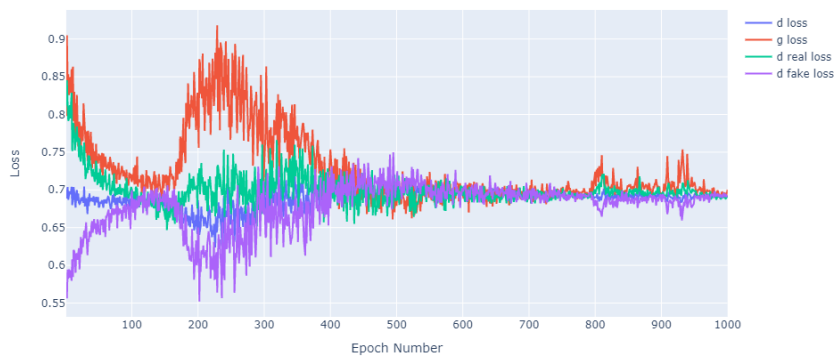


Raw statistics:

GAN accuracy for 2000 epochs



GAN Loss for 2000 epochs

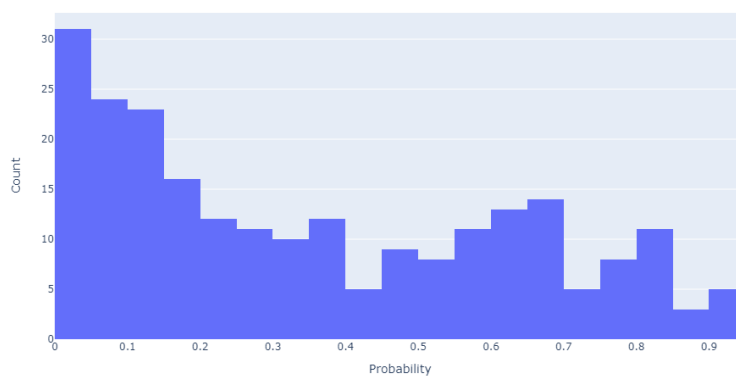


The GAN converged at epoch 1000.

The RandomForest classifier's performance:

- The classifier accuracy on test set (30%): 0.78
Confidence score (Probability) distribution:
Min: 0.004, Max: 0.942, Mean: 0.354

Random Forest probability distribution for Diabetes



- For generated samples (1000 samples):
Confidence score (Probability) distribution:

Fake- Min: 0, Max: 0.998, Mean: 0.487
RF- Min: 0.318, Max: 0.726, Mean: 0.568

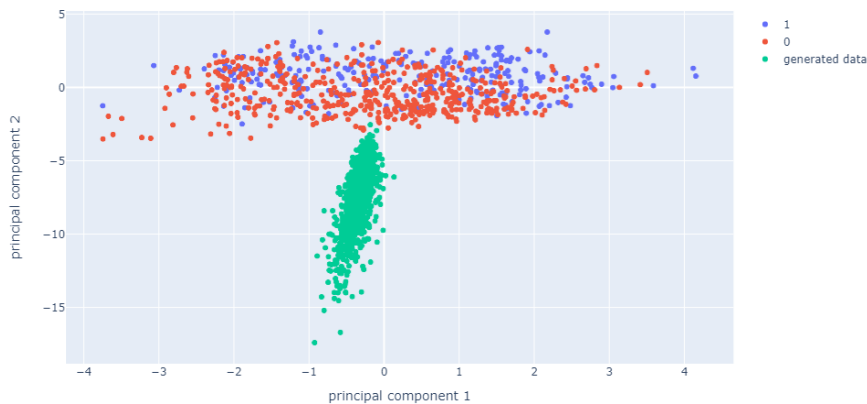
Confidence scores for generated samples on Diabetes



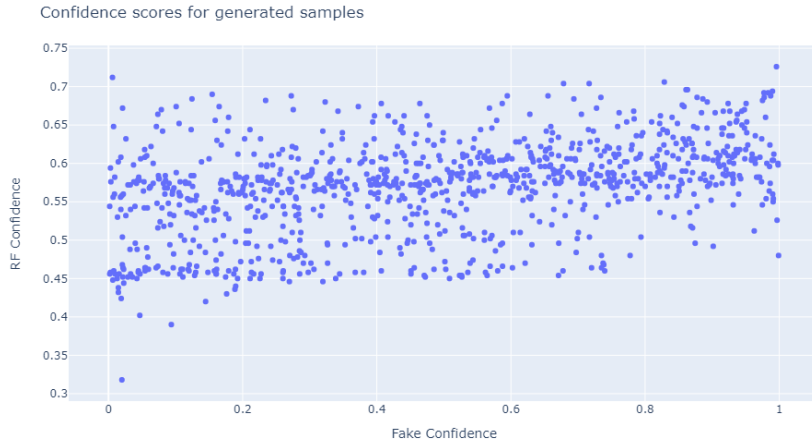
It seems that the generator is generating samples that succeed to confuse the RF classifier, most of the samples receive ~0.5 confidence score.

We visualize both the generated samples and the origin data (using PCA dimension reduction), we can see that there is similarity between the generated data and the origin data (for both classes).

Origin and generated samples - visualization with PCA



Correlation between RF confidence and Fake confidence: 0.421

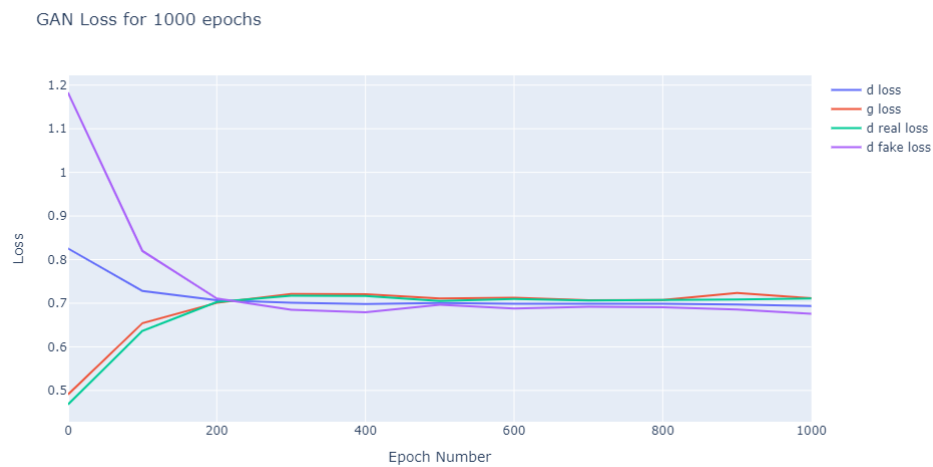


4. German credit Dataset:

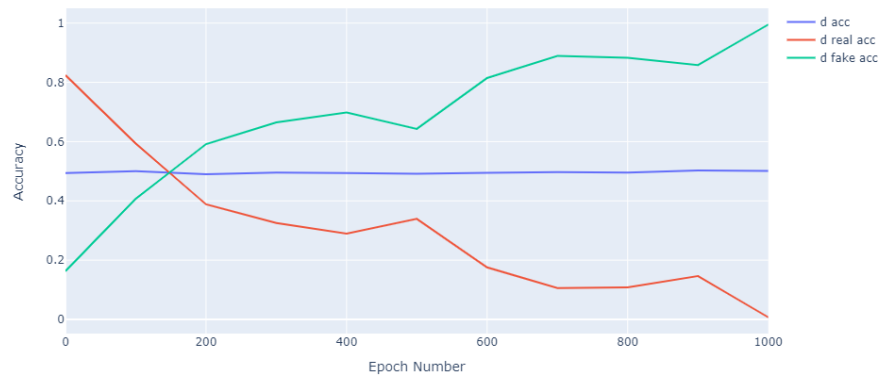
Hyper parameters:

Parameter description	Parameter value
batch size	64
Number of epochs	1000
Learning rate	1e-3
Noise dimension size	48
Number of units in layers factor	20

Average statistics per 100 epochs:

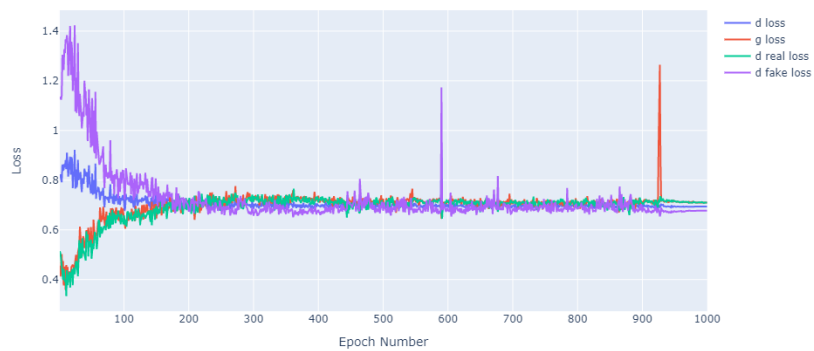


GAN accuracy for 1000 epochs

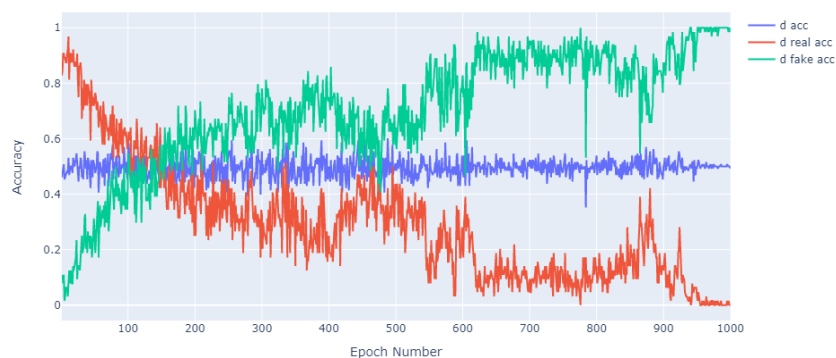


Raw statistics:

GAN Loss for 2000 epochs



GAN accuracy for 2000 epochs

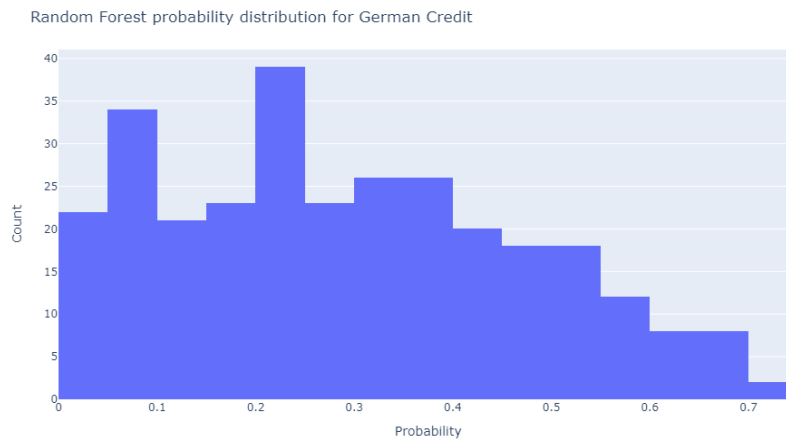


The GAN converged at epoch 1000.

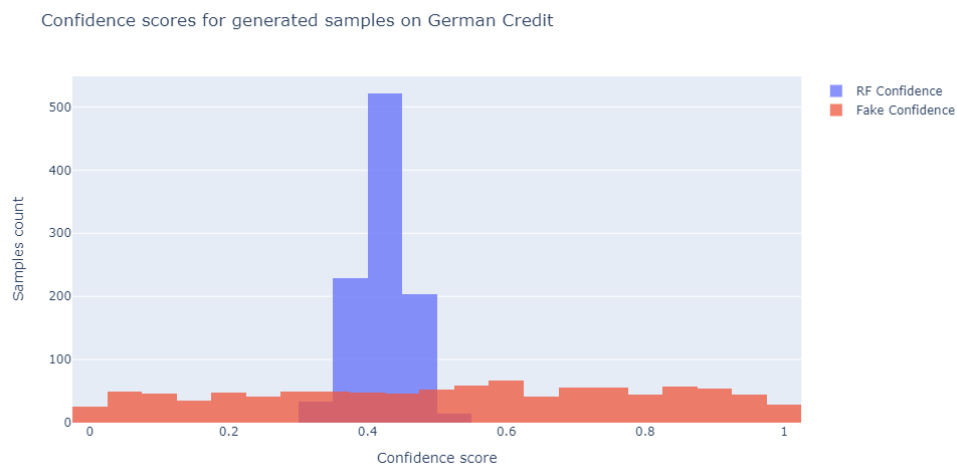
The RandomForest classifier's performance:

- c. The classifier accuracy on test set (30%): 0.73
Confidence score (Probability) distribution:

Min: 0.014, Max: 0.742, Mean: 0.294



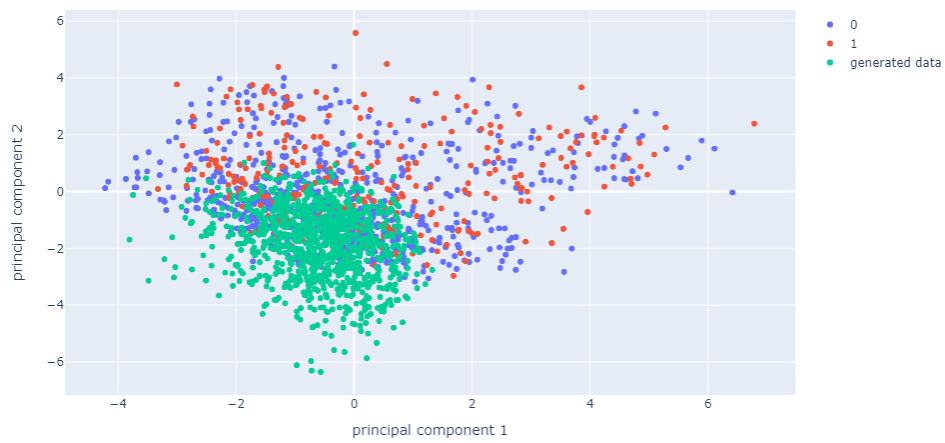
- d. For generated samples (1000 samples):
Confidence score (Probability) distribution:
Fake- Min: 0, Max: 0.997, Mean: 0.516
RF- Min: 0.31, Max: 0.518, Mean: 0.418



It seems that the generator is generating samples that are more biased toward the 0 class, the RF confidence score centers around ~0.4 confidence score.

We visualize both the generated samples and the origin data (using PCA dimension reduction), we can see that there is more similarity between the generated data and the 0 class in the origin data. We can see that the generator suffers from a Mode collapse issue, though it still succeeds in generating samples that are similar to class 1 as well.

Origin and generated samples - visualization with PCA



Correlation between RF confidence and Fake confidence: -0.041

Confidence scores for generated samples

