

While waiting...

1. Try to download data and codes:

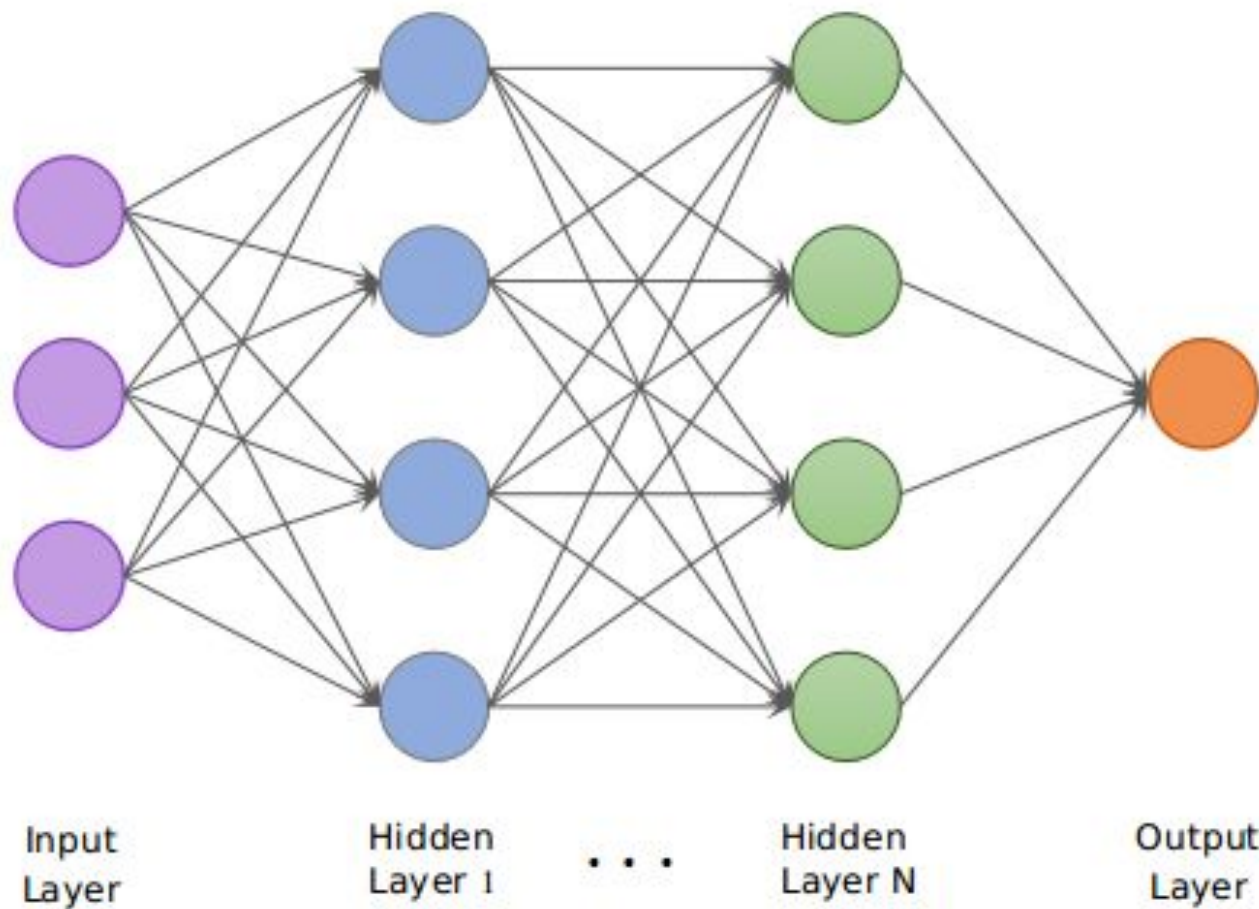
- GitHub repository:

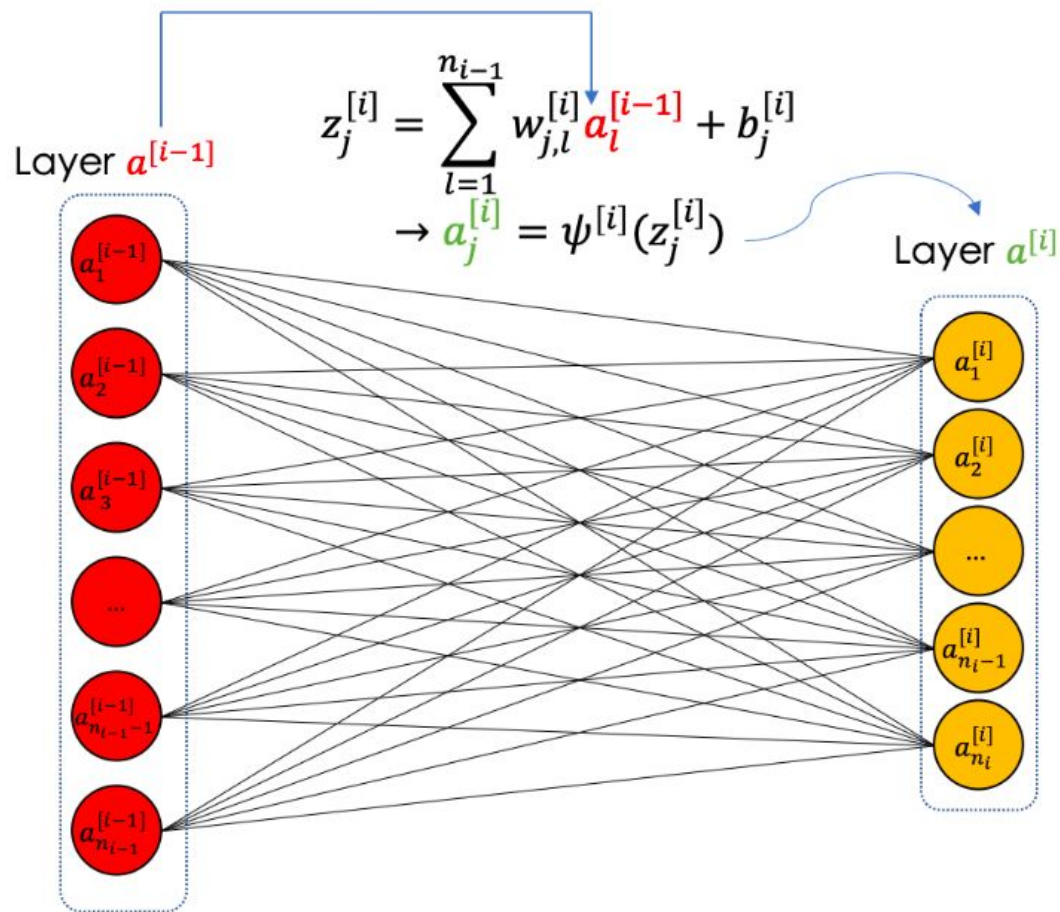
<https://github.com/OBIWOW/OBiWoW-2025/tree/main/08-Monday/deep-learning-for-drug-target-interaction-prediction-and-drug-repurposing>

2. Download conda if you do not have it on your computer

3. Setup your environment (install necessary libraries)

- We will run **Jupyter notebook**. We will need basic bioinformatics libraries such as **pandas**, **numpy**, **scikit-learn**, **matplotlib** and deep learning framework **PyTorch**, also **requests**, **pickle** (or **pickle5**)

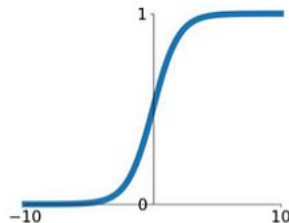




Activation Functions

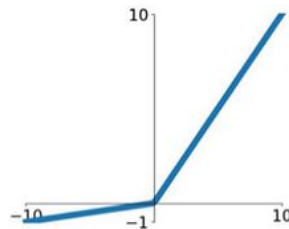
Sigmoid

$$\sigma(x) = \frac{1}{1+e^{-x}}$$



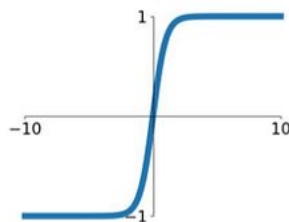
Leaky ReLU

$$\max(0.1x, x)$$



tanh

$$\tanh(x)$$

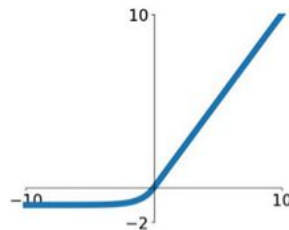


Maxout

$$\max(w_1^T x + b_1, w_2^T x + b_2)$$

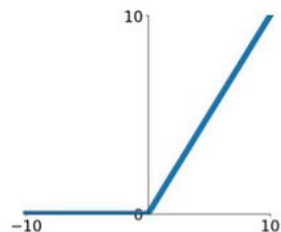
ELU

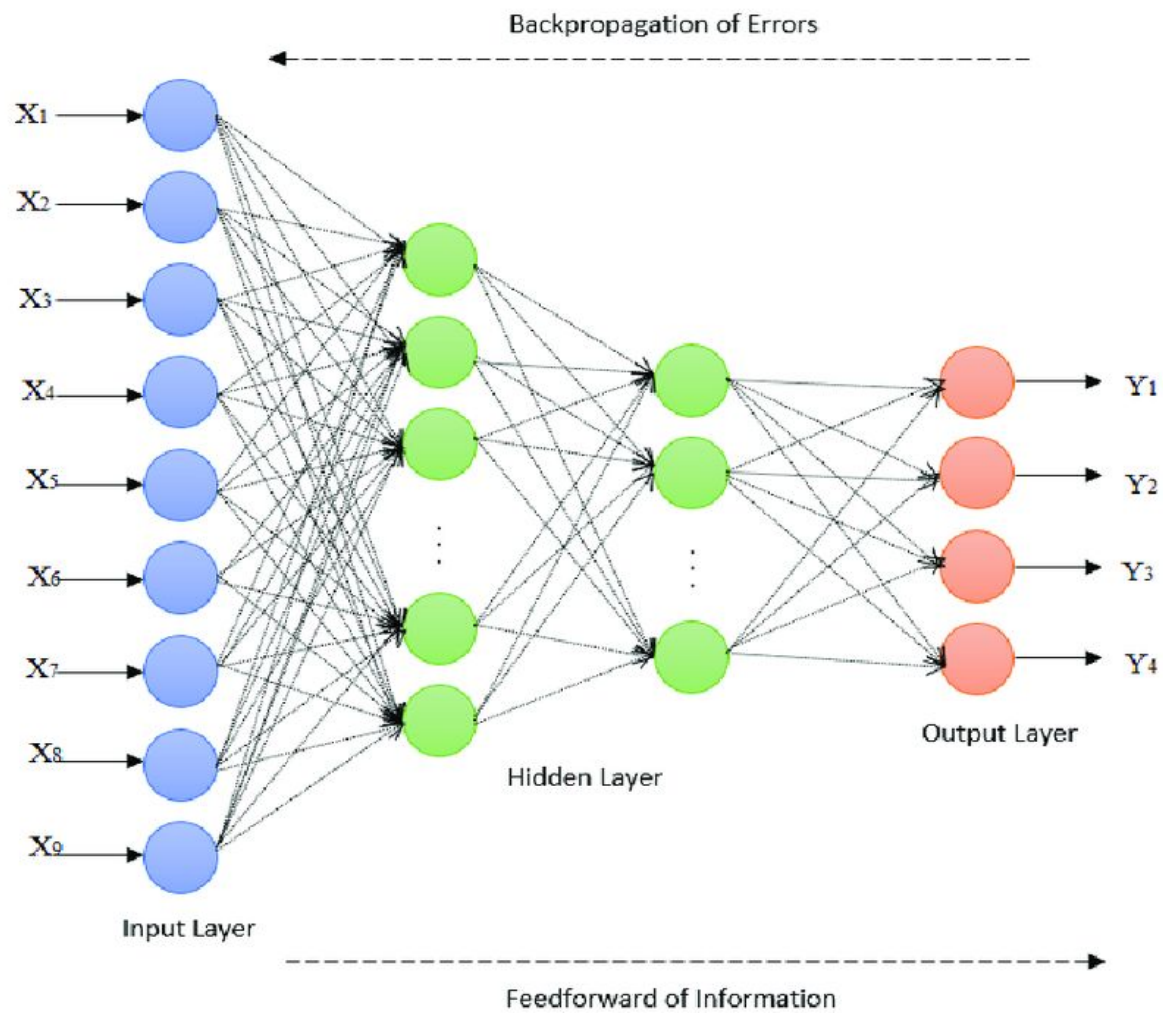
$$\begin{cases} x & x \geq 0 \\ \alpha(e^x - 1) & x < 0 \end{cases}$$



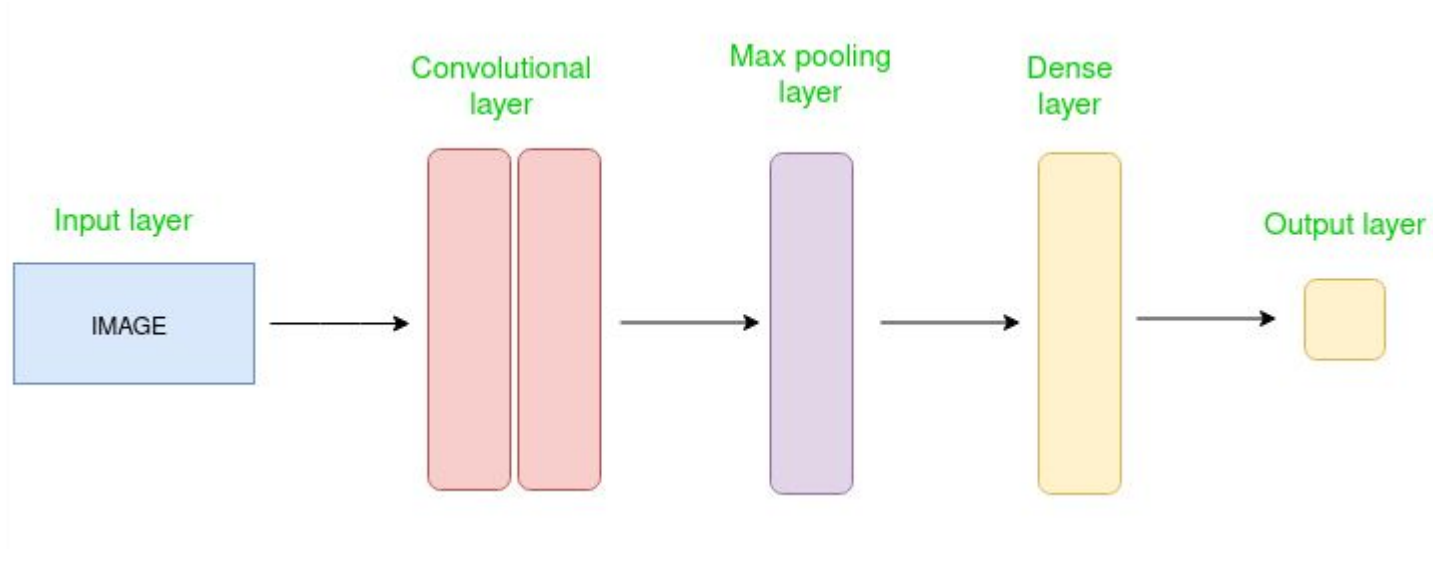
ReLU

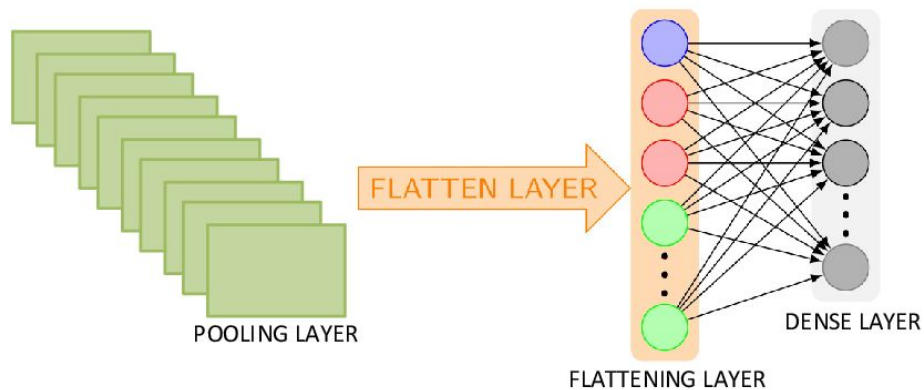
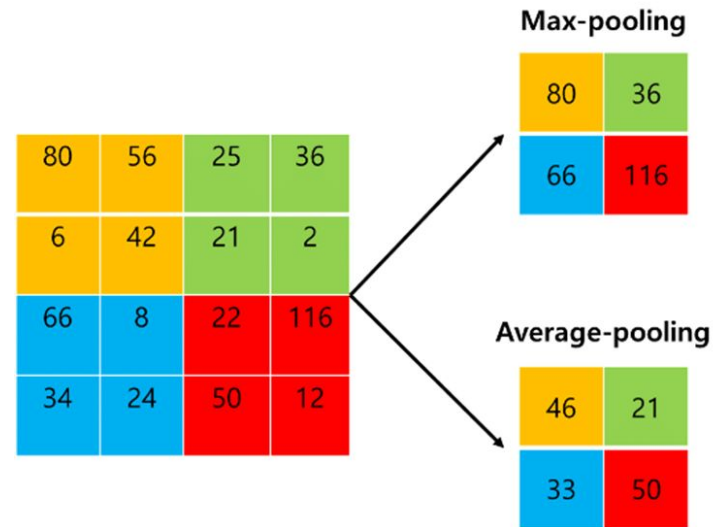
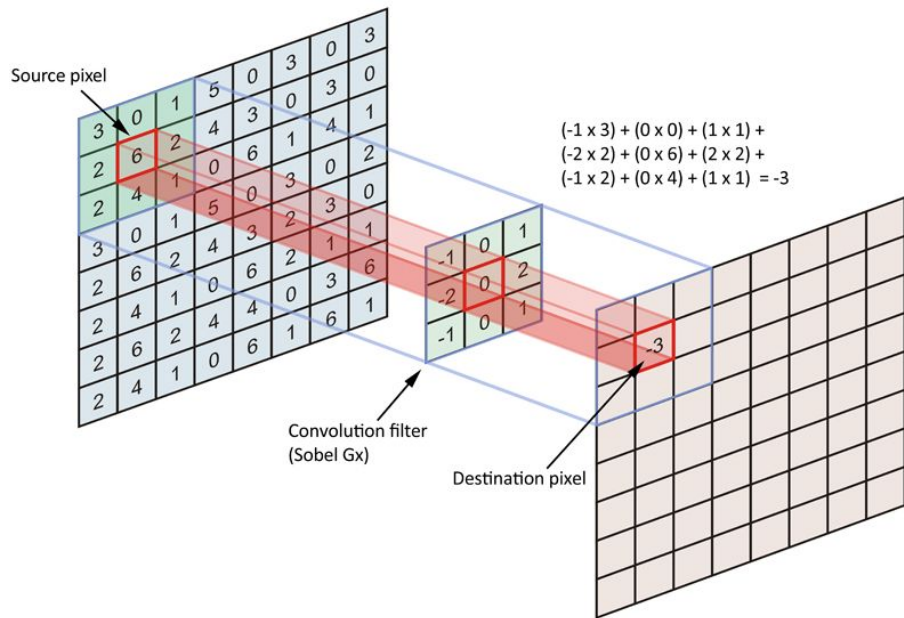
$$\max(0, x)$$

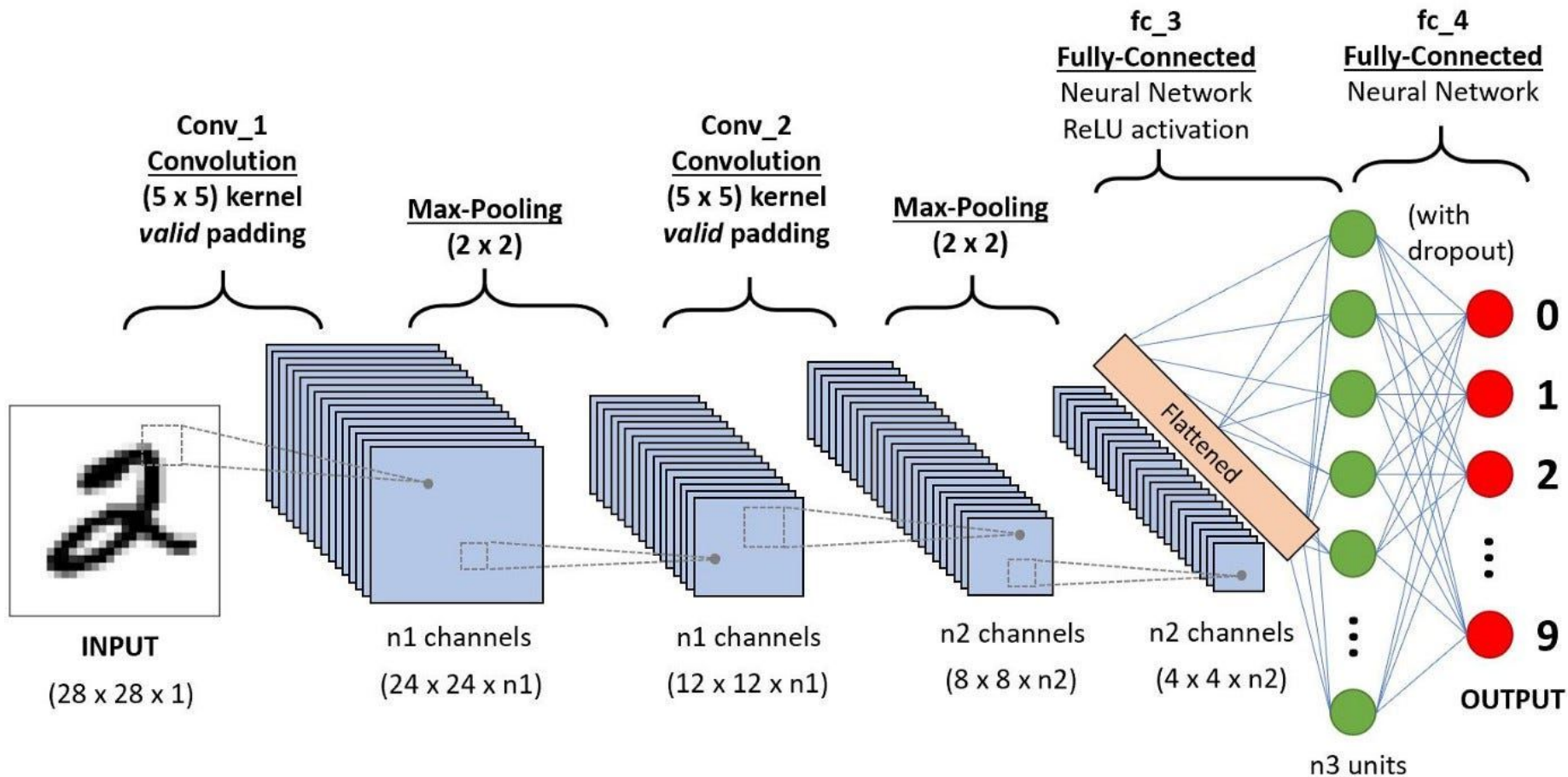




Convolutional Neural Networks







Evaluation
form

