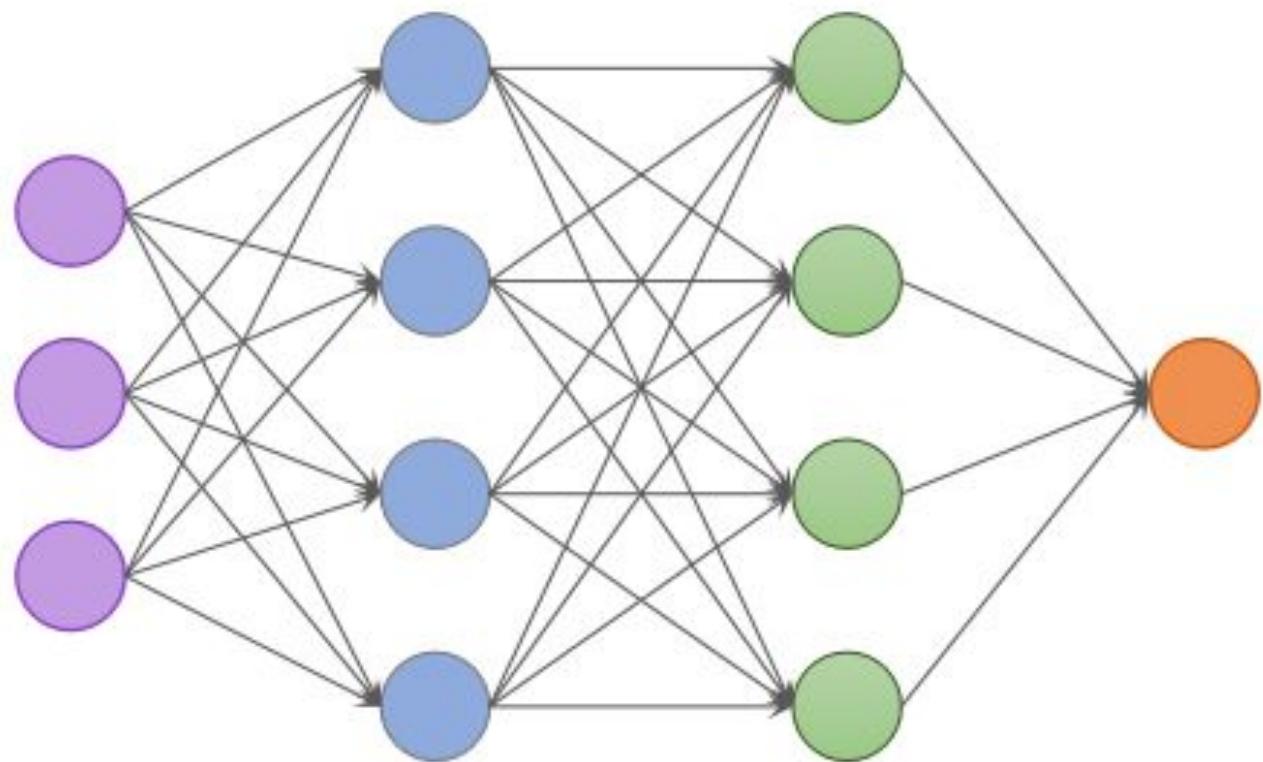


While waiting...

- 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**)



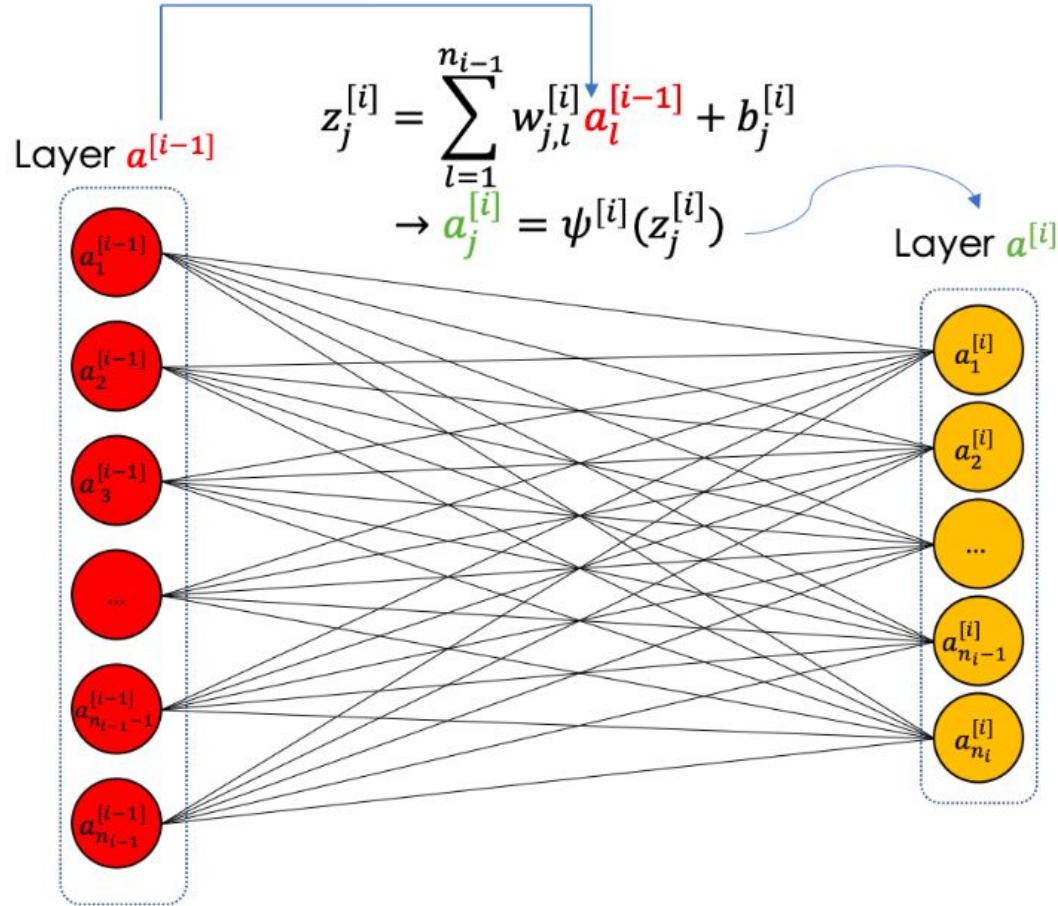
Input
Layer

Hidden
Layer 1

...

Hidden
Layer N

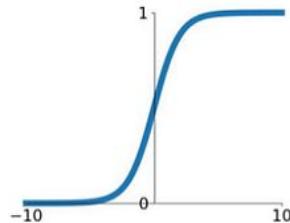
Output
Layer



Activation Functions

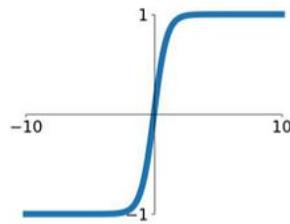
Sigmoid

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



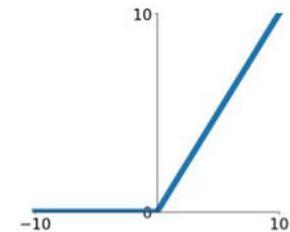
tanh

$$\tanh(x)$$



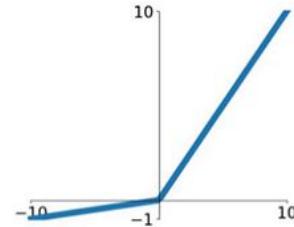
ReLU

$$\max(0, x)$$



Leaky ReLU

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

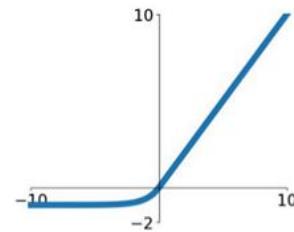


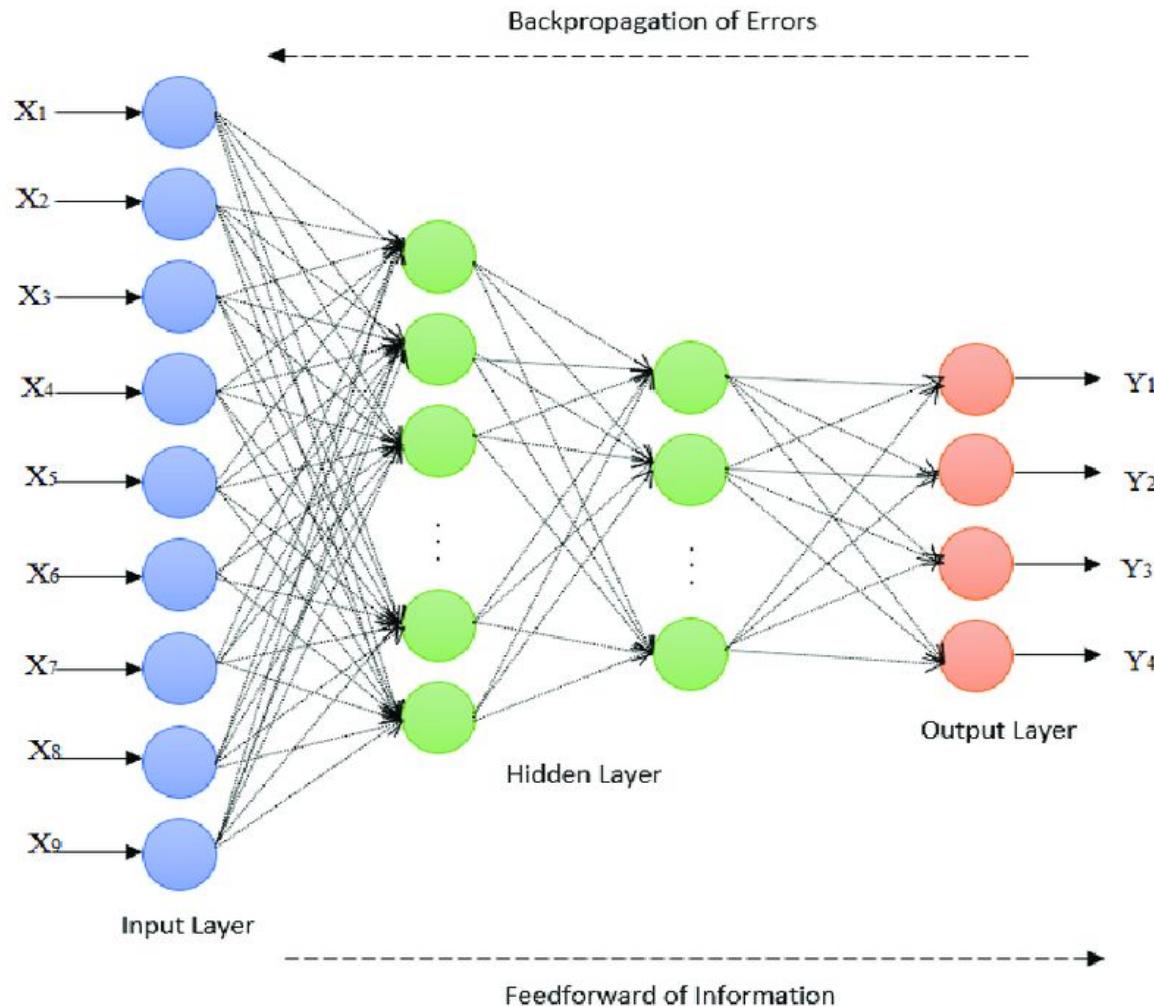
Maxout

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

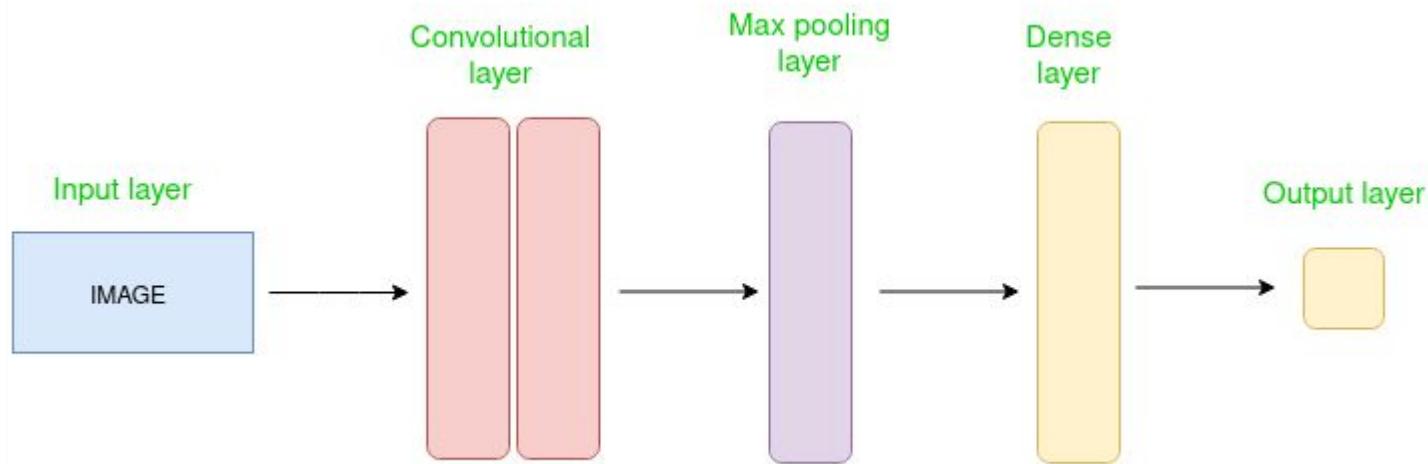
ELU

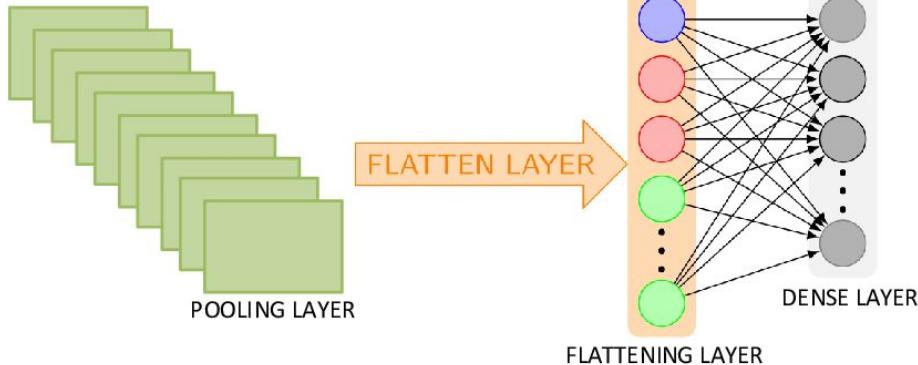
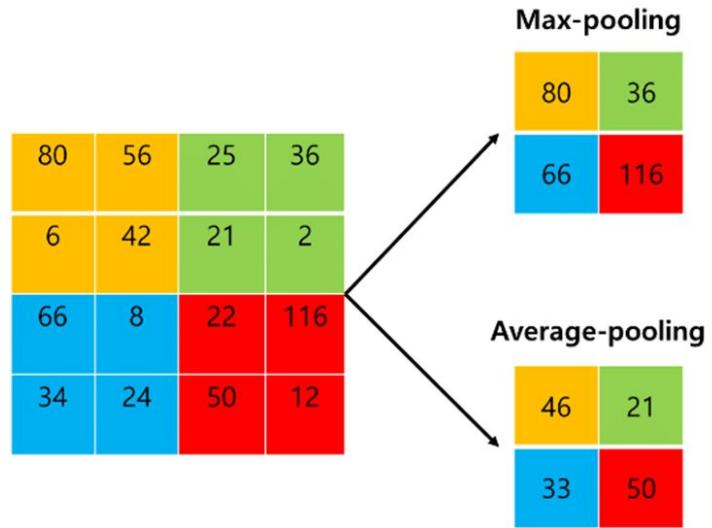
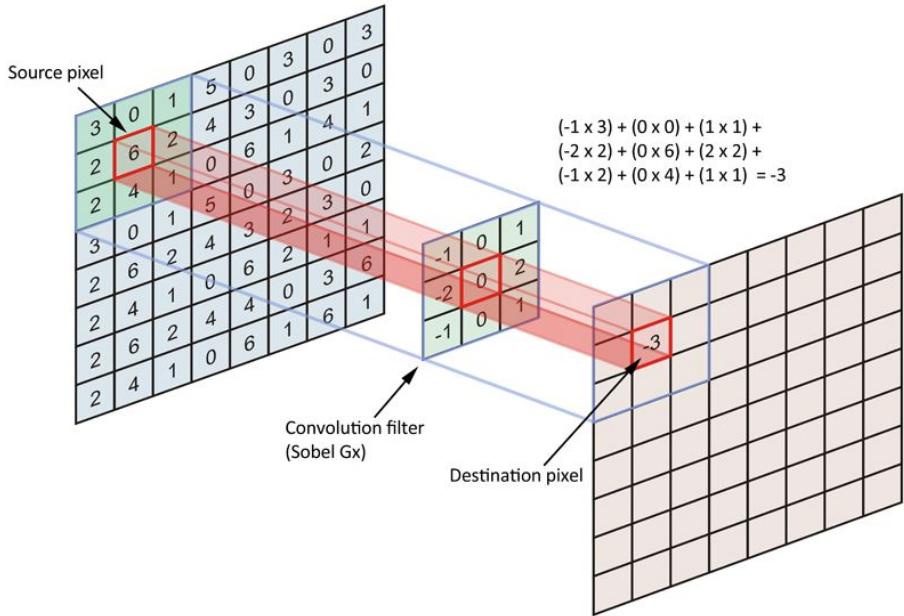
$$\begin{cases} x & x \geq 0 \\ \alpha(e^x - 1) & x < 0 \end{cases}$$

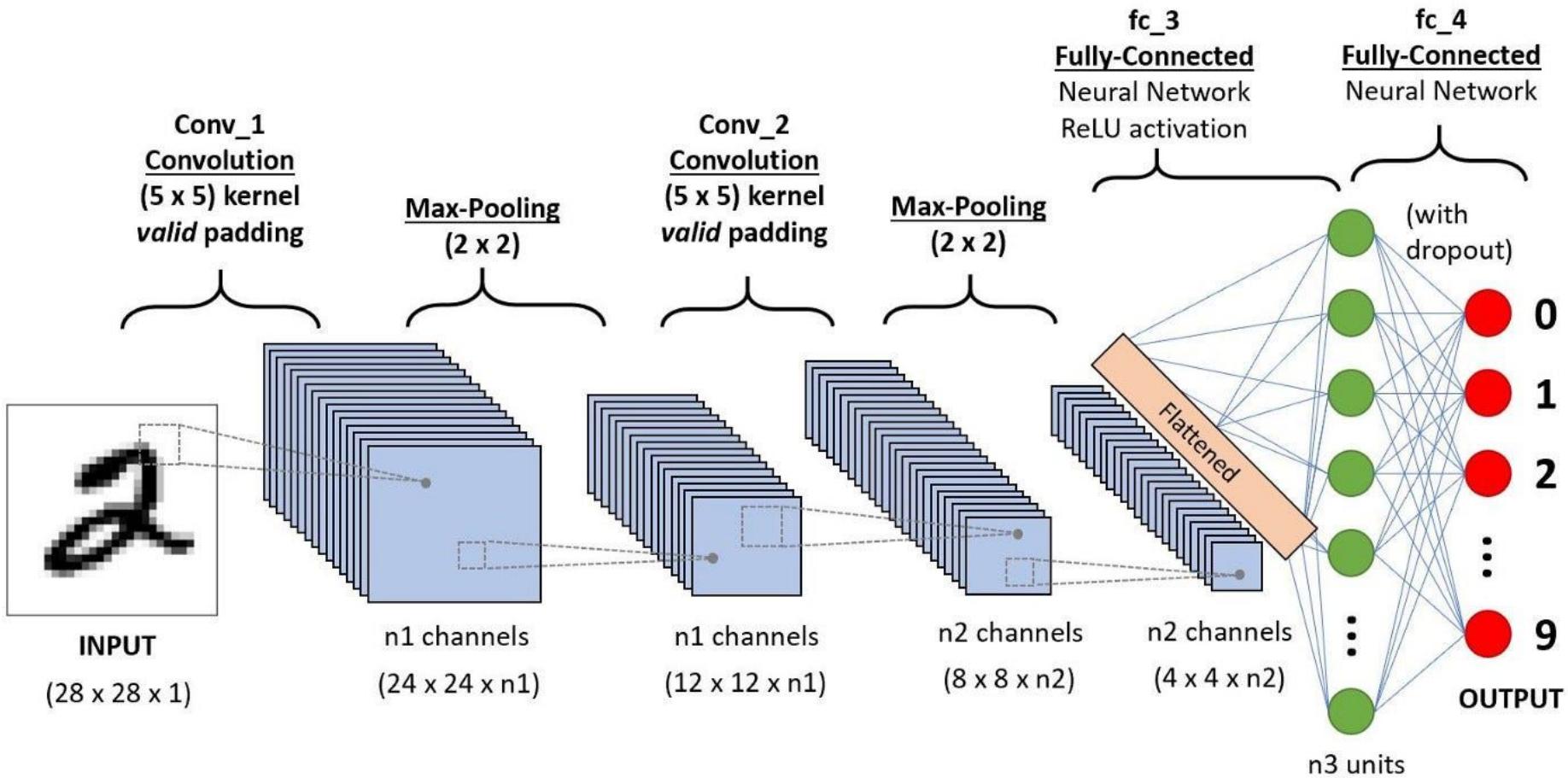




Convolutional Neural Networks







Evaluation
form

