

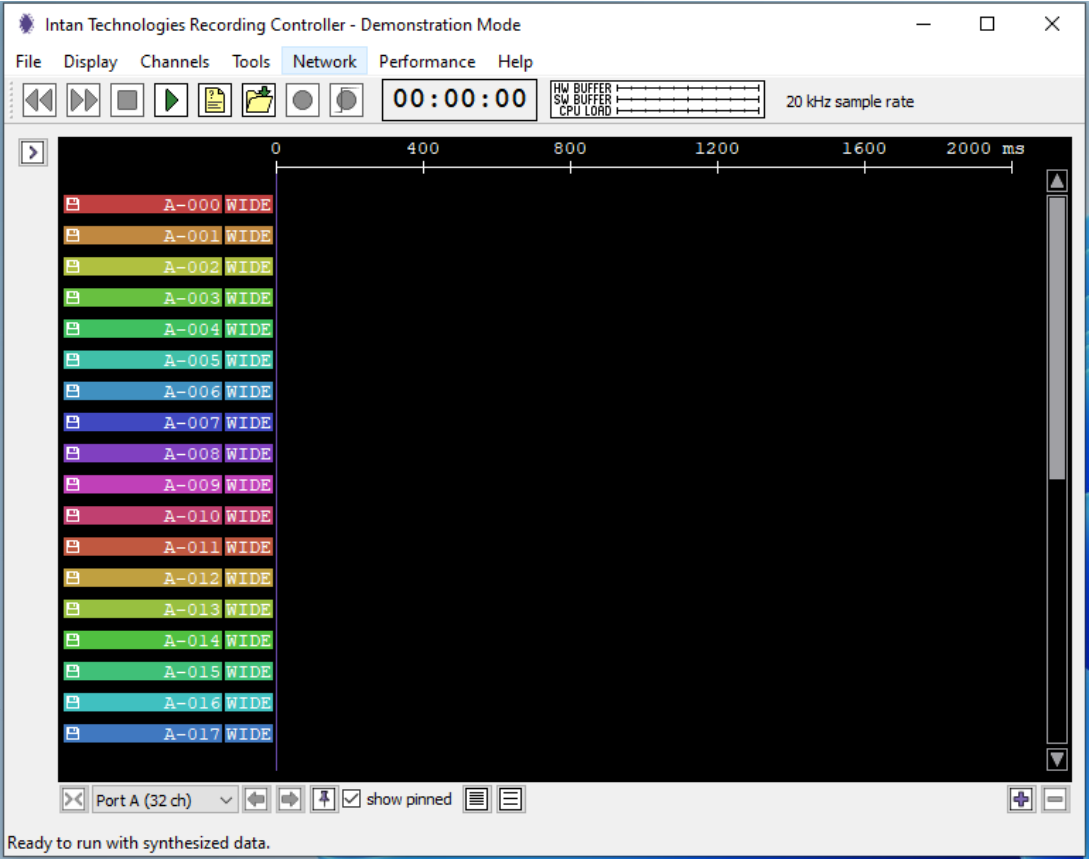
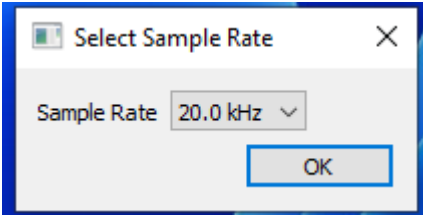
# User Manual to run Plotter.py



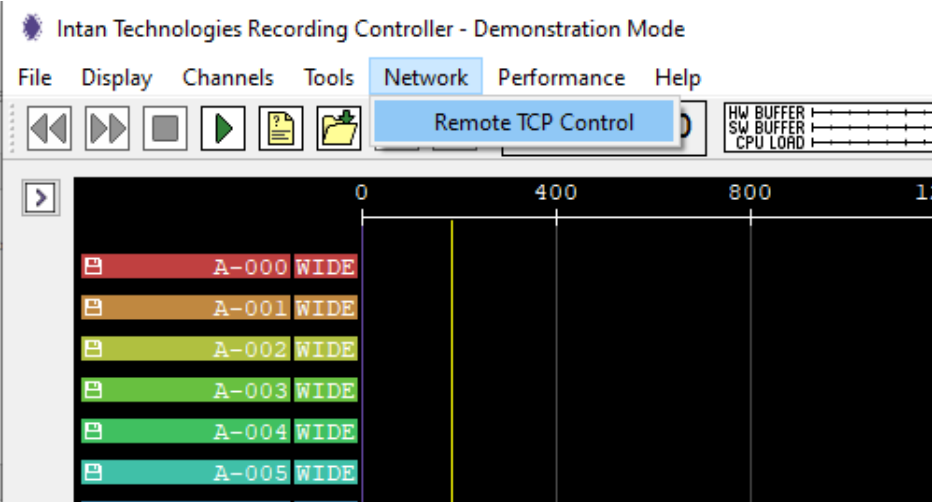
**STEP 1**



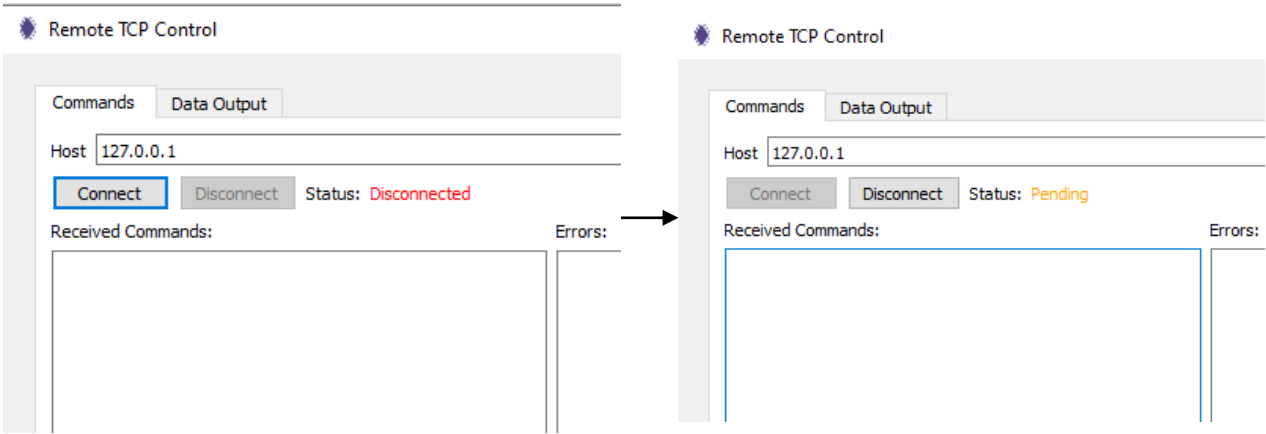
**STEP 2**



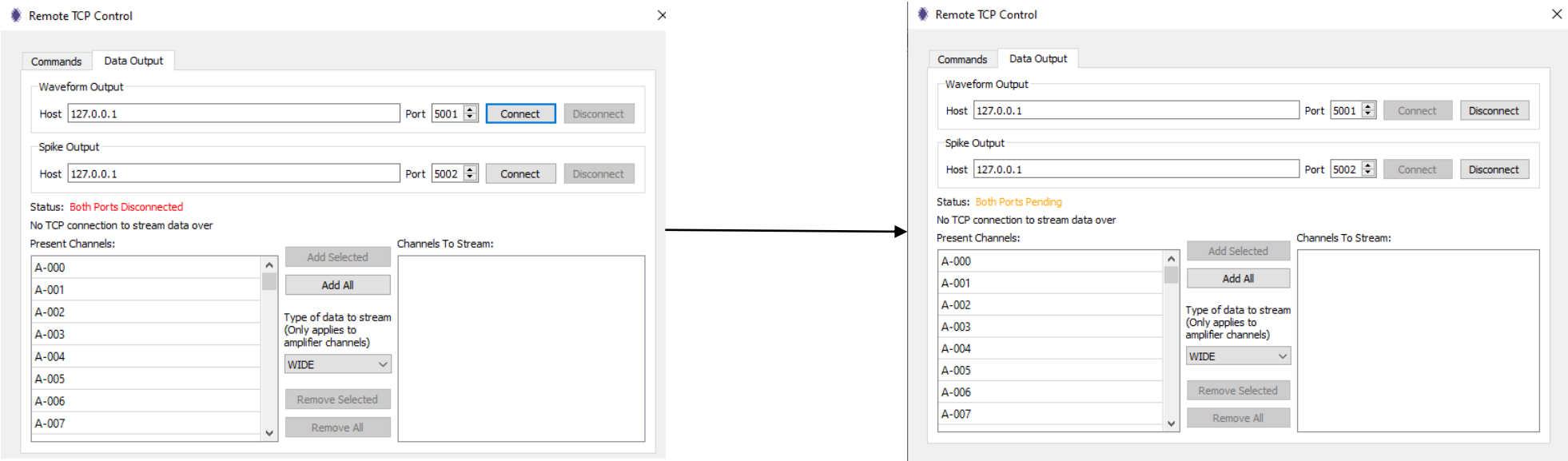
STEP 3



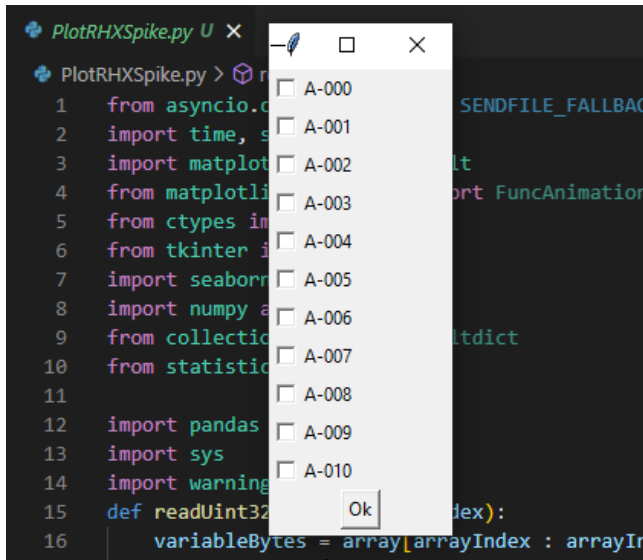
STEP 4



STEP 5



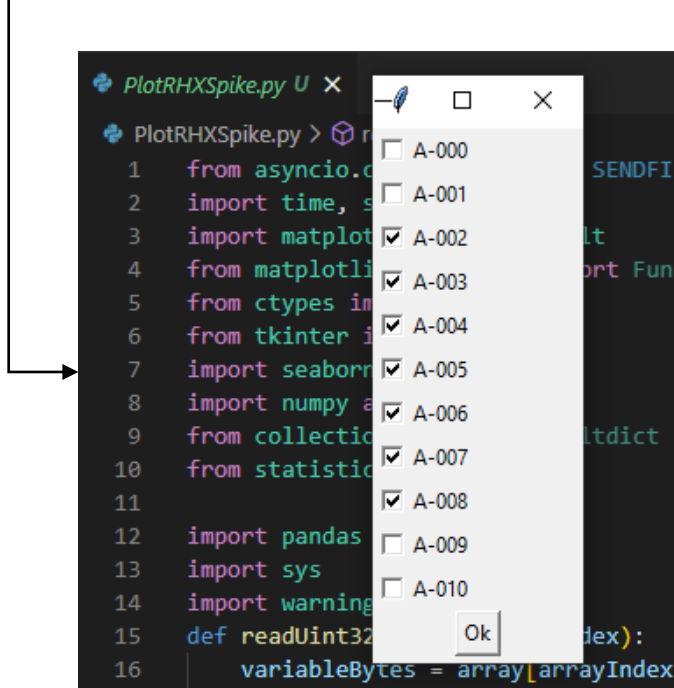
## STEP 6



## STEP 7

Once 'Ok' clicked , you will see

- two empty matplotlib plots (figure 1 & figure 2) open
- Plotter.py waiting for response from `Texture_task_wTCPIP.m`



```
H:\realtime-analysis>C:/Users/sammasi/AppData/Local/miniconda3/envs/RTA/python.exe h:/realtime-analysis/PlotRHXSpike.py
selected channels ['A-001', 'A-002', 'A-003', 'A-004', 'A-005', 'A-007', 'A-007', 'A-008', 'A-007']
opening plot.....
Connecting to TCP command server...
Connecting to TCP SPIKE output server...
```

## STEP 8

Once the connection is established ,  
plotter.py will be ready to receive  
stim\_cond & SPK\_outut ON OFF signal

```
H:\realtime-analysis>C:/Users/sammasi/AppData/Local/miniconda3/envs/RTA/python.exe h:/realtime-analys
selected channels ['A-002', 'A-003', 'A-004', 'A-005', 'A-006']
opening plot.....
Connecting to TCP command server...
Connecting to TCP SPIKE output server...
connected from ('172.27.85.105', 39080) to receive stim conditions
connected to receive t time to collect SPK
stim condition recieved :
```

When the texture\_taskwTCPIP.m is  
terminated, the plotter.py terminal  
would look something like this

```
stop SPK output
stim condition recieved : 2
start SPK output
.
.
.
.
.
.
.
.
stop SPK output
stim condition recieved :
Enter 'q' to quit: 
```

And to quit plotter.py, enter 'q'

```
.
.
stop SPK output
stim condition recieved :
Enter 'q' to quit: q
H:\realtime-analysis>
```

The spike data from the trials will be  
stored in “CH\_stim\_SPK\_data.csv”

To run the code again ,follow steps 3-7