

# Exploring the Impact of Deep Learning Data Augmentation on Medical Imagery Classification

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**Abstract.** We evaluate the impact of data augmentation methods on the performance of a Convolutional Neural Network in 8 medical image classification tasks. We find that in marginally improved performance on some datasets when using either Geometric or Deep Learning data augmentation methods<sup>1</sup>.

## 1 Introduction

MedMNIST is a collection of labelled biomedical image datasets [7] standardized to fit the MNIST format [9] (see Table 1). The authors created MedMNIST to offer a suite of medical data applicable to supervised classification tasks and the comparison of classifier algorithms as part of a benchmark<sup>2</sup>. In this context, we are interested in studying the impact of data augmentation on the performance of such classifiers.

Data augmentation techniques are broadly split in two families: geometric transformations (e.g. rotations, deformations, etc.) and Deep Learning (DL) approaches. In this study, we select a few geometric and DL data augmentation techniques to assess whether potential performance improvements of a given classifier can be observed.

We retain 8 two-dimensional MedMNIST datasets for this study (see Table 1). As such, the next two sections will respectively introduce our methodology and provide preliminary results and observations.

**Table 1.** Retained MedMNIST datasets for a given classification task

Dataset	Type (Fig. with montage)	Task	Subset size		
			Training	Validation	Test
PathMNIST	Colon pathology (Fig. 1)	Multi (9)	89,996	10,004	7,180
DermaMNIST	Dermatoscope (Fig. 2)	Multi (7)	7,007	1,003	2,005
OctMNIST	Retinal OCT (Fig. 3)	Multi (4)	97,477	10,832	1,000
PneumoniaMNIST	Chest X-ray (Fig. 4)	Binary (2)	4,708	524	624
BreastMNIST	Breast Ultrasound (Fig. 5)	Binary (2)	546	78	156
OrganAMNIST	Axial Abdo. CT (Fig. 6)	Multi (11)	34,581	6,491	17,778
OrganCMNIST	Coronal Abdo. CT (Fig. 7)	Multi (11)	13,000	2,392	8,268
OrganSMNIST	Sagittal Abdo. CT (Fig. 8)	Multi (11)	13,940	2,452	8,829

## 2 Methodology

### 2.1 A Baseline Classifier

We reuse the Pytorch Convolutional Neural Network (CNN) [13,14] provided by the MedMNIST team on their GitHub page [8]. The CNN depth and hyperparameters are

<sup>1</sup> All code and results used in this document are available on Google Colab: <https://colab.research.google.com/drive/1J64f1Vq0ALWS7JBd8hj5qF1bwdbHlmeR?usp=sharing>

<sup>2</sup> Benchmark page: <https://medmnist.com/>

fixed to control for each modelization factor besides our selection of data augmentation methods. The only divergence between implementations relates to the datasets' number of classes which modulates a CNN's output size. An example of such CNN implementation for the PathMNIST dataset is available in Fig. 9.

Each CNN implementation is trained for 20 epochs with a Stochastic Gradient Descent optimizer [1] with a 0.001 learning rate and 0.9 momentum. The loss is computed via cross-entropy (binary cross-entropy in the case of binary classification). Batch size is set to 64. We reuse the MedMNIST authors' custom evaluator object, which provides two output metrics to assess model performance: accuracy and area-under-the-curve (AUC). When testing an implementation on a test set, we use the model that yielded the best validation accuracy.

## 2.2 Dealing with Imbalancedness with Weighted Random Sampling

The 8 retained datasets display class imbalances (see Fig. 10,11,12,13,14,15,16,17). Furthermore, the amount of available datapoints in a dataset can be small (e.g. the BreastMNIST dataset has less than 1,000 points). Such imbalance and lack of datapoints may impact a classifier's convergence towards an optimum [15]. As such, image data augmentation techniques may help improve a model's generalization capability by reducing overfitting to the training data [18].

To further help reduce the effect of class imbalance, we will also include to our process PyTorch's Weighted Random Sampling (WRS) sampler object. This allows us to deal with class imbalance at the dataloader level. We will evaluate all our CNN setups with and without WRS.

## 2.3 Augmenting Datasets: Geometric Approaches

To implement setups with geometric data augmentation, we rely on Pytorch's Transforms class at the dataloader level. We select two types of geometric augmentation pipeline that is conditioned on the underlying MedMNIST dataset.

In the case of a black-and-white image dataset, the corresponding geometric data augmentation is a 3-step pipeline: random sharpness adjustment, random contrast then random horizontal flipping, all with default Pytorch parameters. In the case of a color image dataset, the geometric data augmentation is identical to the black-and-white pipeline but with a color hue jitter of 0.05 added after sharpness and contrast adjustments. Each pipeline ends with a normalization step.

## 2.4 Augmenting Datasets: VAE and GAN Approaches

The Manifold Hypothesis states that high-dimensional data are representations of lower dimensional objects [4].

Variational Autoencoders (VAEs) assume that a random process involving a continuous random variable has generated a dataset. A latent space representation of the data can be learned via a pair of encoder and decoder networks [2].

Meanwhile, Generative Adversarial Networks (GANs) generate new data using adversarial training, relying on a pair of generator and discriminator networks to

learn a new data generating process [6]. Deep Convolutional GANs have been shown to improve medical data classification [5,20].

We are interested in implementing both types of methods for our DL data augmentation setups. As such, we settle on 3 methods: Conditional VAE, Joint VAE, and Conditional GAN.

**Conditional VAEs** learn a latent representation of a dataset where the underlying random process is informed by the data classes [16]. Conditional VAEs use classes as input (see Fig. 18 for an example representation of a Conditional VAE, and Fig. 19 and Fig. 20 for our implementation in the PathMNIST case).

Instead of learning the prior distribution  $P_\theta(z)$  of the latent space  $z$  (with the data  $x$  being generated by the distribution  $P_\theta(x|z)$ ), a Conditional VAE learns  $P_\theta(z|y)$  with  $y$  the data’s labels conditioning the latent space. As such, Conditional VAEs allow us to generate new labelled data.

**Joint VAEs** learn both a continuous and discrete representation of a dataset [3]. As such, a high-dimensional dataset can be represented in a low-dimensional latent space with continuous and discrete elements. A Joint VAE learns the prior distribution  $P_\theta(z, c)$  of the continuous space  $z$  and discrete/categorical space  $c$  with the data  $x$  being generated by the distribution  $P_\theta(x|z, c)$  (see Fig. 21 for an example representation of a Joint VAE, and Fig. 22 and Fig. 23 for our implementation in the PathMNIST case).

A Joint VAE may help capture medical images’ characteristics that a purely continuous latent space might not. Indeed, medical images are usually obtained in strictly controlled environments where some variables are discrete such as the left-right orientation of a chest CT scan (discrete).

The main drawback is the absence of conditioning on classes. We can either train one Joint VAE per dataset class or train a single Joint VAE on a whole dataset. The latter case, though less computationally expensive, implies that we do not have the ability to categorically condition the underlying generative process.

**Conditional GANs** are similar to Conditional VAEs. They generate data conditioned on its classes [10] (see Fig. 24 for an example representation of a Conditional GAN, and Fig. 25, and Fig. 26 for our implementation in the PathMNIST case).

Conditional GANs rely on using labels as inputs to guide the generator and discriminator’s learning and the subsequent generative process [11,12,17]. The discriminator is evaluated on the similarity between generated and real data, and the match between the generated image’s predicted and true labels. The rationale behind using a Conditional GAN is similar to that of the Conditional VAE: we want to generate new labelled data as part of a data augmentation process.

## VAE and GAN Training Setup

Our two VAEs rely on existing implementations available online [19]. We use the TorchFusion library for the Conditional GAN [12]. We train the two VAEs for 200 epochs (with early stopping), and the Conditional GAN for 100. The Conditional

VAE is trained with a latent space of dimension 150, the Joint VAE with continuous and discrete latent spaces of dimensions 100 and 50 respectively, and the Conditional GAN with a latent space of dimension 128.

### Data Augmentation Setup

As a result of training (see Fig. 27 for the three methods’ loss per epoch in the PathMNIST case), we generate new images for each retained MedMNIST datasets. We only augment the training sets; validation and test sets are untouched.

The Conditional VAE and GAN generate new labelled data in amounts inversely proportional to the scarcity of the class in each original datasets. We add to each dataset (in the order stated in Table 1) the following amounts of images: 4%, 10%, 4%, 10%, 10%, 5%, 10%, 10%, i.e. 3599, 700, 3899, 470, 54, 1729, 1300, and 1394 images (see Fig. 28 for examples of generated data for the OctMNIST dataset).

The Joint VAE augments the MedMNIST datasets by interweaving the original images with their VAE reconstructions. We expect this setup to increase intra-class variability, which may help our CNN implementations yield better results.

## 3 Results

We reproduce the Accuracy and Area under the Curve test set results for each dataset and model (80 different cases) in the Tables 2, 3, and 4. The best results for each dataset are highlighted in bold.

Our first observation is that WRS generally improves model performance on 6 out of 8 datasets when no data augmentation is involved. This shows that class imbalance is an existing problem to account for in the MedMNIST’s collection and benchmark.

We also find that in all cases but BreastMNIST and OctMNIST, a data augmentation method (whether geometric or deep learning based) improves model performance over the baseline when no WRS is involved. We note that using a Joint VAE never improved performance over the baseline except in the PneumoniaMNIST case.

Given all tested techniques, an accuracy improvement between 0.35 and 8.77 points over the baseline is observed on all the datasets (see Table 5).

Finally, though we could find a data augmentation technique leading to performance improvements for 7 out of 8 datasets, we actually observe a performance decrease for our classifier in most of the cases where data augmentation techniques.

## 4 Conclusion

We demonstrate that it is possible to find a data augmentation technique (furthermore a deep learning one) that improves a simple CNN classifier’s accuracy. The use of Conditional GAN and VAE improved performance on 5 out of 8 datasets, implying that conditioning data generation on classes can be relevant for classification.

The second key takeaway is that data augmentation alone does not guarantee performance improvement, however. We highlight that most implementation have shown worse results.

Finally, we underline that the MedMNIST authors insist on the data not being intended for clinical use. We must be careful with extending any preliminary conclusion especially in the case of possible critical applications in healthcare.

**Table 2.** Test Accuracy & AUC obtained with geometric data augmentation.

Dataset	no GDA or WRS		WRS, no GDA		no WRS, GDA		WRS & GDA	
	AUC	Acc.	AUC	Acc.	AUC	Acc.	AUC	Acc.
PathMNIST	97.99	84.36	96.67	79.93	98.94	87.97	<b>99.14</b>	<b>88.29</b>
DermaMNIST	89.21	65.94	89.62	<b>69.43</b>	<b>89.97</b>	66.53	89.43	64.39
OctMNIST	93.4	70.4	<b>94.29</b>	<b>76.6</b>	91.54	68.8	92.98	74.2
PneumoniaMNIST	95.8	85.58	96.07	87.98	95.74	86.38	<b>96.66</b>	<b>90.38</b>
BreastMNIST	<b>88.35</b>	82.69	86.45	<b>84.62</b>	82.14	78.21	83.02	67.95
OrganAMNIST	99.12	88.1	<b>99.14</b>	<b>88.49</b>	98.87	87.03	98.85	87.43
OrganCMNIST	98.72	86.25	<b>98.74</b>	<b>86.32</b>	98.35	83.19	98.3	81.66
OrganSMNIST	<b>97.46</b>	71.92	96.42	<b>71.94</b>	95.42	68.64	95.27	63.95

**Table 3.** Test Accuracy & AUC obtained with DL data augmentation.

Dataset	Conditional VAE		Joint VAE		Conditional GAN		Improved perf.	
	AUC	Acc.	AUC	Acc.	AUC	Acc.	over no GDA	over GDA
PathMNIST	<b>98.11</b>	<b>87.49</b>	97.9	85.08	97.76	85.52	Yes	No
DermaMNIST	90.32	<b>74.71</b>	<b>90.34</b>	73.22	90.13	71.67	Yes	Yes
OctMNIST	<b>93.53</b>	<b>72.5</b>	93.2	73.0	92.93	67.3	Yes	Yes
PneumoniaMNIST	<b>96.81</b>	<b>86.38</b>	93.52	85.58	96.27	85.42	Yes	eq.
BreastMNIST	87.49	78.85	84.02	77.56	<b>83.31</b>	<b>80.77</b>	No	Yes
OrganAMNIST	99.0	87.87	98.98	88.08	<b>99.07</b>	<b>88.54</b>	Yes	Yes
OrganCMNIST	<b>98.66</b>	<b>86.59</b>	98.43	85.87	<b>98.66</b>	85.43	Yes	Yes
OrganSMNIST	96.01	70.98	95.86	69.66	<b>96.46</b>	<b>72.27</b>	Yes	Yes

**Table 4.** Test Accuracy & AUC obtained with DL data augmentation with WRS.

Dataset	Conditional VAE		Joint VAE		Conditional GAN		Improved perf.	
	AUC	Acc.	AUC	Acc.	AUC	Acc.	over no GDA	over GDA
PathMNIST	<b>97.95</b>	<b>86.16</b>	96.12	74.81	96.28	79.54	Yes	No
DermaMNIST	<b>90.21</b>	<b>73.02</b>	88.68	66.38	88.54	68.93	Yes	Yes
OctMNIST	91.99	<b>74.1</b>	<b>93.7</b>	72.3	92.17	70.0	No	No
PneumoniaMNIST	<b>95.3</b>	84.78	94.77	<b>89.42</b>	95.87	87.02	Yes	No
BreastMNIST	85.69	82.69	86.93	80.13	<b>89.33</b>	<b>85.9</b>	Yes	Yes
OrganAMNIST	<b>99.15</b>	<b>88.57</b>	99.06	88.33	99.08	88.2	Yes	Yes
OrganCMNIST	98.53	86.07	98.54	85.27	<b>98.83</b>	<b>86.94</b>	Yes	Yes
OrganSMNIST	96.01	<b>71.57</b>	95.6	68.86	<b>96.16</b>	71.47	No	Yes

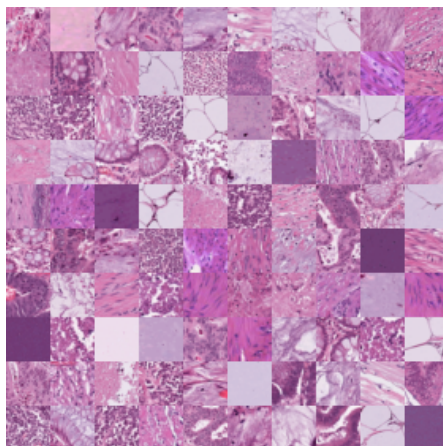
**Table 5.** Best model per dataset (based on test Accuracy results)

Dataset	Model	Acc. gain over no GDA/WRS	Loss Convergence	Confusion Matrix
PathMNIST	WRS + geometric DA	+3.93	Fig. 29	Fig. 30
DermaMNIST	Conditional VAE	+8.77	Fig. 31	Fig. 32
OctMNIST	WRS only	+6.2	Fig. 33	Fig. 34
PneumoniaMNIST	WRS + geometric DA	+4.8	Fig. 35	Fig. 36
BreastMNIST	WRS + Conditional GAN	+3.21	Fig. 37	Fig. 38
OrganAMNIST	WRS + Conditional VAE	+0.47	Fig. 39	Fig. 40
OrganCMNIST	WRS + Conditional GAN	+0.69	Fig. 41	Fig. 42
OrganSMNIST	Conditional GAN	+0.35	Fig. 43	Fig. 44

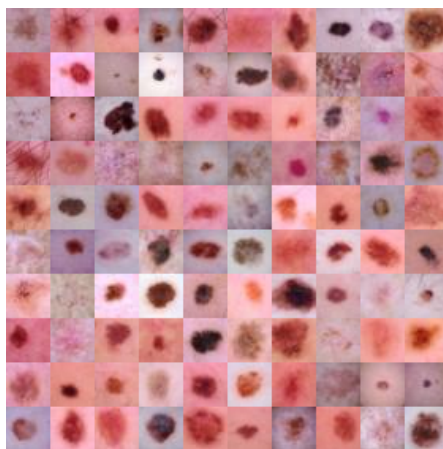
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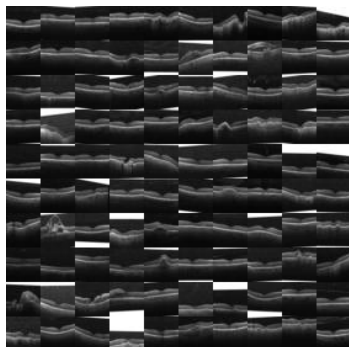
## Annex



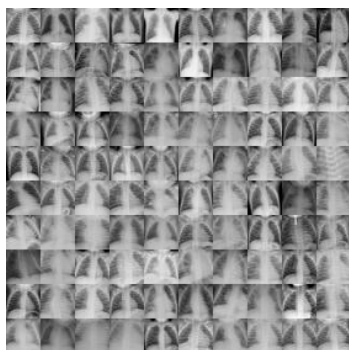
**Fig. 1.** Montage of training samples from the PathMNIST dataset.



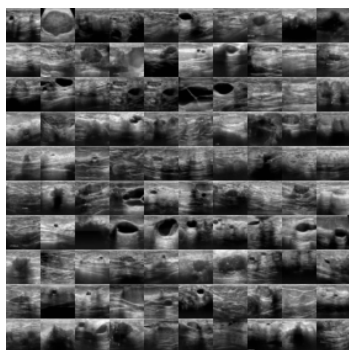
**Fig. 2.** Montage of training samples from the DermaMNIST dataset.



**Fig. 3.** Montage of training samples from the OctMNIST dataset.

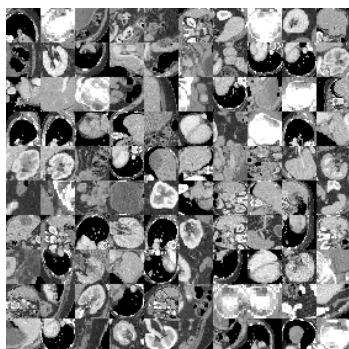


**Fig. 4.** Montage of training samples from the PneumoniaMNIST dataset.

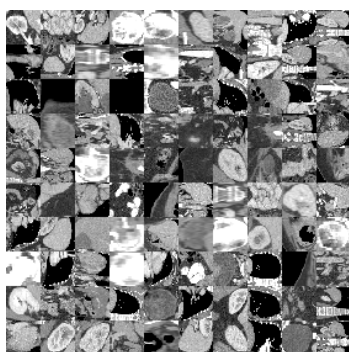


**Fig. 5.** Montage of training samples from the BreastMNIST dataset.

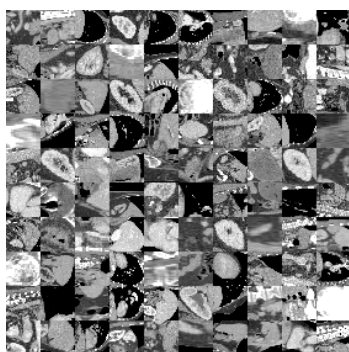




**Fig. 6.** Montage of training samples from the OrganMNIST\_Axial dataset.



**Fig. 7.** Montage of training samples from the OrganMNIST\_Coronal dataset.



**Fig. 8.** Montage of training samples from the OrganMNIST\_Sagittal dataset.

Layer (type)	Output Shape	Param #
Conv2d-1	[-1, 16, 26, 26]	448
BatchNorm2d-2	[-1, 16, 26, 26]	32
ReLU-3	[-1, 16, 26, 26]	0
Conv2d-4	[-1, 16, 24, 24]	2,320
BatchNorm2d-5	[-1, 16, 24, 24]	32
ReLU-6	[-1, 16, 24, 24]	0
MaxPool2d-7	[-1, 16, 12, 12]	0
Conv2d-8	[-1, 64, 10, 10]	9,280
BatchNorm2d-9	[-1, 64, 10, 10]	128
ReLU-10	[-1, 64, 10, 10]	0
Conv2d-11	[-1, 64, 8, 8]	36,928
BatchNorm2d-12	[-1, 64, 8, 8]	128
ReLU-13	[-1, 64, 8, 8]	0
Conv2d-14	[-1, 64, 8, 8]	36,928
BatchNorm2d-15	[-1, 64, 8, 8]	128
ReLU-16	[-1, 64, 8, 8]	0
MaxPool2d-17	[-1, 64, 4, 4]	0
Linear-18	[-1, 128]	131,200
ReLU-19	[-1, 128]	0
Linear-20	[-1, 128]	16,512
ReLU-21	[-1, 128]	0
Linear-22	[-1, 9]	1,161
Total params: 235,225		
Trainable params: 235,225		
Non-trainable params: 0		
Input size (MB): 0.01		
Forward/backward pass size (MB): 0.82		
Params size (MB): 0.90		
Estimated Total Size (MB): 1.73		

**Fig. 9.** Convolutional Neural Network classifier used on the PathMNIST dataset.

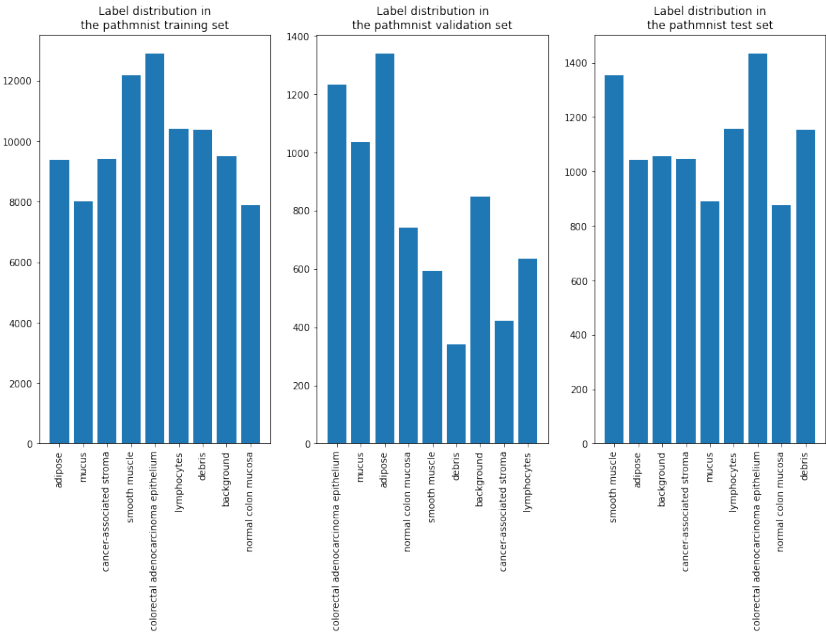


Fig. 10. Class distribution of the PathMNIST dataset.

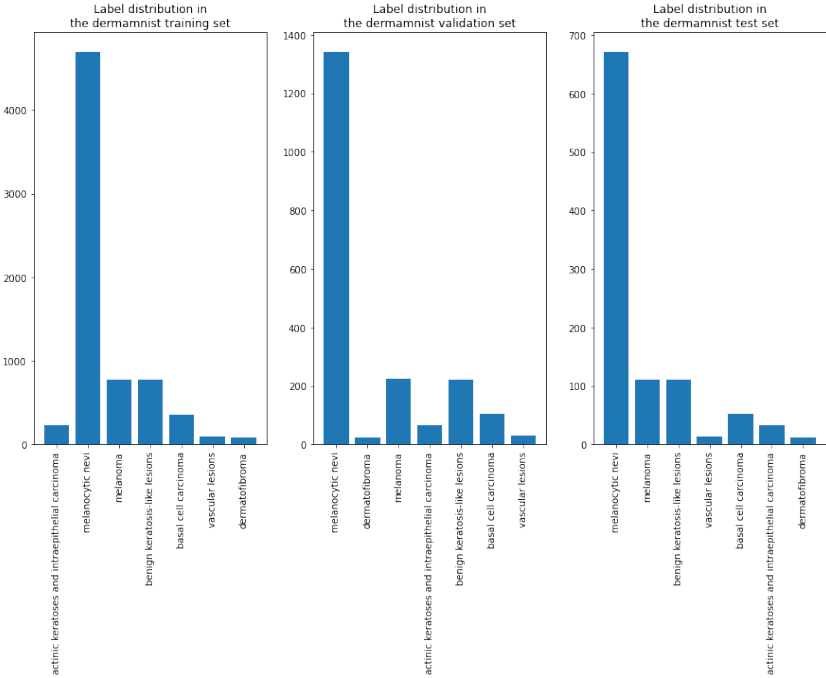
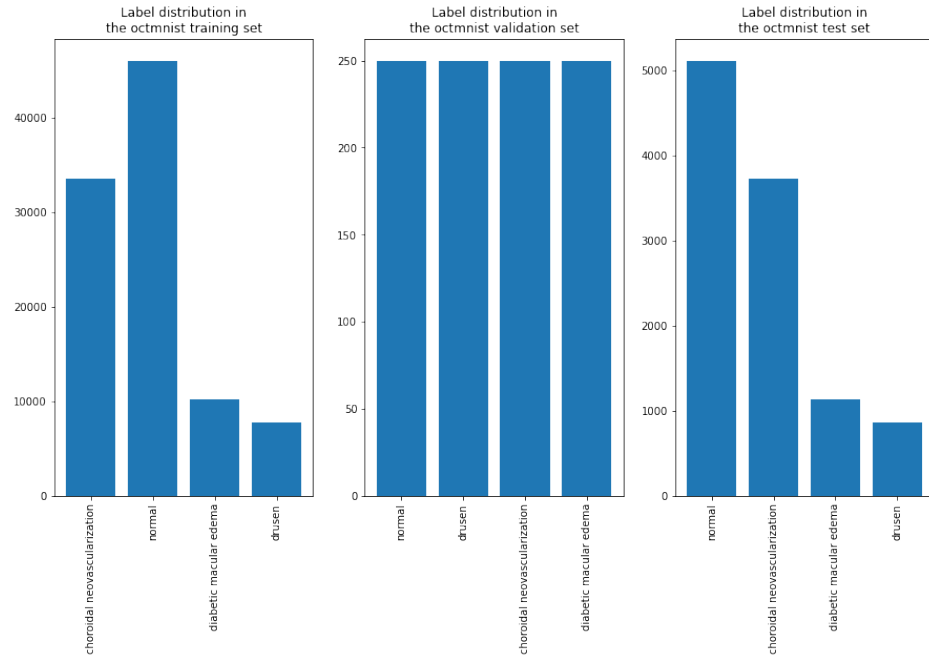
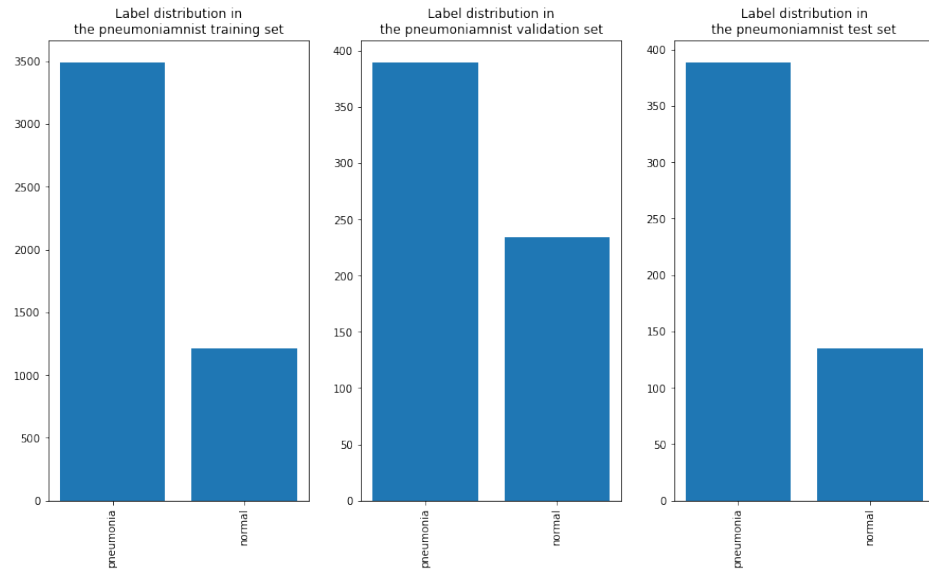


Fig. 11. Class distribution of the DermaMNIST dataset.



**Fig. 12.** Montage of training samples from the OctMNIST dataset.



**Fig. 13.** Montage of training samples from the PneumoniaMNIST dataset.

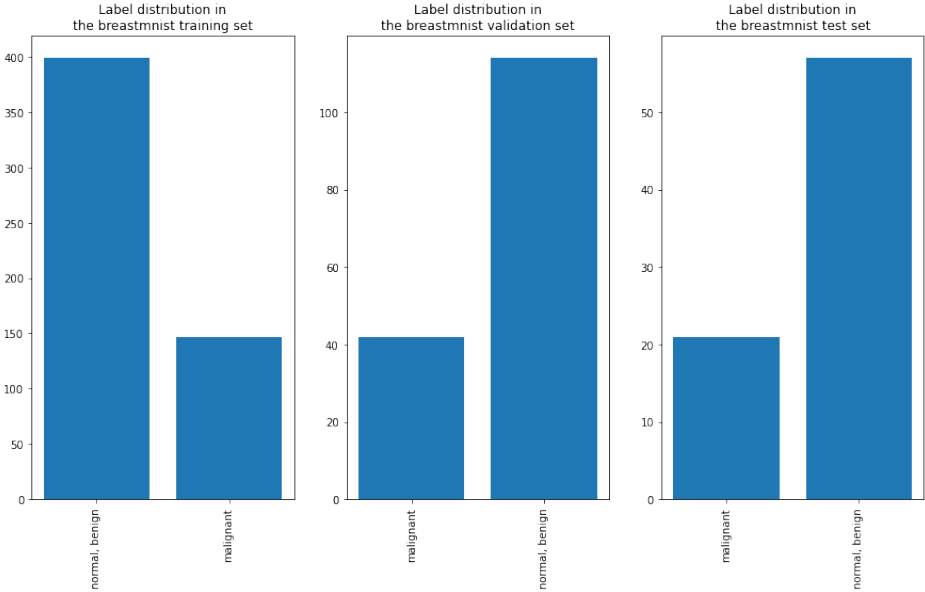


Fig. 14. Class distribution of the BreastMNIST dataset.

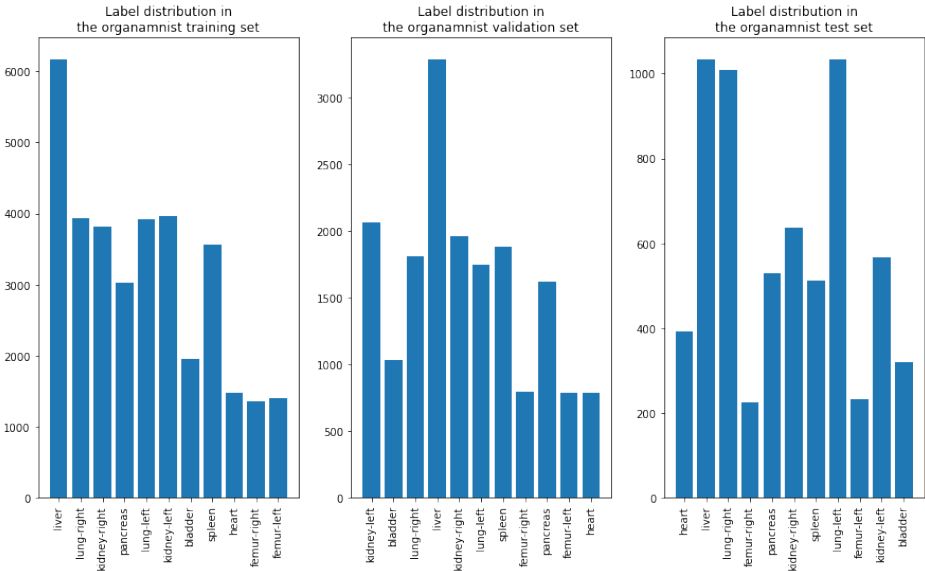
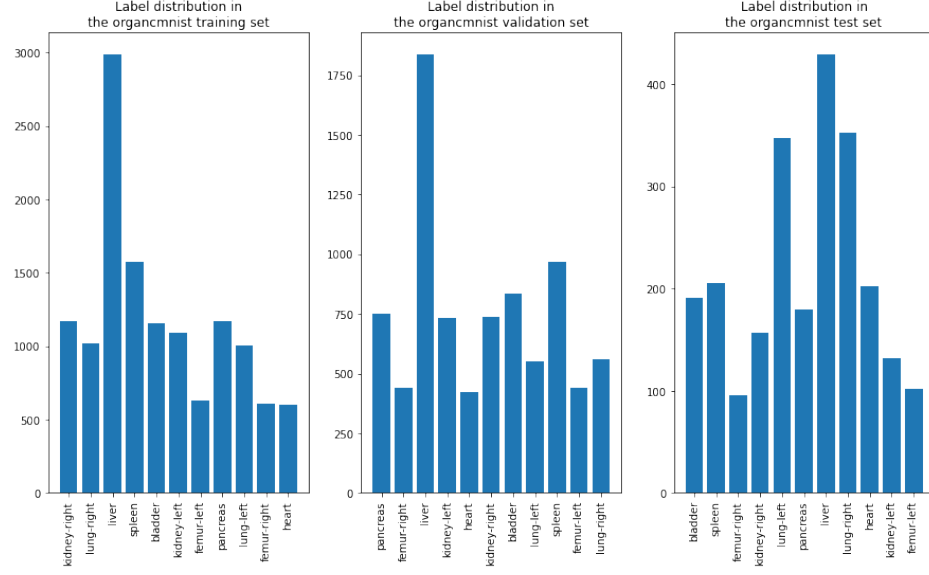
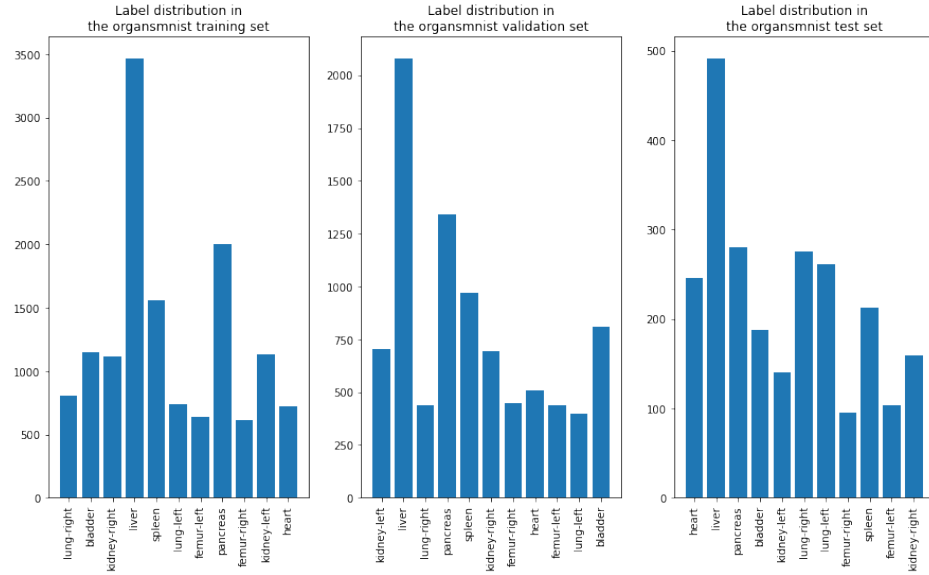


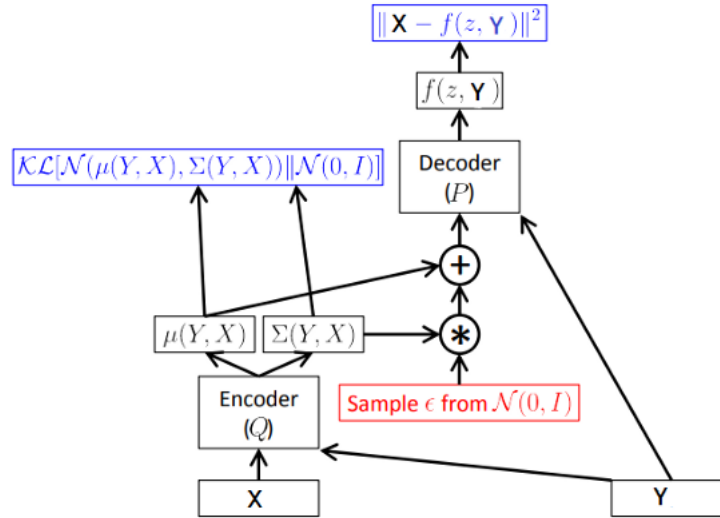
Fig. 15. Class distribution of the OrganMNIST\_Axial dataset.



**Fig. 16.** Class distribution of the OrganMNIST\_Coronal dataset.



**Fig. 17.** Class distribution of the OrganMNIST\_Sagittal dataset.



**Fig. 18.** Reproduction of the graphical representation of a Conditional Variational Auto-Encoder[2].

Decoder input layer:

Linear(in\_features=109, out\_features=2048, bias=True)

Decoder hidden layers:

Layer (type)	Output Shape	Param #
ConvTranspose2d-1	[-1, 256, 4, 4]	1,179,904
BatchNorm2d-2	[-1, 256, 4, 4]	512
LeakyReLU-3	[-1, 256, 4, 4]	0
ConvTranspose2d-4	[-1, 128, 8, 8]	295,040
BatchNorm2d-5	[-1, 128, 8, 8]	256
LeakyReLU-6	[-1, 128, 8, 8]	0
ConvTranspose2d-7	[-1, 64, 16, 16]	73,792
BatchNorm2d-8	[-1, 64, 16, 16]	128
LeakyReLU-9	[-1, 64, 16, 16]	0
ConvTranspose2d-10	[-1, 32, 32, 32]	18,464
BatchNorm2d-11	[-1, 32, 32, 32]	64
LeakyReLU-12	[-1, 32, 32, 32]	0

Total params: 1,568,160

Trainable params: 1,568,160

Non-trainable params: 0

Input size (MB): 0.01

Forward/backward pass size (MB): 1.41

Params size (MB): 5.98

Estimated Total Size (MB): 7.40

Decoder output layer:

Layer (type)	Output Shape	Param #
ConvTranspose2d-1	[-1, 32, 61, 61]	9,248
BatchNorm2d-2	[-1, 32, 61, 61]	64
LeakyReLU-3	[-1, 32, 61, 61]	0
Conv2d-4	[-1, 32, 30, 30]	9,248
BatchNorm2d-5	[-1, 32, 30, 30]	64
LeakyReLU-6	[-1, 32, 30, 30]	0
Conv2d-7	[-1, 3, 28, 28]	867
Tanh-8	[-1, 3, 28, 28]	0

Total params: 19,491

Trainable params: 19,491

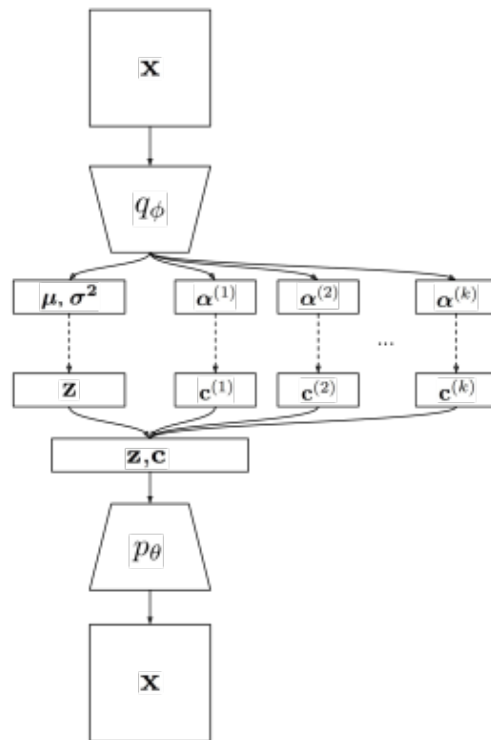
Non-trainable params: 0

**Fig. 19.** Decoder architecture of the Conditional VAE in the case of the PathMNIST dataset.



Layer (type)	Output Shape	Param #
Conv2d-1	[-1, 32, 14, 14]	1,184
BatchNorm2d-2	[-1, 32, 14, 14]	64
LeakyReLU-3	[-1, 32, 14, 14]	0
Conv2d-4	[-1, 64, 7, 7]	18,496
BatchNorm2d-5	[-1, 64, 7, 7]	128
LeakyReLU-6	[-1, 64, 7, 7]	0
Conv2d-7	[-1, 128, 4, 4]	73,856
BatchNorm2d-8	[-1, 128, 4, 4]	256
LeakyReLU-9	[-1, 128, 4, 4]	0
Conv2d-10	[-1, 256, 2, 2]	295,168
BatchNorm2d-11	[-1, 256, 2, 2]	512
LeakyReLU-12	[-1, 256, 2, 2]	0
Conv2d-13	[-1, 512, 1, 1]	1,180,160
BatchNorm2d-14	[-1, 512, 1, 1]	1,024
LeakyReLU-15	[-1, 512, 1, 1]	0
Total params: 1,570,848		
Trainable params: 1,570,848		
Non-trainable params: 0		

**Fig. 20.** Encoder architecture of the Conditional VAE in the case of the PathMNIST dataset.



**Fig. 21.** Reproduction of the graphical representation of a Joint Variational Auto-Encoder[3].

Decoder input layer:

Linear(in\_features=110, out\_features=2048, bias=True)

Decoder hidden layers:

Layer (type)	Output Shape	Param #
ConvTranspose2d-1	[-1, 256, 4, 4]	1,179,904
BatchNorm2d-2	[-1, 256, 4, 4]	512
LeakyReLU-3	[-1, 256, 4, 4]	0
ConvTranspose2d-4	[-1, 128, 8, 8]	295,040
BatchNorm2d-5	[-1, 128, 8, 8]	256
LeakyReLU-6	[-1, 128, 8, 8]	0
ConvTranspose2d-7	[-1, 64, 16, 16]	73,792
BatchNorm2d-8	[-1, 64, 16, 16]	128
LeakyReLU-9	[-1, 64, 16, 16]	0
ConvTranspose2d-10	[-1, 32, 32, 32]	18,464
BatchNorm2d-11	[-1, 32, 32, 32]	64
LeakyReLU-12	[-1, 32, 32, 32]	0

Total params: 1,568,160

Trainable params: 1,568,160

Non-trainable params: 0

Input size (MB): 0.01

Forward/backward pass size (MB): 1.41

Params size (MB): 5.98

Estimated Total Size (MB): 7.40

Decoder output layer:

Layer (type)	Output Shape	Param #
ConvTranspose2d-1	[-1, 32, 61, 61]	9,248
BatchNorm2d-2	[-1, 32, 61, 61]	64
LeakyReLU-3	[-1, 32, 61, 61]	0
Conv2d-4	[-1, 32, 30, 30]	9,248
BatchNorm2d-5	[-1, 32, 30, 30]	64
LeakyReLU-6	[-1, 32, 30, 30]	0
Conv2d-7	[-1, 3, 28, 28]	867
Tanh-8	[-1, 3, 28, 28]	0

Total params: 19,491

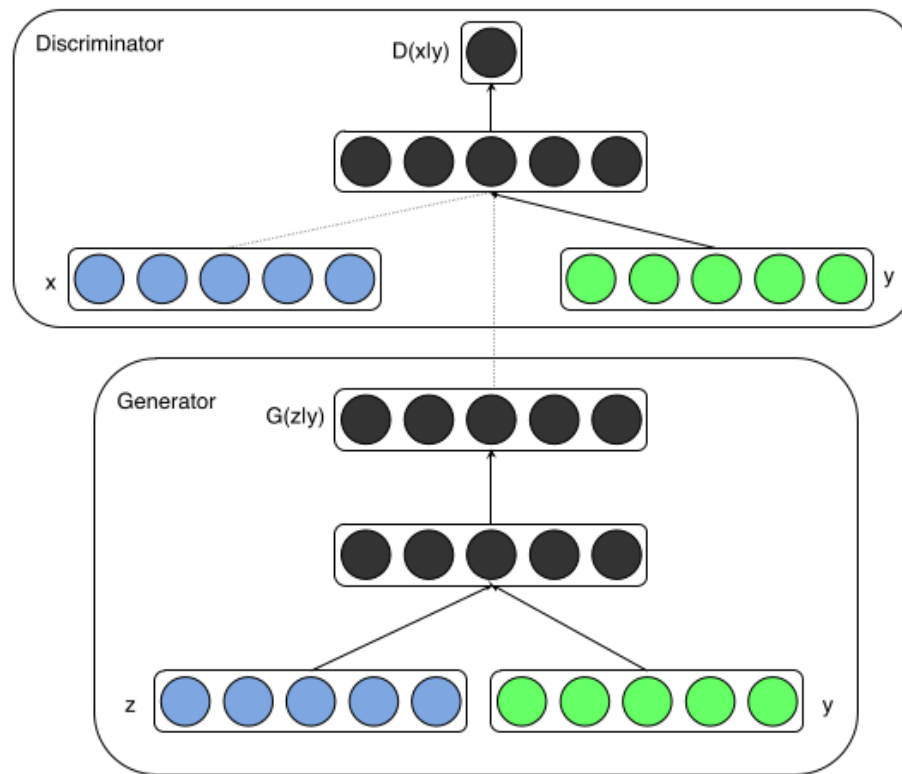
Trainable params: 19,491

Non-trainable params: 0

**Fig. 22.** Decoder architecture of the Joint VAE in the case of the PathMNIST dataset.

Layer (type)	Output Shape	Param #
Conv2d-1	[-1, 32, 14, 14]	896
BatchNorm2d-2	[-1, 32, 14, 14]	64
LeakyReLU-3	[-1, 32, 14, 14]	0
Conv2d-4	[-1, 64, 7, 7]	18,496
BatchNorm2d-5	[-1, 64, 7, 7]	128
LeakyReLU-6	[-1, 64, 7, 7]	0
Conv2d-7	[-1, 128, 4, 4]	73,856
BatchNorm2d-8	[-1, 128, 4, 4]	256
LeakyReLU-9	[-1, 128, 4, 4]	0
Conv2d-10	[-1, 256, 2, 2]	295,168
BatchNorm2d-11	[-1, 256, 2, 2]	512
LeakyReLU-12	[-1, 256, 2, 2]	0
Conv2d-13	[-1, 512, 1, 1]	1,180,160
BatchNorm2d-14	[-1, 512, 1, 1]	1,024
LeakyReLU-15	[-1, 512, 1, 1]	0
=====		
Total params: 1,570,560		
Trainable params: 1,570,560		
Non-trainable params: 0		

**Fig. 23.** Encoder architecture of the Joint VAE in the case of the PathMNIST dataset.



**Fig. 24.** Reproduction of the graphical representation of a Conditional Generative Adversarial Network[2].

Layer (type)	Output Shape	Param #
Conv2d-1	[-1, 32, 32, 32]	896
LeakyReLU-2	[-1, 32, 32, 32]	0
StandardDiscriminatorBlock-3	[-1, 32, 32, 32]	
Conv2d-4	[-1, 64, 16, 16]	32,832
LeakyReLU-5	[-1, 64, 16, 16]	0
LeakyReLU-6	[-1, 64, 16, 16]	0
LeakyReLU-7	[-1, 64, 16, 16]	0
StandardDiscriminatorBlock-8	[-1, 64, 16, 16]	
Dropout-9	[-1, 64, 16, 16]	0
Conv2d-10	[-1, 8, 16, 16]	512
Conv2d-11	[-1, 8, 16, 16]	512
Conv2d-12	[-1, 64, 16, 16]	4,096
Softmax-13	[-1, 256, 256]	0
SelfAttention-14	[-1, 64, 16, 16]	0
Conv2d-15	[-1, 128, 8, 8]	131,200
LeakyReLU-16	[-1, 128, 8, 8]	0
LeakyReLU-17	[-1, 128, 8, 8]	0
LeakyReLU-18	[-1, 128, 8, 8]	0
StandardDiscriminatorBlock-19	[-1, 128, 8, 8]	
Dropout-20	[-1, 128, 8, 8]	0
Conv2d-21	[-1, 256, 4, 4]	524,544
LeakyReLU-22	[-1, 256, 4, 4]	0
LeakyReLU-23	[-1, 256, 4, 4]	0
LeakyReLU-24	[-1, 256, 4, 4]	0
StandardDiscriminatorBlock-25	[-1, 256, 4, 4]	
Dropout-26	[-1, 256, 4, 4]	0
Flatten-27	[-1, 4096]	0
Linear-28	[-1, 1]	4,097
Embedding-29	[-1, 1, 4096]	36,864
Linear-28	[-1, 9]	36,873
=====		
Total params: 772,426		
Trainable params: 772,426		
Non-trainable params: 0		

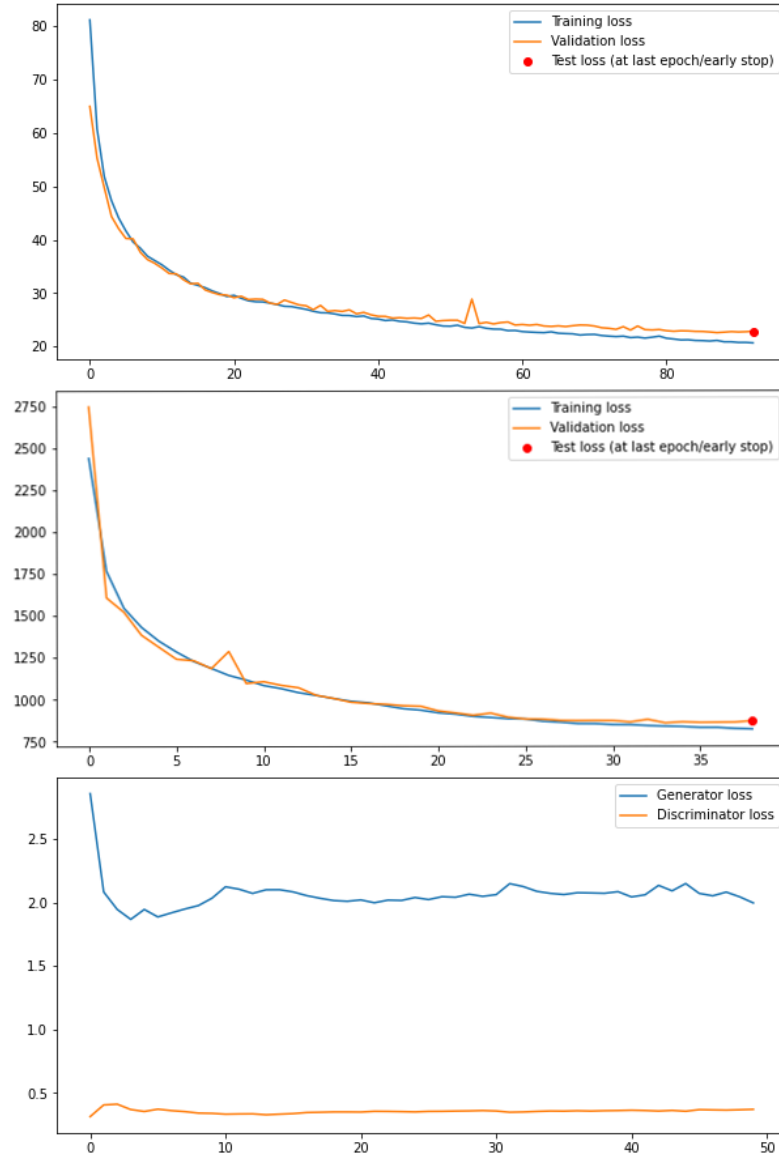
**Fig. 25.** Discriminator architecture of the Conditional GAN<sup>†</sup> in the case of the PathMNIST dataset.

<sup>†</sup>To work with the TorchFusion library, the width and height of the datasets' images are upscaled from 28 to 32

Layer (type)	Output Shape	Param #
ConvTranspose2d-1	[-1, 256, 4, 4]	409,856
BatchNorm2d-2	[-1, 256, 4, 4]	0
Embedding-3	[-1, 1, 256]	2,304
Embedding-4	[-1, 1, 256]	2,304
ConditionalBatchNorm2d-5	[-1, 256, 4, 4]	
LeakyReLU-6	[-1, 256, 4, 4]	0
LeakyReLU-7	[-1, 256, 4, 4]	0
LeakyReLU-8	[-1, 256, 4, 4]	0
StandardGeneratorBlock-9	[-1, 256, 4, 4]	
ConvTranspose2d-10	[-1, 128, 8, 8]	524,416
BatchNorm2d-11	[-1, 128, 8, 8]	0
Embedding-12	[-1, 1, 128]	1,152
Embedding-13	[-1, 1, 128]	1,152
ConditionalBatchNorm2d-14	[-1, 128, 8, 8]	
LeakyReLU-15	[-1, 128, 8, 8]	0
LeakyReLU-16	[-1, 128, 8, 8]	0
LeakyReLU-17	[-1, 128, 8, 8]	0
StandardGeneratorBlock-18	[-1, 128, 8, 8]	
Dropout-19	[-1, 128, 8, 8]	0
ConvTranspose2d-20	[-1, 64, 16, 16]	131,136
BatchNorm2d-21	[-1, 64, 16, 16]	0
Embedding-22	[-1, 1, 64]	576
Embedding-23	[-1, 1, 64]	576
ConditionalBatchNorm2d-24	[-1, 64, 16, 16]	
LeakyReLU-25	[-1, 64, 16, 16]	0
LeakyReLU-26	[-1, 64, 16, 16]	0
LeakyReLU-27	[-1, 64, 16, 16]	0
StandardGeneratorBlock-28	[-1, 64, 16, 16]	
Dropout-29	[-1, 64, 16, 16]	0
ConvTranspose2d-30	[-1, 3, 32, 32]	3,075
Total params: 1,076,547		
Trainable params: 1,076,547		
Non-trainable params: 0		

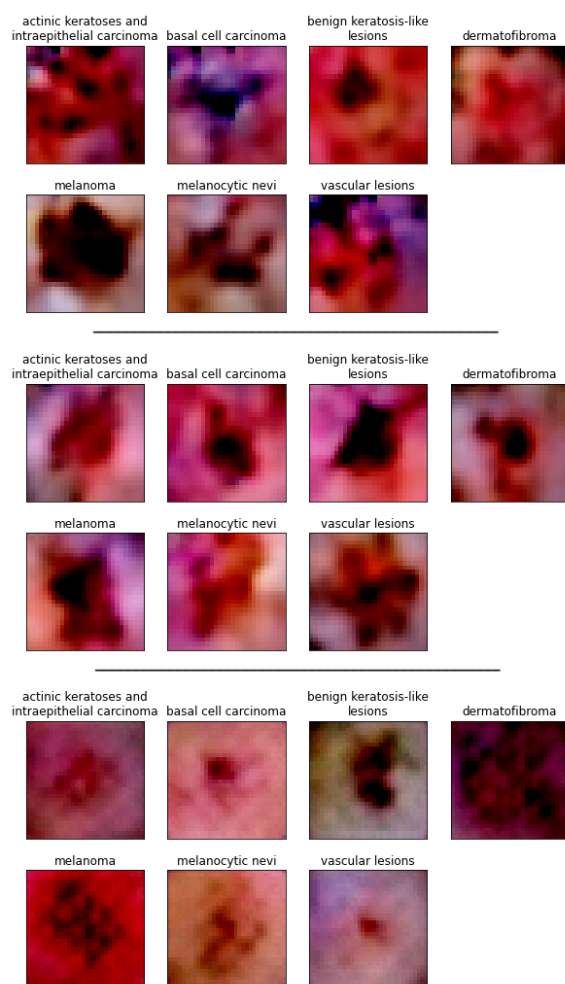
**Fig. 26.** Generator architecture of the Conditional GAN<sup>†</sup> in the case of the PathMNIST dataset.

<sup>†</sup>To work with the TorchFusion library, the width and height of the generated images are set to 32, and are subsequently downsampled to 28 to work with the baseline classifier

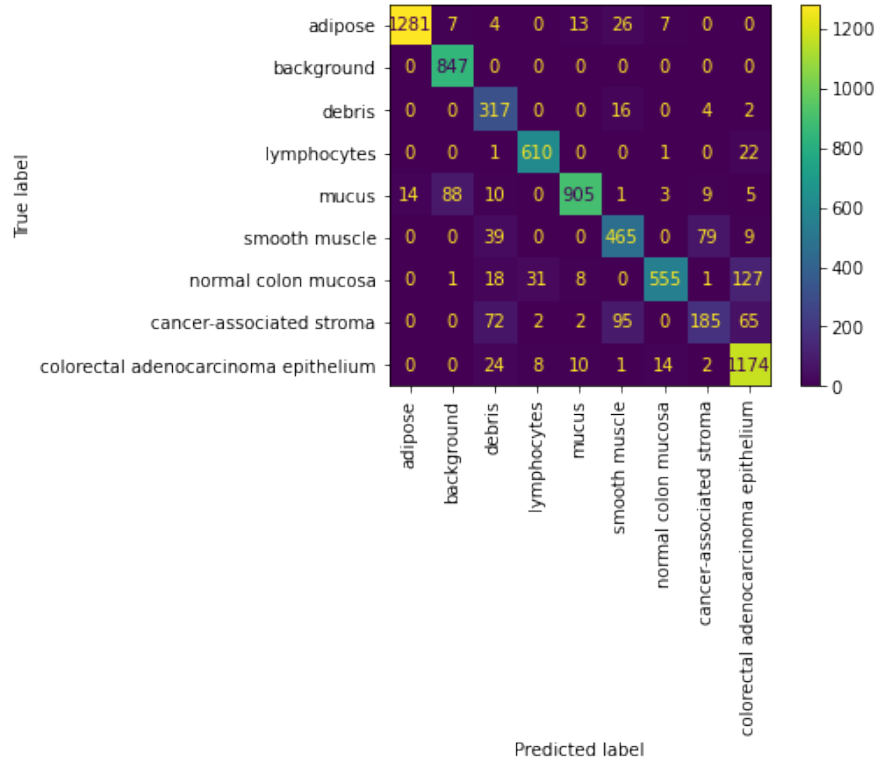


**Fig. 27.** Training loss of (in order) the conditional VAE, joint VAE, and conditional GAN on the PathMNIST dataset

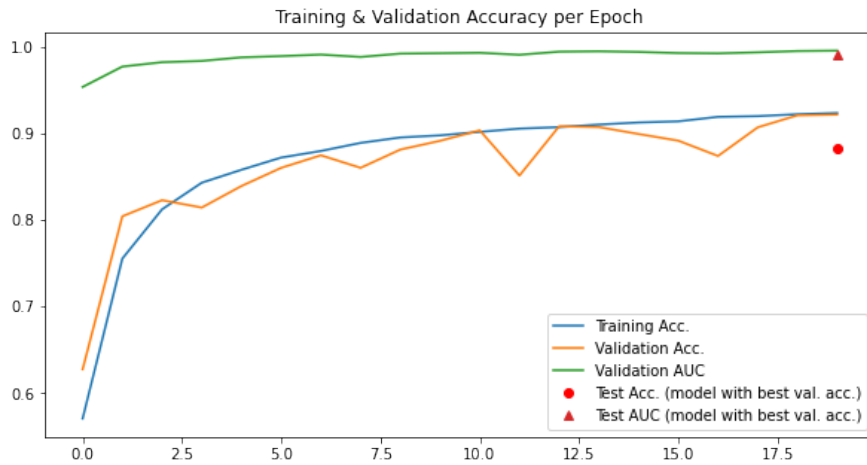




**Fig. 28.** Example of generated data for the DermaMNIST dataset for each data augmentation models in order: Conditional VAE, Joint VAE, Conditional GAN



**Fig. 29.** Confusion Matrix for the best PathMNIST model (Weighted Random Sampling + Geometric Data Augmentation)



**Fig. 30.** Accuracy and AUC per epoch for the best PathMNIST model (Weighted Random Sampling + Geometric Data Augmentation)

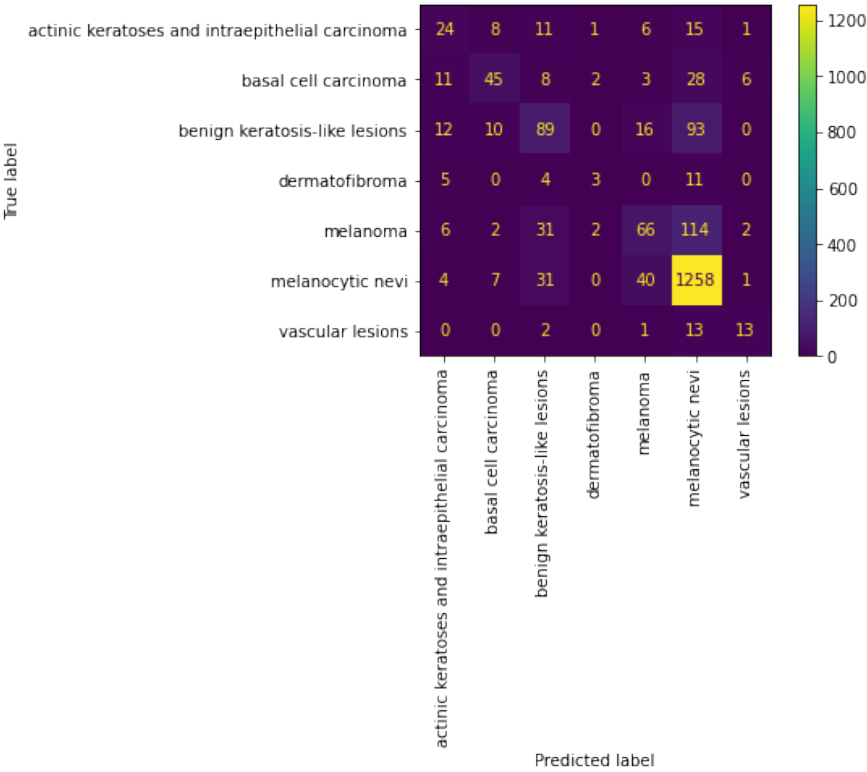


Fig. 31. Confusion Matrix for the best DermaMNIST model (Conditional VAE)

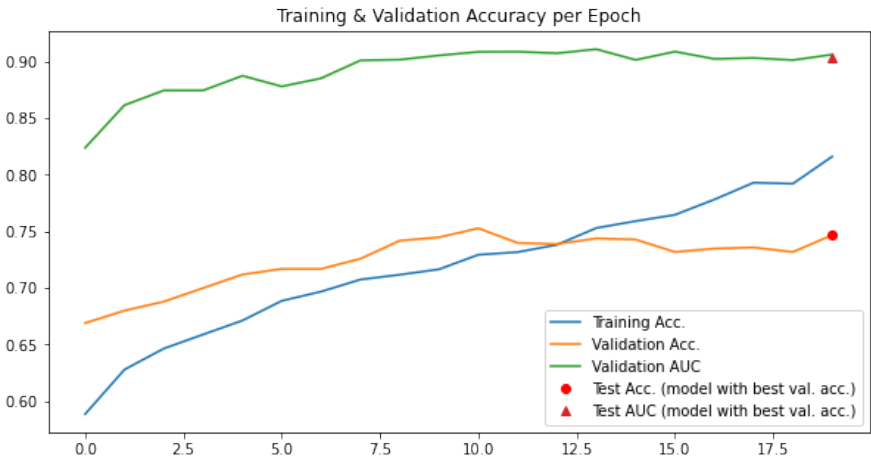
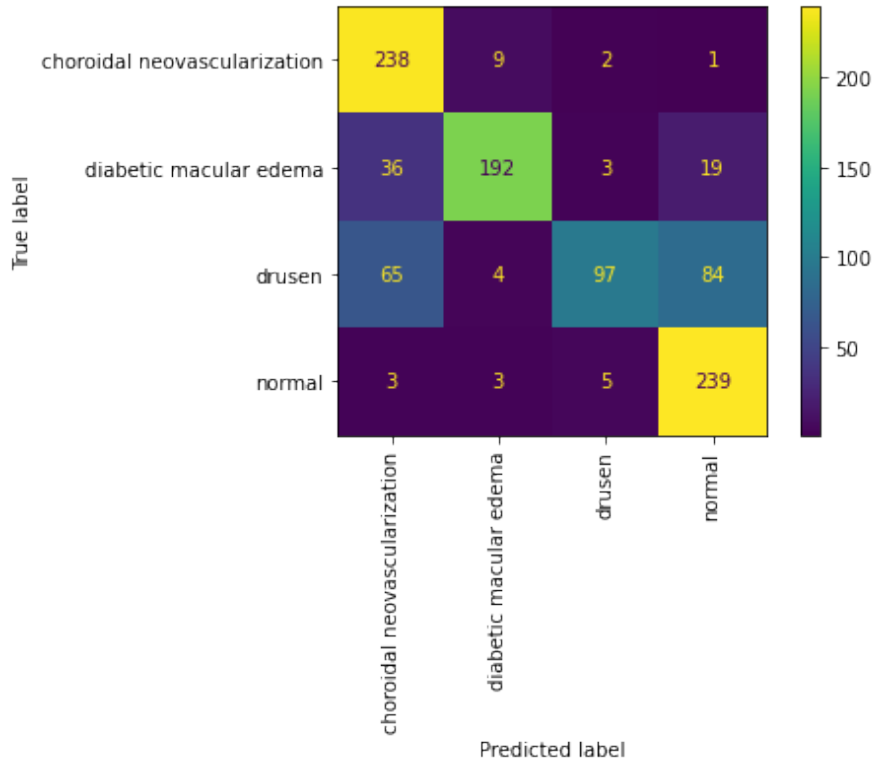
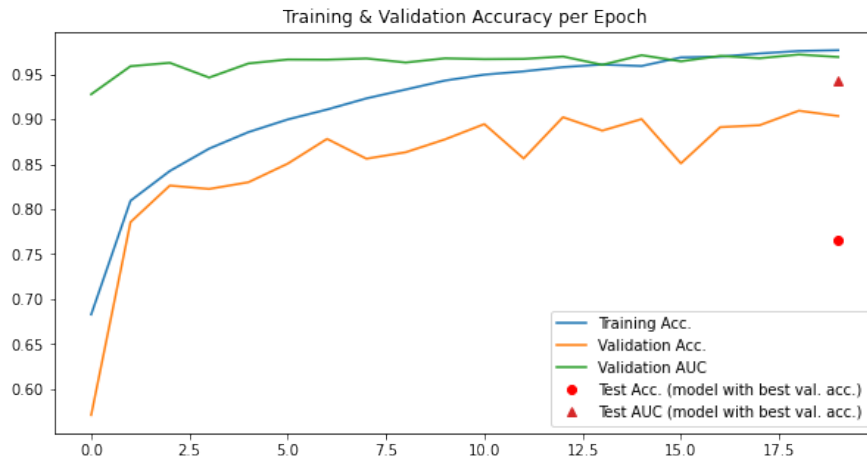


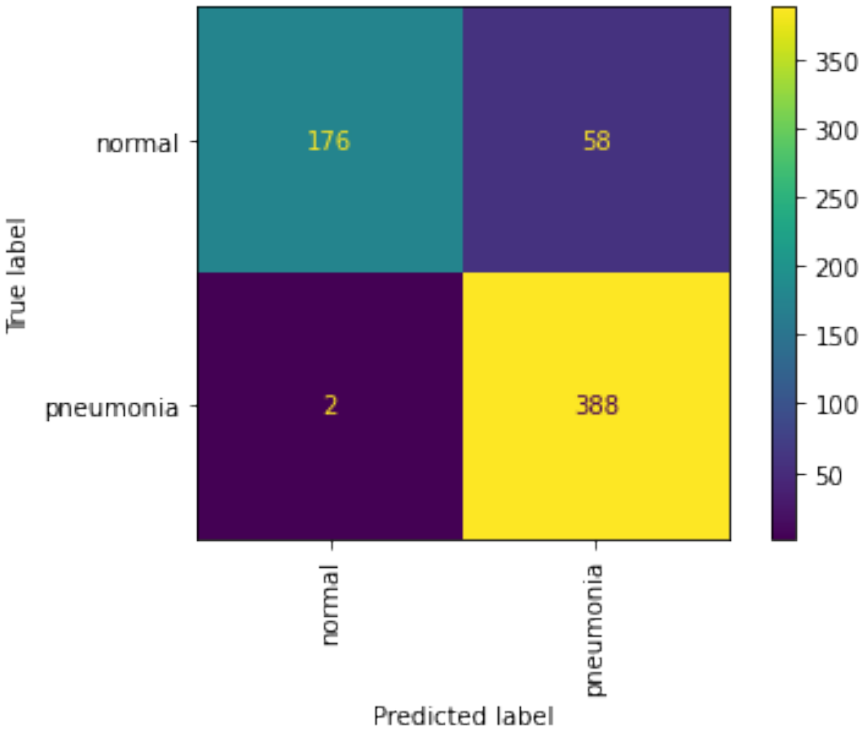
Fig. 32. Accuracy and AUC per epoch for the best DermaMNIST model (Conditional VAE)



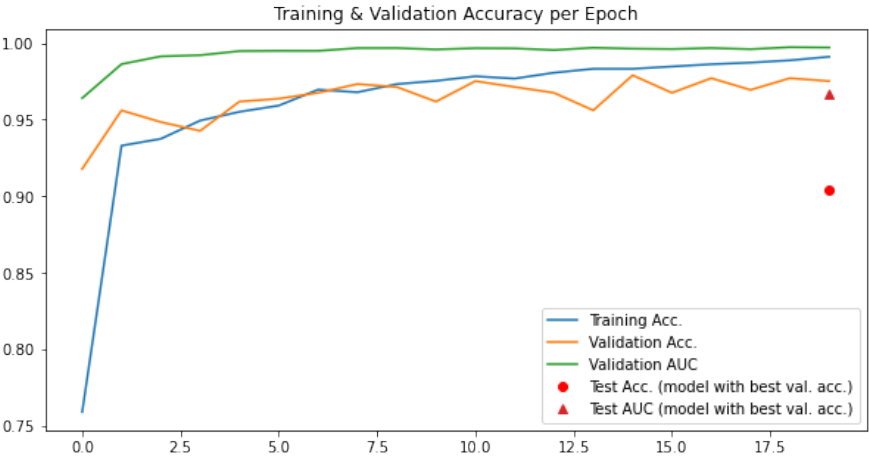
**Fig. 33.** Confusion Matrix for the best DermaMNIST model (Weighted Random Sampling)



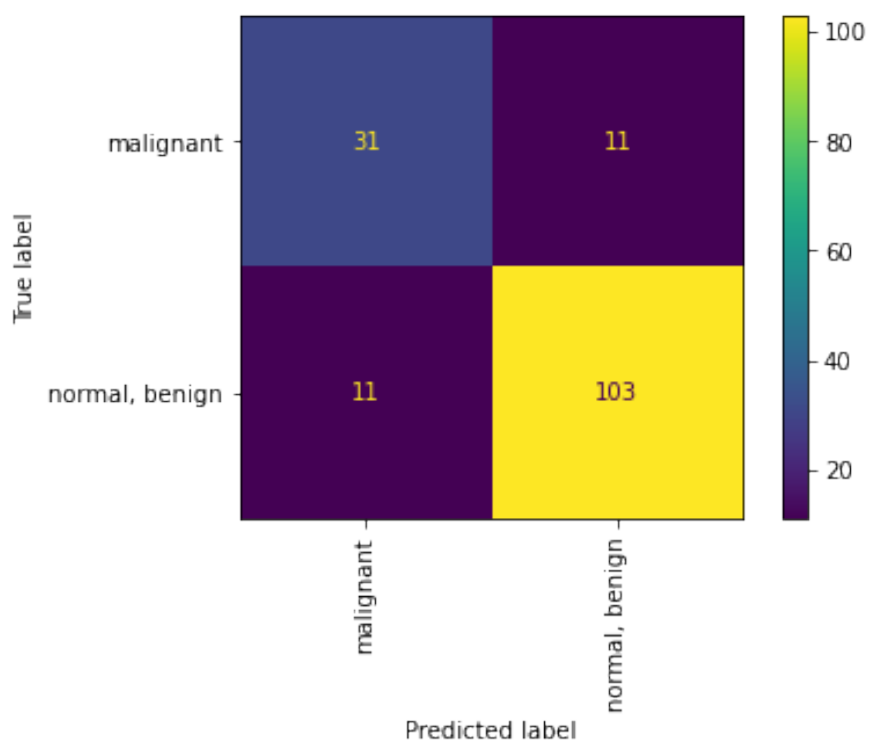
**Fig. 34.** Accuracy and AUC per epoch for the best DermaMNIST model (Weighted Random Sampling)



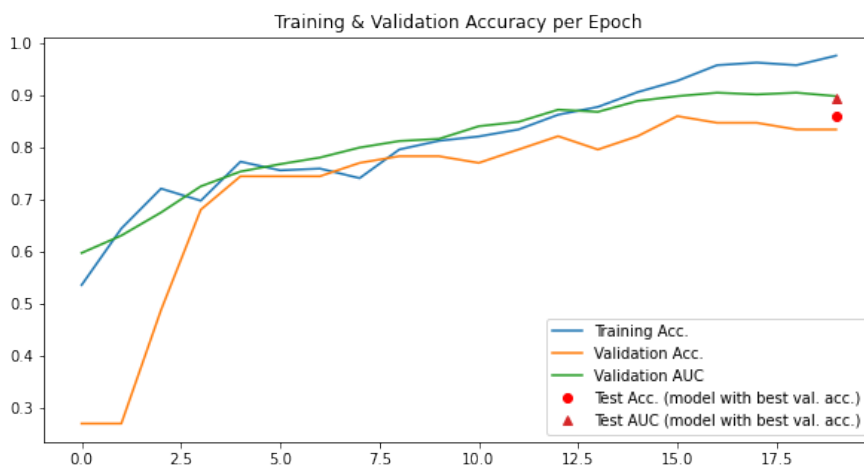
**Fig. 35.** Confusion Matrix for the best PneumoniaMNIST model (Weighted Random Sampling + geometric data augmentation)



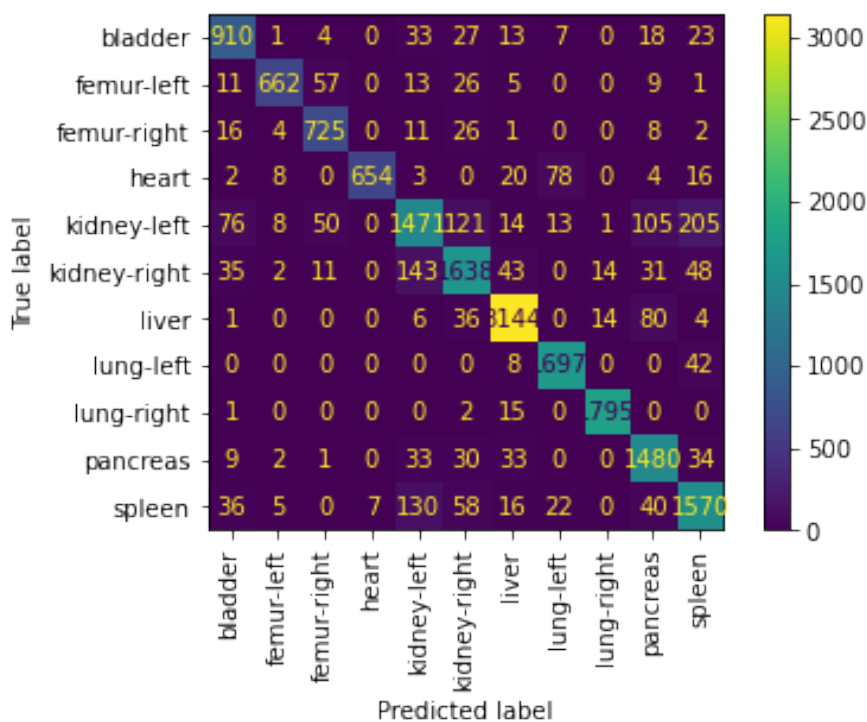
**Fig. 36.** Accuracy and AUC per epoch for the best PneumoniaMNIST model (Weighted Random Sampling + geometric data augmentation)



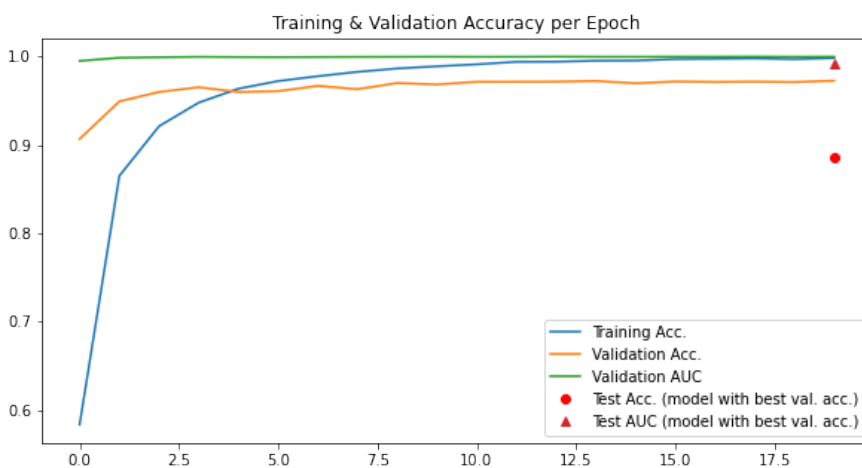
**Fig. 37.** Confusion Matrix for the best BreastMNIST model (Weighted Random Sampling + Conditional GAN)



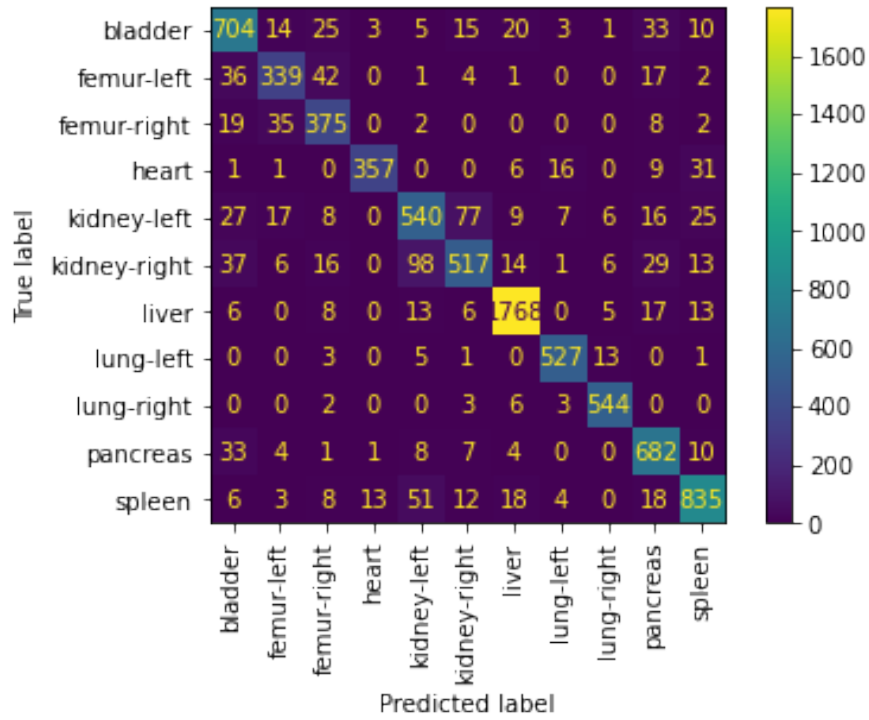
**Fig. 38.** Accuracy and AUC per epoch for the best BreastMNIST model (Weighted Random Sampling + Conditional GAN)



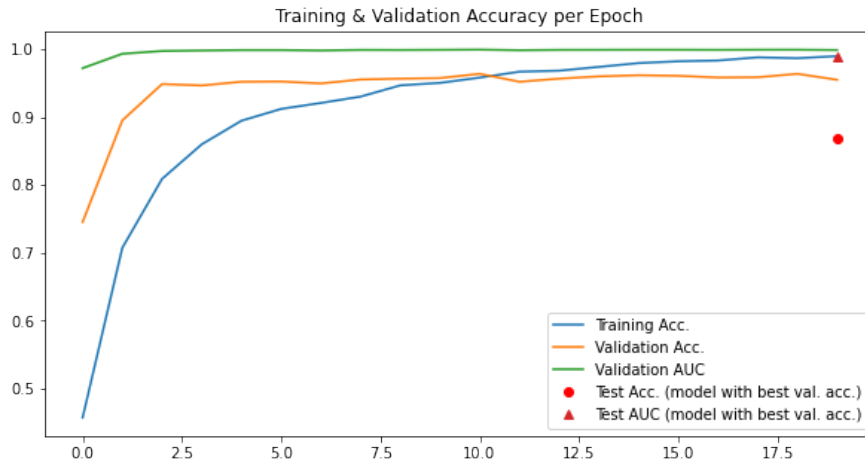
**Fig. 39.** Confusion Matrix for the best OrganaMNIST model (Weighted Random Sampling + Conditional VAE)



**Fig. 40.** Accuracy and AUC per epoch for the best OrganaMNIST model (Weighted Random Sampling + Conditional VAE)



**Fig. 41.** Confusion Matrix for the best OrgancMNIST model (Weighted Random Sampling + Conditional GAN)



**Fig. 42.** Accuracy and AUC per epoch for the best OrgancMNIST model (Weighted Random Sampling + Conditional GAN)



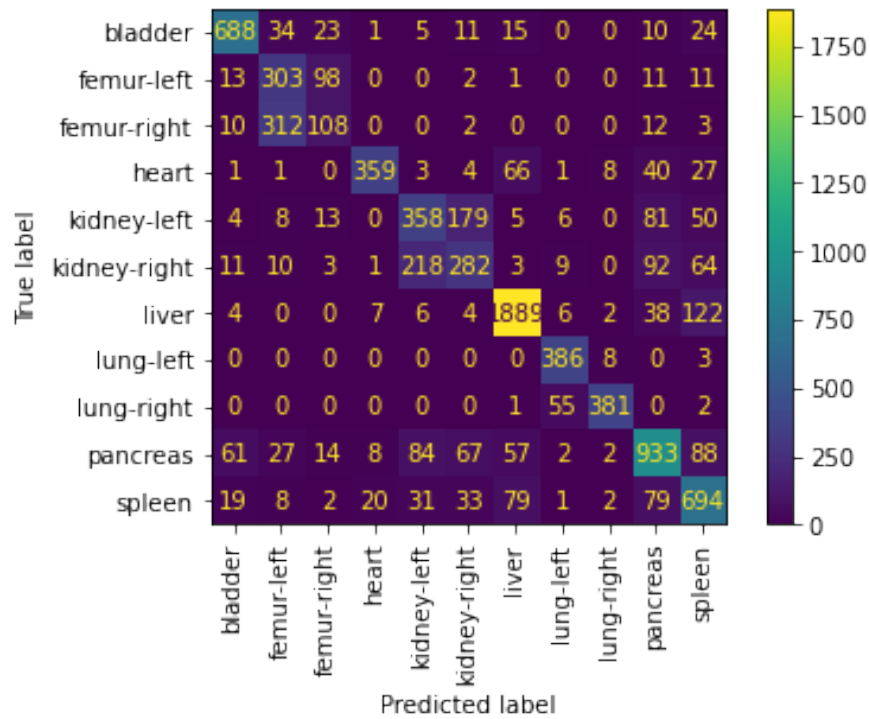


Fig. 43. Confusion Matrix for the best OrgansMNIST model (Conditional GAN)

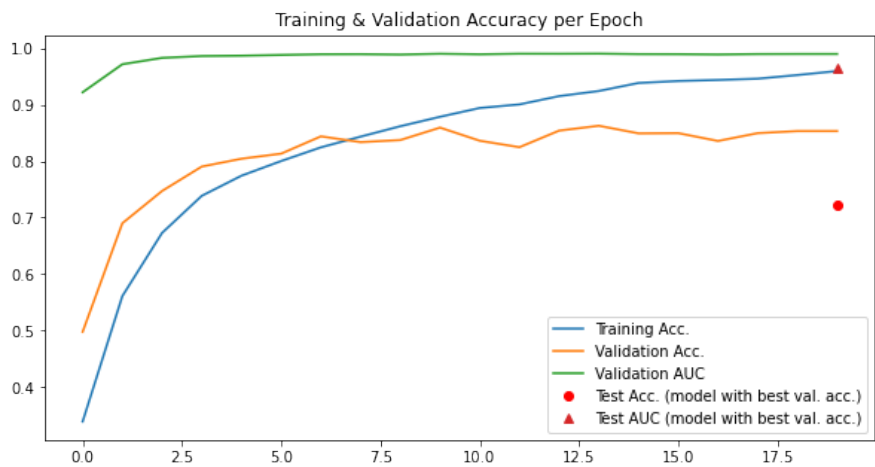


Fig. 44. Accuracy and AUC per epoch for the best OrgansMNIST model (Conditional GAN)