

## Exercise 1

**Deadline: 6.11.2024 16:00.**

Ask questions in discord to `#ask-your-tutor`

In this exercise, you will implement and test classical generative modelling methods.

## Regulations

Please implement your solutions in form of *Jupyter notebooks* (\*.ipynb files), which can mix executable code, figures and text in a single file. They can be created, edited, and executed in the web browser, in the stand-alone app JupyterLab, or in the cloud via Google Colab and similar services.

Create a Jupyter notebook `generative-baselines.ipynb` for your solution and export the notebook to HTML as `generative-baselines.html`. Zip all files into a single archive `ex01.zip` and upload this file to MaMPF before the given deadline.

Moreover, please set your **Anzeigename/display name** and **Name in Uebungsgruppen/name in tutorials** in MaMPF to your real name, which should be identical to your name in `muesli` and make sure you **join the submission** of your team via the invitation code before the submission deadline. Check out <https://mampf.blog/handing-in-homework-assignments> for instructions. Also note that joining a submission takes a while when you do it for the first time (later it will be just a single click), so don't wait until the last minute before the deadline.

## 1 Two-dimensional data

Use the function `sklearn.datasets.make_moons()` to create 2-dimensional training data sets of varying sizes. Implement and train the following models (do not use pre-defined models and training algorithms from sklearn!) :

1. a two-dimensional histogram
2. a single Gaussian
3. a Gaussian mixture model (GMM)
4. a kernel density estimator (KDE) with squared exponential kernel

Implement the maximum mean discrepancy (MMD) metric with squared exponential and inverse multi-quadratic kernels for evaluation. Evaluate the accuracy of your models by calculating the MMD between a test dataset from `make_moons()` and the data generated by each model. Visualize the accuracies as a function of model hyperparameters (histogram: bin size, GMM: number of components, KDE: kernel bandwidth) and training set size. Comment on your findings.

For a number of representative models (both good and bad ones), create two 2D plots that (i) visualize the numerical values of the learned density, and (ii) visualize a generated dataset from the model. Comment on model strengths and weaknesses. Bonus: Add some representation of the model solution to your plots (e.g. the grid of the histogram, some selected mixture components of the GMM).

## 2 Higher-dimensional data

Repeat the same tasks with the digits dataset (`sklearn.datasets.load_digits()`). Use the models and algorithms from sklearn this time. You may consider [sklearn's KDtrees](#) for speeding up computations in GMMs and KDE. Replace histograms (which do not scale to higher dimensions) with density

forests, e.g. using the code from <https://pypi.org/project/quantile-forest/> or [https://github.com/kfritsch/density\\_forest](https://github.com/kfritsch/density_forest).

Again, check model accuracy by MMD and visualize generated data for some representative models (you do not need to visualize the numerical density values – this is hard in 64 dimensions). In addition, train a `sklearn.ensemble.RandomForestClassifier` on the original dataset to distinguish the 10 digit classes. Use this classifier to check for the models which create recognisable output if the 10 digits are generated in equal proportions.