

# Distributed Algorithms (UAI/503): Berkeley Algorithm

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## Abstract

In this report, the Berkeley Algorithm and its implementation using Java RMI technology is explained. The source file can be viewed under Github.

## 1 Introduction

In the Distributed Systems (DS) the nodes are communicating with each other using message passing. To achieve real time applications working in ordered manner the reference parameter used is time. Therefore synchronisation of time is essential for allocating the available resources. Synchronization in DS can be achieved by using physical clock of the node. For synchronization purpose, each node in the system needs to share their local clock time with another node in the system.

## 2 Berkeley Algorithm

The Berkeley Algorithm follows a master slave communication operation. There is a central computer that serves as the master or a time server. The master periodically sends a request message to all other slaves or nodes which asks the time of the destination nodes. The master will receive the round trip time value (RTT) from the slaves, and the master will average the time values including its own clock value and readjusts its own clock accordingly. The master will also eliminate readings from faulty clocks that is values far outside the range. It does this by taking a subset of the returned times with a small variance. After this, the master will then send the amount by which each individual's clock requires adjustment to each individual clock. This value can be positive or negative. If the master fails at any point, then a new master is elected to take over and function exactly like its predecessor.

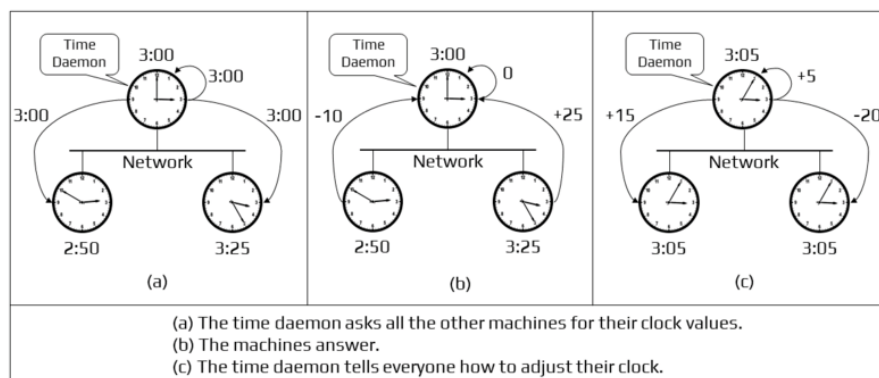


Figure 1: Berkeley Algorithm Workflow[source: see below]

### 3 Methods

- *client.py*: serial script
- *server.py*: File reading option 1 parallel version C script

### 4 Results

To test the script run *server.py* on the machine that will receive the connections and *client.py* on the machine that will send the connections.

Libraries used in the script:

- *time* - time library
- *socket* - library for making connections to other hosts

```
(c) 2019 Microsoft Corporation. All rights reserved.
D:\USB_SEMESTER\503-DistributedAlgorithms\semester_project\503-DistributedAlgorithms\berkeley_algorithm\src>java server/slaveOne
Slave 01 started at port: 1501
Updated Time is: 12:03:23

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D:\USB_SEMESTER\503-DistributedAlgorithms\semester_project\503-DistributedAlgorithms\berkeley_algorithm\src>java server/slaveTwo
Slave 02 started at port: 1502
Updated Time is: 12:03:23

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D:\USB_SEMESTER\503-DistributedAlgorithms\semester_project\503-DistributedAlgorithms\berkeley_algorithm\src>java server/slaveThree
Slave 03 started at port: 1503
Updated Time is: 12:03:23

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D:\USB_SEMESTER\503-DistributedAlgorithms\semester_project\503-DistributedAlgorithms\berkeley_algorithm\src>java server/slaveFour
Slave 04 started at port: 1504
Updated Time is: 12:03:23

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D:\USB_SEMESTER\503-DistributedAlgorithms\semester_project\503-DistributedAlgorithms\berkeley_algorithm\src>java client/master
Local time: 12:00:00
##### Berkeley Algorithm #####
##### Fetched the Master Clock #####
Connected successfully to: SLAVE 01
time at slave 01 is: 12:01:10
Connected successfully to: SLAVE 02
time at slave 02 is: 12:02:50
Connected successfully to: SLAVE 03
time at slave 03 is: 12:03:50
Connected successfully to: SLAVE 04
time at slave 04 is: 12:04:00
Connected successfully to: SLAVE 05
time at slave 05 is: 12:05:00
Average Delta Computed is: 203 seconds
Successfully Updated the Clock...
Master/Local Time: 12:03:23
new time at slave 01 is: 12:03:23
new time at slave 02 is: 12:03:23
new time at slave 03 is: 12:03:23
new time at slave 04 is: 12:03:23
new time at slave 05 is: 12:03:23
```

Figure 2: Updating 5 Slaves and a master clock time using Berkeley Algorithm

[Link to Github](#)

## A Serial Code

```
import os
import io
os.chdir("C:/jithis_drive/zhaw/ACLS/Genomics_track2/miniJATI/batch_J/params")

#f = io.open('file.txt', 'w', newline='\n')

files = []
for i in range(1,601):
    files.append("params_" + str(i))

for i in range(0,6):
    for j in range(0,100):
        x = (i*100) + (j)
        f = io.open(files[x], "w", newline='\n')
        f.write("alphabet=DNA\nalignment=true\n")
        f.write("init.tree=user\ninput.sequence.sites_to_use=all\n")
        if i == 0 or i == 3:
            f.write("model=PIP(model=K80(kappa=2.0),lambda=100.0,mu=0.1)\n")
        if i == 1 or i == 4:
            f.write("model=PIP(model=K80(kappa=2.0),lambda=316.22776601683796,"
                    + "mu=0.31622776601683794)\n")
        if i == 2 or i == 5:
            f.write("model=PIP(model=K80(kappa=2.0),lambda=447.21359549995793"
                    + ",mu=0.4472135954999579)\n")
        f.write("rate_distribution=Constant\n")
        f.write("optimization=ND-BFGS(derivatives=BFGS)\n")
        f.write("optimization.ignore_parameters=BrLen,Model\n")
        f.write("optimization.reparametrization=false\n")
        f.write("optimization.alignment=false\n")
        f.write("optimization.topology=false\n")
        f.write("input.sequence.file=/scratch/IAS/AnisGroup/peechjit/execs/"+str(i)+
                ".exec/output/sim-"+str(j)+"_sequence.txt\n")
        if i==0 or i==1 or i==2:
            f.write("input.tree.file=/scratch/IAS/AnisGroup/common/data/trees/"
                    + "ultrameric_tree_b0.01.nwk\n")
        else:
            f.write("input.tree.file=/scratch/IAS/AnisGroup/common/data/trees/"
                    + "unb_ultrameric_tree_b0.01.nwk\n")

        f.write("output.msa.file=/scratch/IAS/AnisGroup/peechjit/mJATI_output/"
                + "msa_out"+str(i)+"_"+str(j)+".fa\n")
        f.write("output.tree.file=/scratch/IAS/AnisGroup/peechjit/mJATI_output/"
                + "trees/tree_out"+str(i)+"_"+str(j)+".nwk\n")
        f.write("output.estimates=/scratch/IAS/AnisGroup/peechjit/mJATI_output/log"
                + str(i)+"_"+str(j)+"\n")
    f.close()
```