

# Tools & Simulation Structure ( Before Coding)

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## 1. Why NumPy is used ( fast math )

Python alone is slow for:

- arrays
- loops
- math on thousands of time steps
- large vectors

Numpy lets you do:

- ✓ fast vector operations
- ✓ handle voltage arrays
- ✓ handle spike arrays
- ✓ simulate in milliseconds
- ✓ run thousands of dt steps instantly

Without Numpy, our simulation would be painfully slow

## 2. Why Matplotlib is used ( graph plotting )

Neuroscience = LOTS of graphs

We need:

- Voltage vs time graphs
- Spike raster plots
- Emotional patterns
- Firing rate
- Feature visualizations

Matplotlib makes these super easy, clean and scientific looking

## 3. Simulations Loop Logic ( concept )

Every simulation has this loop:

```
for t in range(total_steps):  
    I = input_functions(t)  
    V = update_voltage(V, I)  
    if V crosses threshold:  
        record spike  
        reset voltage  
        store results
```

This loop runs every dt

This is the entire heart of our simulations

## 4. Data Structure to Store Voltage & Spikes

Voltage (V)

Voltage = np.zeros( total\_steps )

Each index = voltage at a time step

Spikes

Spikes = np.zeros( total\_steps )

when neuron fires:

spikes[t] = 1

This becomes spike train dataset

## 5. Why Jupyter Notebooks are Useful

✓ We can see graphs instantly

✓ We can run cell-by-cell

✓ We can keep notes + diagrams next to code

✓ We can debug more easily

## 6. Understanding Basic Project folder Layout

Now the simulation files will be organized:

neuron\_models /

    lif-single-neuron.ipynb

    lif-single-neuron.py

    multi-neuron-circuit.ipynb

emotional-simulation /

    spike-trains /

    spike-plots /

## \* Summary

- NumPy → fast math & arrays
- Matplotlib → graphs
- Simulation loop = core of neuron modeling
- Voltage stored in arrays
- Spikes stored in binary arrays
- Jupyter = perfect for scientific simulations
- Folder layout keeps project clean.