For Windows OS and Linux (e.g. Ubuntu,...):

Step 1: Download the Algorithms.rar file from the link and extract it

Step 2:

- + For Windows: Go to the windows start menu and open the **Command Prompt application**.
- + For Linux: Open **Terminal application**

Step 3: Go to the extracted Algorithms folder and type the following command:

java -jar TestAlgorithms.jar

Notes: To run the program for the low values of minsup on datasets (e.g. $minsup \le 0.059\%$ on the BMS dataset,...), you should make sure that the heap memory of the program is at least 12GB.

Below are two examples to show how to run the program:

1) The first example of running the program on **OS Ubuntu 22** as follows:

```
+
                                         hai@hai-virtual-machine: ~/Downloads/Algorithms
                                                                                                  Q
                                                                                                            _ _
                                            ɪ Ş java -jar TestAlgorithms.jar
Are you running OS Windows or Linux (e.g. Ubuntu,...) {1: Windows; 2: Linux}?
    -----Datasets to test------
1: RunningExample
2: BMS
3: Kosarak10k
4: D2C7T10N2.5S6I4
5: D5C10T6N5S6I4
Enter the minimum support threshold ([0, 1]):
NOTES: + If you have already generated 50 constraints for the selected dataset when running any algorithm for the first t
ime,
         and you want to ensure that subsequent algorithms produce the identical result set using the same constraints,
         then you should select 'n' for the following option.
Do you want to generate 50 new constraints (y or n)?
-----Generating 50 constraints randomly-----
-----Finish Generating Constraints-----
   -----Algorithms to test-
0: MFS-SubSC (first run FGenCloSM, then partition the FCS and FGS sets into equivalence classes, and then run MFS-SubSC)
1: CM-SPADE-SubSC
2: CM-SPAM-SubSC
3: PrefixSpan-SubSC
4: Baseline
5: FGenCloSM
Enter an algorithm to be tested:
  ------Running FGenCloSM, please wait...-----
                      _FGenCloSM finished_
Runtime of FGenCloSM (seconds):3.701
Freq. Closed Sequence Count:1
Freq. Generator Sequence Count:1
```

2) The second example of running the program on **OS Ubuntu 22** as follows:

```
[+]
                                         hai@hai-virtual-machine: ~/Downloads/Algorithms
                                                                                                        =
                                                                                                                 hms$ java -jar TestAlgorithms.jar
Are you running OS Windows or Linux (e.g. Ubuntu,...) {1: Windows; 2: Linux}?
-----Datasets to test-----
1: RunningExample
2: BMS
3: Kosarak10k
4: D2C7T10N2.5S6I4
5: D5C10T6N5S6I4
Enter the minimum support threshold ([0, 1]):
NOTES: + If you have already generated 50 constraints for the selected dataset when running any algorithm for the first t
ime,
         and you want to ensure that subsequent algorithms produce the identical result set using the same constraints,
         then you should select 'n' for the following option.
Do you want to generate 50 new constraints (y or n)?
------Generating 50 constraints randomly-----
-----Finish Generating Constraints----
------Algorithms to test------
0: MFS-SubSC (first run FGenCloSM, then partition the FCS and FGS sets into equivalence classes, and then run MFS-SubSC)
1: CM-SPADE-SubSC
2: CM-SPAM-SubSC
3: PrefixSpan-SubSC
4: Baseline
5: FGenCloSM
Enter an algorithm to be tested:
------Running FGenCloSM, please wait...----
FGenCloSM finished_____
Runtime of FGenCloSM (seconds):1.987
Freq. Closed Sequence Count:1
Freq. Generator Sequence Count:1
```

```
------The program is partitioning FCS and FGS into equivalence classes, please wait...------
Runtime for partitioning FCS and FGS into equivalence Classes (seconds):57.817
          Finish partitioning
-----MFS-SubSC-----
Constraint sequence = MC: {18427}
Runtime for discovering frequent sequences with this constraint: 0.009 (seconds)
Memory usage for discovering frequent sequences with this constraint: 0.0 (MB)
Number of frequent sequences statisfying this constraint: 0
 -----Testing the constraint sequence 2 per 50 constraints-----
               -----MFS-SubSC-----
Constraint sequence = MC: {18659}
Runtime for discovering frequent sequences with this constraint: 0.001 (seconds)
Memory usage for discovering frequent sequences with this constraint: 0.0 (MB)
Number of frequent sequences statisfying this constraint: 0
--------Testing the constraint sequence 3 per 50 constraints-------
-----MFS-SubSC-----
Constraint sequence = MC: {18635}
Runtime for discovering frequent sequences with this constraint: 0.001 (seconds)
Memory usage for discovering frequent sequences with this constraint: 0.0 (MB)
Number of frequent sequences statisfying this constraint: 0
```

•••••

Testing the co	onstraint sequence 49	per 50 constraints				
Constraint sequence = MC Runtime for discovering f Memory usage for discover Number of frequent sequer	: {12743} frequent sequences with ring frequent sequence	th this constraint: es with this constr)		
Constraint sequence = MC Runtime for discovering Memory usage for discover Number of frequent sequen	· MFS-SubSC : {12603} frequent sequences with ring frequent sequence	th this constraint: es with this constr	0.0 (seconds)			
Statistical inf (a) Dataset B)	formation for 50 const (b) Algorithm			(d) Average Runtim	e (second)	(e) Peak Memory (N
(a) BMS Press any key to finish	(b)FGenCloSM + Parti	tion + MFS-SubSC	(c) 59.865	(d) 1.197	(e) 0.0	