# Copulas as High-Dimensional Generative Models: Vine Copula Autoencoders

Natasa Tagasovska, Damien Ackerer and Thibault Vatter. NeurIPS, 2019.

#### Brief introduction of VCAE

#### Vine-copula-Autoencoder

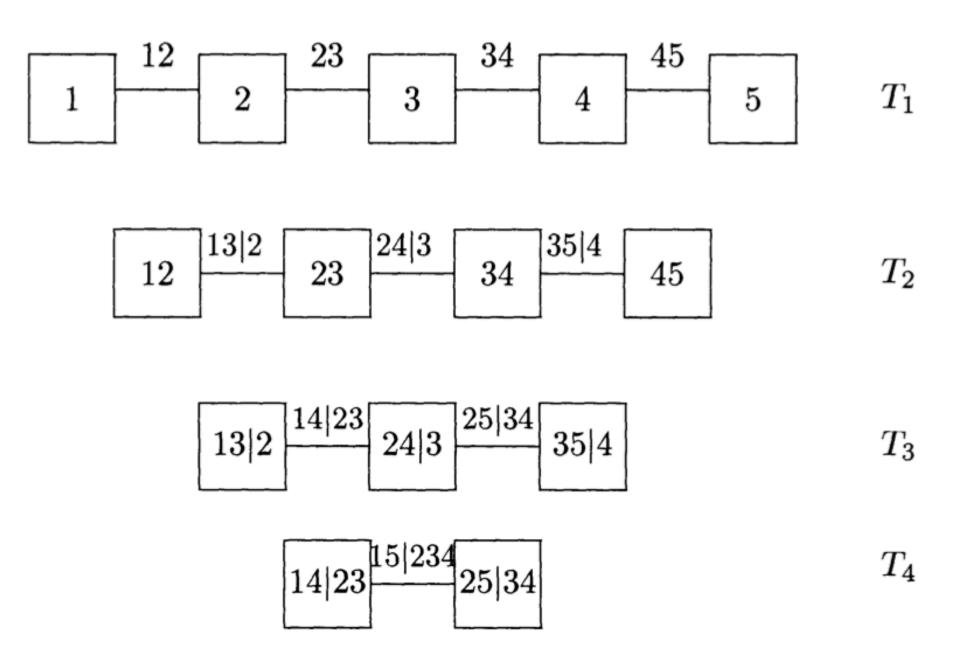


Figure 1. A five-dimensional D-vine, with  $T_j$  denoting the *j*th tree for j = 1, 2, 3, 4.

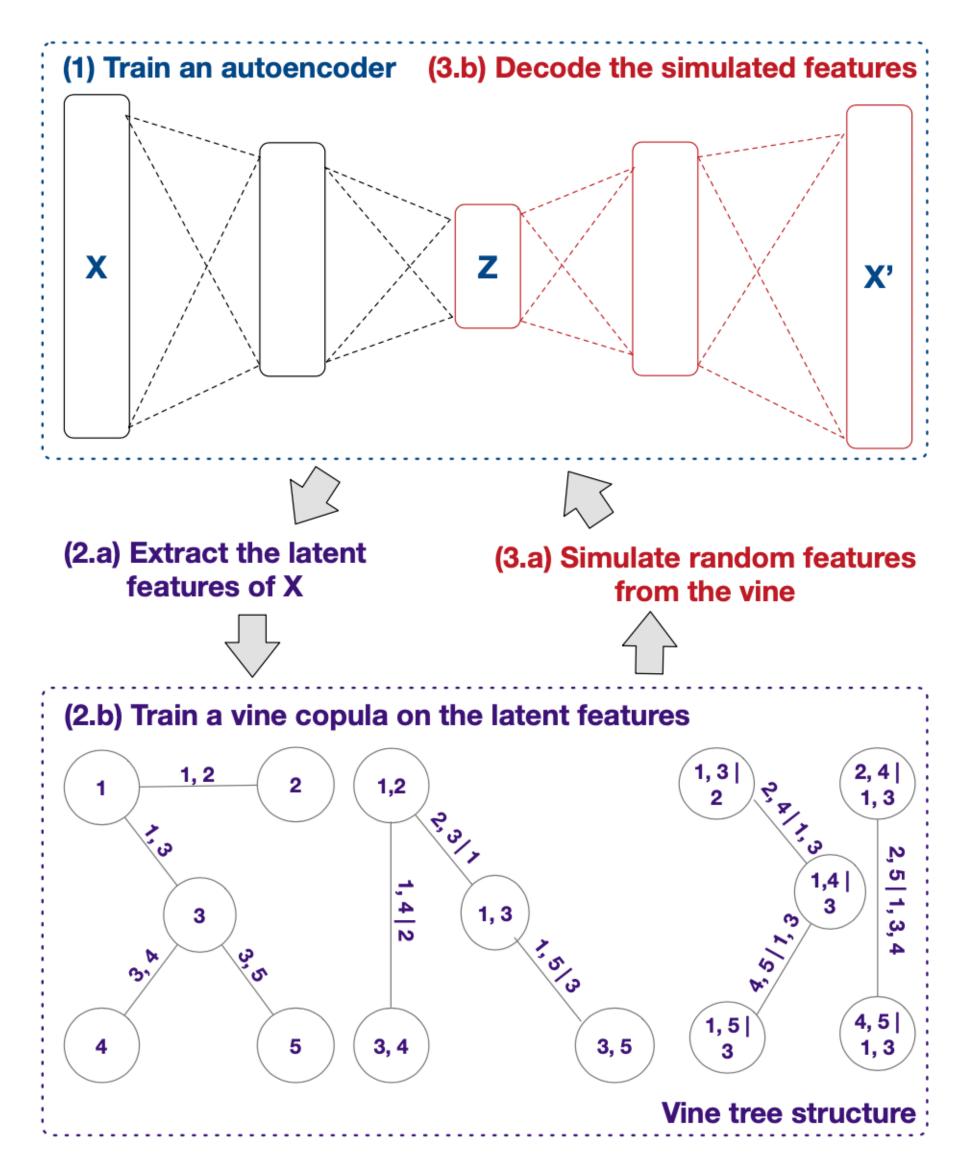


Figure 1: Conceptual illustration of a VCAE.

### VCAE algorithm

#### Algorithm 1 Vine Copula Autoencoder

```
Input: train set X of \{x_1, x_2, ...x_n\} images.
```

1. Train autoencoder component with X:

$$\begin{array}{l} f \leftarrow encoder \\ g \leftarrow decoder \end{array}$$

2. Encode train set with f:

$$\phi(X) \leftarrow f(X)$$

3. Fit a vine copula c using encoded features:

$$c \leftarrow \{\phi_1, \phi_2, ... \phi_n\}$$
 (as described in Sec 2.2 and 2.3).

4. Sample random observations form c:

$$\phi' \leftarrow c(\phi)$$
 (as in Sec 2.4)

5. Decode the random features:

$$X' \leftarrow g(\phi')$$

Output: generated images X'.

- n=5: truncated at 5th tree
- Using nonparametric copula
- Using the 'Rvinecopulib' package in R
- Combining R package with Python code by 'rpy2' package in python

#### CNN for Decoder and Encoder

• Encoder:

$$x \in R^{32 \times 32} \to Conv_{32} \to BN \to ReLU$$
  
  $\to Conv_{64} \to BN \to ReLU$   
  $\to Conv_{128} \to BN \to ReLU$   
  $\to FC_{10}$ 

• Decoder:

$$z \in R^{10} \to FC_{100} \to ConvT_{128} \to BN \to ReLU$$
$$\to ConvT_{64} \to BN \to ReLU$$
$$\to ConvT_{128} \to BN \to ReLU$$
$$\to FC_{1}$$

#### Experiments in paper

MNIST, Street View House Numbers, and one large scale - CelebA.

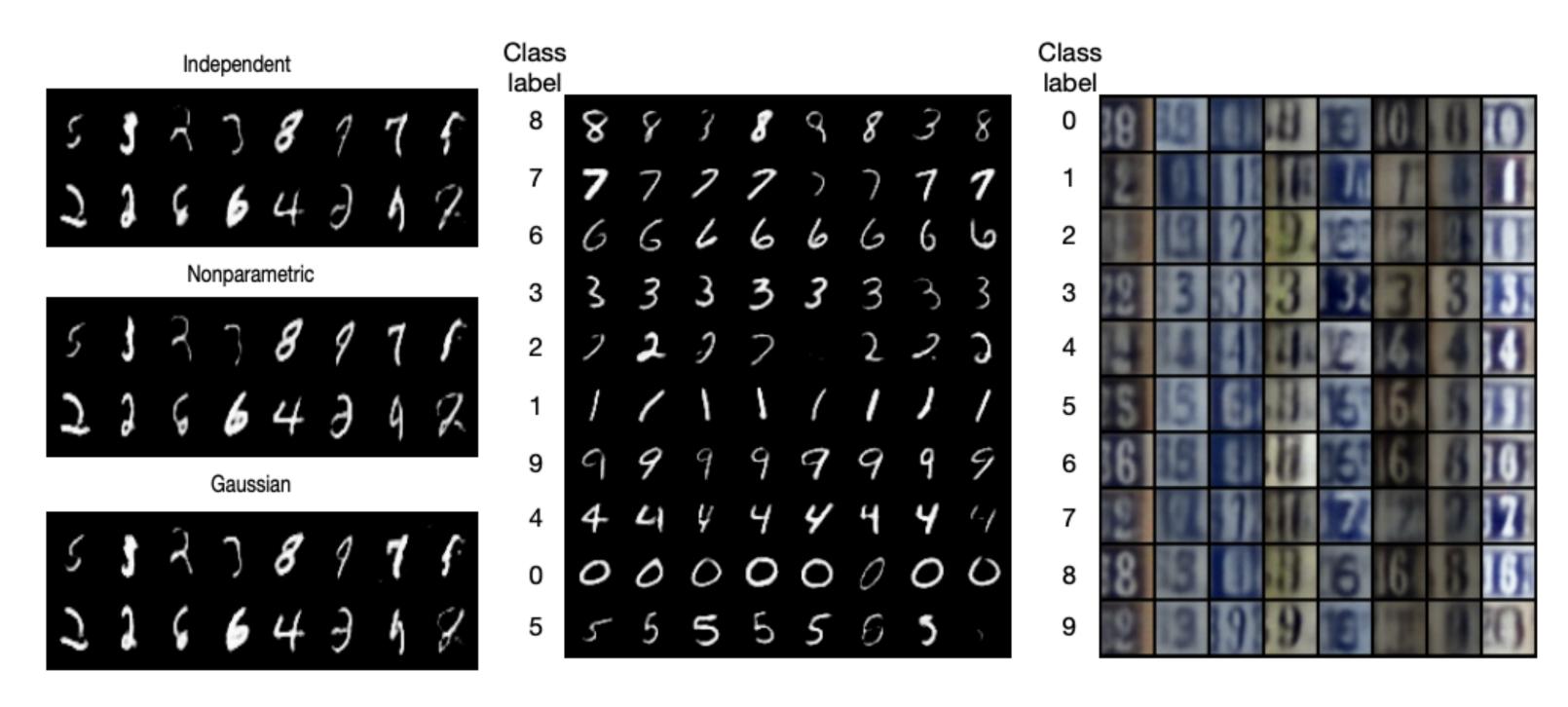


Figure 4: Left - impact of copula family selection in image sampling. Right - Random samples of Conditional *VCAE* on **MNIST** and **SVHN** 

#### VCAE as generative models

Comparing with DCGAN(deep convolutional GAN) and variational autoencoders (VAEs)

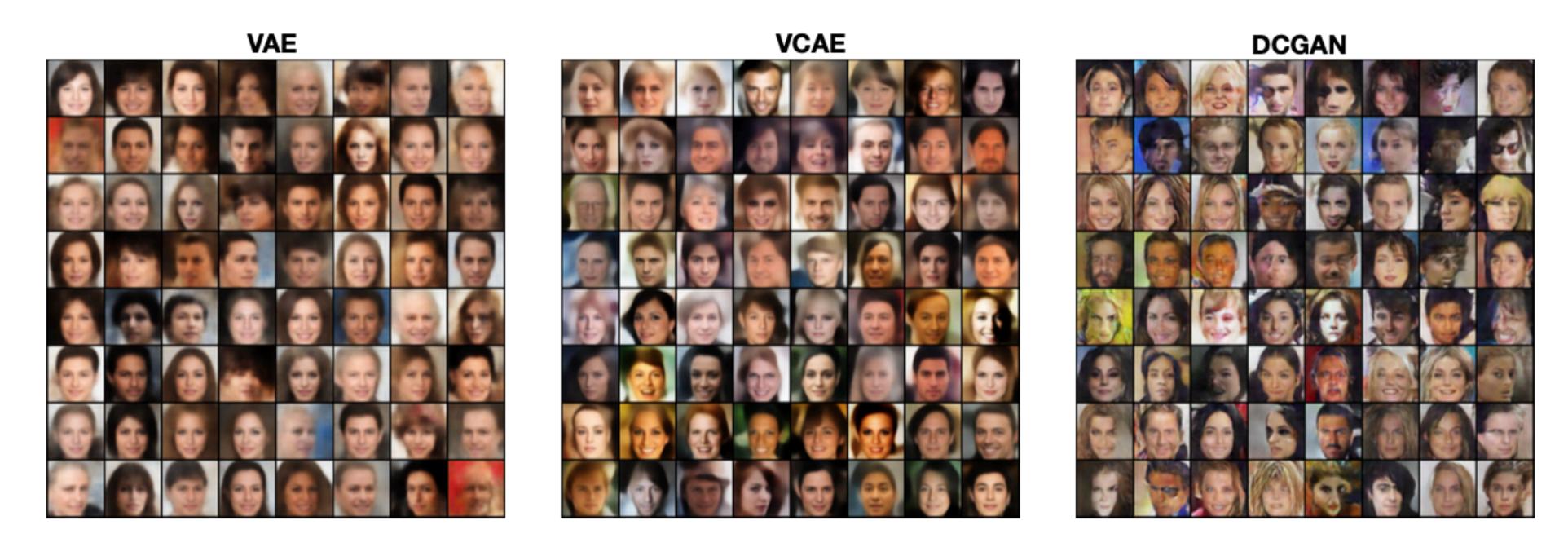


Figure 7: Random samples from left - VAE, - VCAE on CelebA, both trained for 200 epochs. - DCGAN best results at 30 epochs.

## Our Aim: perform VCAE on MedMNIST data

#### organmnist\_axial

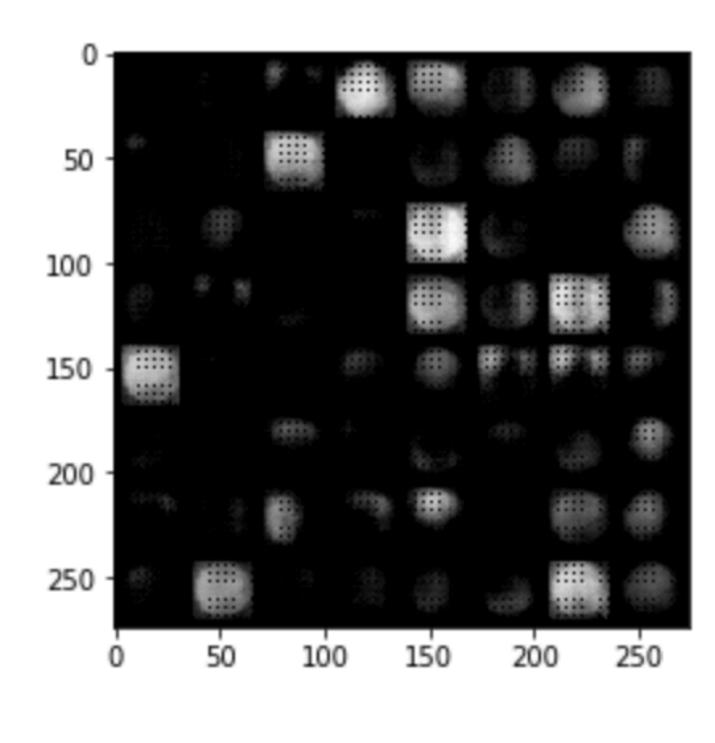
- Epoch=4
- Loss figure

```
import matplotlib.pyplot as plt
In [14]:
          plt.plot(lossseq)
Out[14]: [<matplotlib.lines.Line2D at 0x7fece1ce1350>]
           800
           780
           760
           740
           720
           700
           680
```

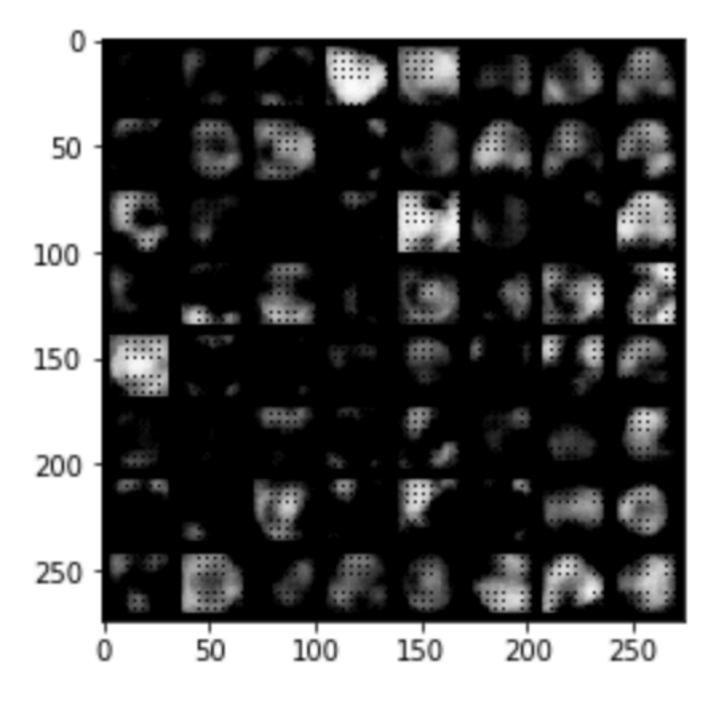
#### Results

<matplotlib.image.AxesImage at 0x7fec7ba06c10>

<matplotlib.image.AxesImage at 0x7fec7b07ed50>



epoch=1

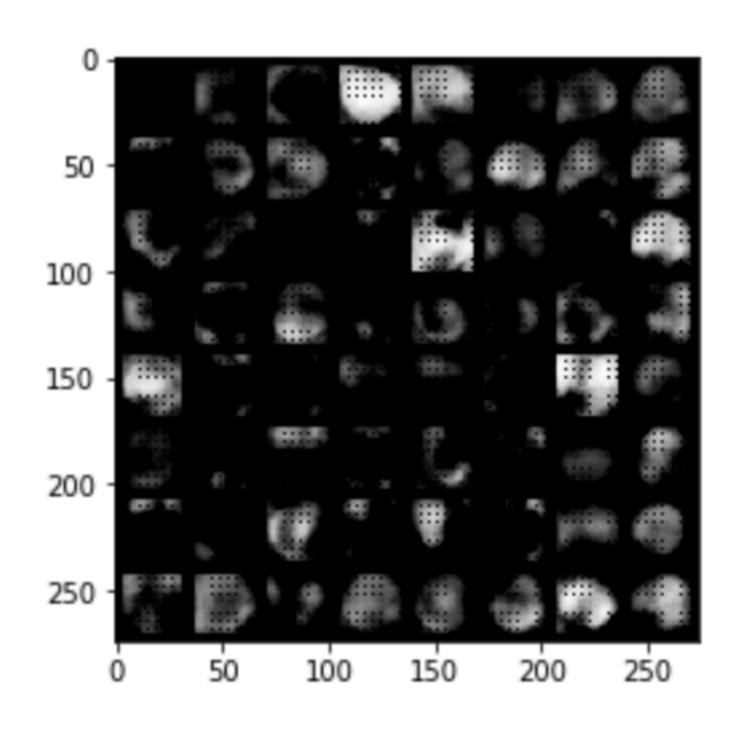


epoch=2

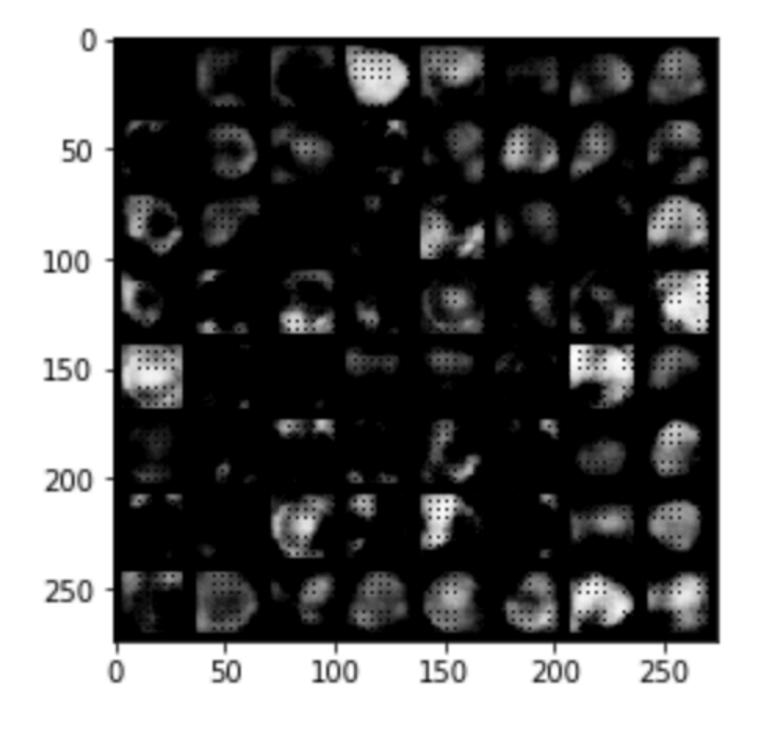
#### Results

<matplotlib.image.AxesImage at 0x7fec7a699e90>

<matplotlib.image.AxesImage at 0x7fec7ab21f90>



epoch=3



epoch=4