

# Assignment 4 README

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**NOTE:** Running ns3 simulation requires **python2**, while the analysis scripts use **python3**. Please ensure both are available on the system.

## Setup of ns3

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Download ns3 3.28.1 : <https://www.nsnam.org/release/ns-allinone-3.28.1.tar.bz2>

Build ns3 by following these instructions <https://www.nsnam.org/docs/release/3.28/tutorial/single.html/index.html#building-ns3>

In the `ns-3.28.1` directory in the file `waf`, change the first line to `#!/usr/bin/env python2` to ensure it is run using python2.

## Compiling simulation.cc

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Modify `simulation.cc` by changing line 37 to "0","256","512","1000" to run experiment for different RTS Thresholds

Copy the `simulation.cc` file into `ns-allinone-3.28.1/ns-3.28.1/scratch/`

Now in the `ns-3.28.1` directory run the following commands :

```
./waf --run scratch/simulation
```

## Analysis

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Now there will be 4 files produced in this directory itself :

1. Westwood\_256.xml (for threshold=256 we have named this .xml see line 206 )
2. AccessPoint-1-0.pcap
3. Station-0-0.pcap
4. Station-2-0.pcap

## Analysing .xml

All relevant files are in `xmls` folder. or you can find them by below instructions :

Copy the following file `ns-3.28.1/src/flow-monitor/examples/flowmon-parse-results.py` and .xml file produced above, into a common directory

Run

```
python2 flowmon-parse-results.py Westwood_256.xml
```

You can see the output on terminal now

**Note :** Please ensure both python2 and python3 are installed on your system. The ns3 core uses python2 whilst the packet analysis is done using python3.

# Analysing .pcap

Scripts are in `scripts/data` folder in submission

in that folder make a folder named exactly `assignment-4-data` and 4 subfolders named :  
"0", "256", "512", "1000". Copy the pcaps in each subfolder. [Or see note below]

The scripts will be run using python3. We will also need to install `dpkt` package.

```
pip install dpkt
```

Now to run the scripts:

```
python3 gen_ack.py  
python3 gen_cts.py  
python3 gen_rts.py  
python3 gen_tcp_ack_seg.py
```

Now they will produce .txt files which will have labelled data.

Paste the text file data into graph scripts located in `scripts/graph` folder

```
python3 graph_ack.py  
python3 graph_cts.py  
python3 graph_rts.py  
python3 graph_tcp_ack.py  
python3 graph_tcp_seg.py
```

Hence we get output of graphs, you can export them in any image or pdf format.

We have done the above process and saved the txt files and graphs in `/data` and `/graphs` folder in our submission.

**Note :** The pcap files are 1.2 GB in size per experiment so they are uploaded here : [https://iitgoffice-my.sharepoint.com/personal/atrivedi\\_iitg\\_ac\\_in/\\_layouts/15/onedrive.aspx?id=%2Fpersonal%2Fatrivedi%5Fiitg%5Fac%5Fin%2FDocuments%2Fcs342%2Fassign4%2Fdata](https://iitgoffice-my.sharepoint.com/personal/atrivedi_iitg_ac_in/_layouts/15/onedrive.aspx?id=%2Fpersonal%2Fatrivedi%5Fiitg%5Fac%5Fin%2FDocuments%2Fcs342%2Fassign4%2Fdata)

All the outputs and graphs are mentioned in report as well.