

# UNIVERSIDAD REGIONAL AMAZONICA IKIAM BIOINFORMATICA

FECHA: 09/Mayo/2023

**Nombre: Stalin Daniel Guaigua Puedma** 

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MINGW64:/c/Users/LENOVO/downloads/Stalin/CSB-master/unix/sandbox

LENOVO@StalinPC MINGW64 ~/downloads

LENOVO@StalinPC MINGW64 ~/downloads/Stalin

LENOVO@StalinPC MINGW64 ~/downloads/Stalin

$ cd CSB-master

LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master

$ cd unix

LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix

$ cd sandbox

LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix

$ cd sandbox

LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox

$ cd sandbox
```

```
NINGW64:/c/Users/LENOVO/downloads/Stalin/CSB-master/unix/sandbox
                                                                                    X
_ENOVO@StalinPC MINGW64 ~
$ cd downloads
ENOVO@StalinPC MINGW64 ~/downloads
$ cd Stalin
LENOVO@StalinPC MINGW64 ~/downloads/Stalin
$ cd CSB-master
_ENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master
$ cd unix
LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix
$ cd sandbox
LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox
$ ls -lh ../data/Marra2014_data.fasta
-rw-r--r-- 1 LENOVO 197121 553K May 10 09:59 ../data/Marra2014_data.fasta
LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox
```



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MINGW64:/c/Users/LENOVO/downloads/Stalin/CSB-master/unix/sandbox
                                                                                         X
_ENOVO@StalinPC MINGW64 ~
$ cd downloads
_ENOVO@StalinPC MINGW64 ~/downloads
$ cd Stalin
LENOVO@StalinPC MINGW64 ~/downloads/Stalin
$ cd CSB-master
_ENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master
LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix
$ cd sandbox
LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox
$ ls -lh ../data/Marra2014_data.fasta
-rw-r--r-- 1 LENOVO 197121 553K May 10 09:59 ../data/Marra2014_data.fasta
LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox
$ cp ../data/Marra2014_data.fasta my_file.fasta
LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox
'Papers and reviews'/ my_file.fasta
_ENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox
$
```



```
MINGW64:/c/Users/LENOVO/downloads/Stalin/CSB-master/unix/sandbox
                                                                    X
ENOVO@StalinPC MINGW64 ~
$ cd downloads
_ENOVO@StalinPC MINGW64 ~/downloads
$ cd Stalin
_ENOVO@StalinPC MINGW64 ~/downloads/Stalin
$ cd CSB-master
_ENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master
$ cd unix
.ENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix
$ cd sandbox
$ ls -lh ../data/Marra2014_data.fasta
-rw-r--r-- 1 LENOVO 197121 553K May 10 09:59 ../data/Marra2014_data.fasta
$ cp ../data/Marra2014_data.fasta my_file.fasta
'Papers and reviews'/ my_file.fasta
_ENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox
 grep -c isogroup00036 my_file.fasta | wc -l
.ENOVO@StalinPC MINGw64 ~/downloads/Stalin/CSB-master/unix/sandbox
$ grep -c isogroup00036 my_file.fasta
NINGW64:/c/Users/LENOVO/downloads/Stalin/CSB-master/unix/sandbox
                                                                    X
LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox

$ cat my_file.fasta | tr -s " " "," | head -n 3

>contig00001,length=527,numreads=2,gene=isogroup00001,status=it_thresh
```

.ENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox



```
LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox

$ grep '>' my_file.fasta | cut -d "," -f 4 | sort | uniq | wc -l

955
```

```
MINGW64:/c/Users/LENOVO/downloads/Stalin/CSB-master/unix/sandbox

LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox

$ grep '>' my_file.fasta | cut -d "," -f 1,3 | head -n 3
>contig00001 length=527 numreads=2 gene=isogroup00001 status=it_thresh
>contig00002 length=551 numreads=8 gene=isogroup00001 status=it_thresh
>contig00003 length=541 numreads=2 gene=isogroup00001 status=it_thresh
```

```
LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox $ grep '>' my_file.fasta | cut -d "," -f 1,3 | sort -t "-" -k 2 -n
                              numreads=2
>contig00001
                                            gene=isogroup00001
                length=527
                                                                    status=it_thresh
                length=551
                                                                   status=it_thresh
status=it_thresh
status=it_thresh
>contig00002
                                            gene=isogroup00001
                              numreads=8
>contig00003
                length=541
                              numreads=2
                                            gene=isogroup00001
>contig00004
                length=291
                              numreads=3
                                            gene=isogroup00001
                length=580
contig00005
                              numreads=12
                                             gene=isogroup00001 status=it_thresh
                length=3288
                               numreads=35
                                              gene=isogroup00001 status=it_thresh
gene=isogroup00001 status=it_thresh
>contig00006
                length=1119
>contig00008
                               numreads=10
>contig00010
                length=202 numreads=4 gene=isogroup00001 status=it_thresh
                length=5563
                                              gene=isogroup00001 status=it_thresh
                               numreads=61
>contig00011
>contig00012
>contig00013
                length=824
                                             gene=isogroup00001 status=it_thresh
                              numreads=10
                length=707
                                            gene=isogroup00001 status=it_thresh
                              numreads=6
                                                                   status=it_thresh
>contig00014
                length=568
                              numreads=2
                                            gene=isogroup00001
>contig00015
                length=123
                              numreads=4
                                            gene=isogroup00001
                                                                   status=it_thresh
                               numreads=22
                                              gene=isogroup00001 status=it_thresh
>contig00016
                length=1721
>contig00017
                length=788 numreads=15
                                             gene=isogroup00001 status=it_thresh
                length=1886
                              numreads=12
>contig00018
                                              gene=isogroup00001 status=it_thresh
                              numreads=3 gene=isogroup00001 status=it_thresh
numreads=10 gene=isogroup00001 status=it_thresh
>contig00022
>contig00023
                length=802
                length=885
>contig00025
                length=142
                              numreads=3
                                            gene=isogroup00001 status=it_thresh
                length=301
                                                                    status=it_thresh
                              numreads=4
>contig00026
                                            gene=isogroup00001
>contig00027
>contig00028
                                                                   status=it_thresh
status=it_thresh
                length=178
                              numreads=7
                                            gene=isogroup00001
                length=135
                                            gene=isogroup00001
                              numreads=2
>contig00029
                length=465
                              numreads=5
                                            gene=isogroup00001
                                                                    status=it_thresh
>contig00030
                length=121
                              numreads=3
                                            gene=isogroup00001
                                                                    status=it_thresh
                                                                    status=it_thresh
status=it_thresh
>contig00031
                length=595
                              numreads=9
                                            gene=isogroup00001
>contig00032
                length=1202
                              numreads=5
                                             gene=isogroup00001
>contig00033
                length=540
                              numreads=27
                                             gene=isogroup00001
                                                                     status=it_thresh
                length=1004 numreads=6 gene=isogroup00001 status=it_thresh
length=566 numreads=8 gene=isogroup00001 status=it_thresh
length=3571 numreads=8 gene=isogroup00001 status=it_thresh
>contig00034
contig00035>
>contig00036
                               numreads=10
                                              gene=isogroup00001 status=it_thresh
                length=1168
>contig00037
>contig00039
>contig00040
                                             gene=isogroup00001 status=it_thresh
gene=isogroup00001 status=it_thresh
                length=371 numreads=94
                length=2505
                               numreads=15
>contig00042
                length=620
                              numreads=11
                                             gene=isogroup00001 status=it_thresh
                                             gene=isogroup00001 status=it_thresh
>contig00044
                length=1301
                               numreads=4
>contig00045
                length=563
                              numreads=6
                                           gene=isogroup00001 status=it_thresh
>contig00046
                length=1046
                                              gene=isogroup00001 status=it_thresh
                               numreads=44
                length=211
>contig00048
                              numreads=4
                                            gene=isogroup00001 status=it_thresh
>contig00049
                length=740
                              numreads=3
                                            gene=isogroup00001
                                                                   status=it_thresh
```

```
LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox

$ grep '>' my_file.fasta | cut -d "," -f 1,3 | sort -t "-" -k 2 -n | head -n 5
>contig00001 length=527 numreads=2 gene=isogroup00001 status=it_thresh
>contig00002 length=551 numreads=8 gene=isogroup00001 status=it_thresh
>contig00003 length=541 numreads=2 gene=isogroup00001 status=it_thresh
>contig00004 length=291 numreads=3 gene=isogroup00001 status=it_thresh
>contig00005 length=580 numreads=12 gene=isogroup00001 status=it_thresh
```



```
LENOVO@StalinPC MINGW64 ~/downloads/Stalin/CSB-master/unix/sandbox

$ grep '>' my_file.fasta | cut -d "," -f 1,3 | sort -t "-" -k 2 -n -r | head -n 1

>contig01385 length=1965 numreads=7 gene=isogroup00043 status=isotig
```

```
MINGW64:/c/Users/LENOVO/downloads/stalin/CSB-master/unix/data — X

GNU nano 7.2 Ejercicio1_10_2.sh
#!/bin/bash

#¿Cuántas veces fueron los niveles de los individuos 3 y 27 grabados?

#Para los individuos 3

cut -f 1 Gesquiere2011_data.csv | grep -w 3 | grep -c 3

#Para los individuos 27

cut -f 1 Gesquiere2011_data.csv | grep -w 27 | grep -c 27
```

```
MINGW64:/c/Users/LENOVO/downloads/stalin/CSB-master/unix/data
                                                                              X
 .ENOVO@StalinPC MINGW64 ~/downloads/stalin/CSB-master/unix/data
$ nano Ejercicio1_10_2.sh
_ENOVO@StalinPC MINGW64 ~/downloads/stalin/CSB-master/unix/data
$ nano Ejercicio1_10_2.sh
LENOVO@StalinPC MINGW64 ~/downloads/stalin/CSB-master/unix/data
$ cat Ejercicio1_10_2.sh
#!/bin/bash
#¿Cuántas veces fueron los niveles de los individuos 3 y 27 grabados?
#Para los individuos 3
cut -f 1 Gesquiere2011_data.csv | grep -w 3 | grep -c 3
#Para los individuos 27
cut -f 1 Gesquiere2011_data.csv | grep -w 27 | grep -c 27
LENOVO@StalinPC MINGW64 ~/downloads/stalin/CSB-master/unix/data
$ bash Ejercicio1_10_2.sh
```



```
NINGW64:/c/Users/LENOVO/downloads/stalin/CSB-master/unix/data
                                                                              X
  GNU nano 7.2
                                                                           Modified
                                   Ejercicio1_10_2.sh
 !/bin/bash
¿Cuántas veces fueron los niveles de los individuos 3 y 27 grabados?
#Para los individuos 3
cut -f 1 Gesquiere2011_data.csv | grep -w 3 | grep -c 3
#Para los individuos 27
cut -f 1 Gesquiere2011_data.csv | grep -w 27 | grep -c 27
#Luego se crea un vector para los IDS del documento
vdownloads=`tail -n +2 Gesquiere2001_data.csv | cut -f 1 | uniq
for x in $vdownloads
ids=`bash Conteo_Ejercicio1_10_2.sh Gesquiere2011_data.csv $x`
echo "ID:" $x "conteo:" $ids
```

```
MINGW64:/c/Users/LENOVO/downloads/stalin/CSB-master/unix/data — X

LENOVO@StalinPC MINGW64 ~/downloads/stalin/CSB-master/unix/data
$ bash Ejercicio1_10_2.sh
61
5
tail: cannot open 'Gesquiere2001_data.csv' for reading: No such file or directory

LENOVO@StalinPC MINGW64 ~/downloads/stalin/CSB-master/unix/data
$ |
```



```
NINGW64:/c/Users/LENOVO/downloads/stalin/CSB-master/unix/data
                                                                                                X
                                                                                             Modified
  GNU nano 7.2
                                                 netsize.sh
 #!/bin/bash
echo "Filename:"
echo $1
echo "Number of rows:"
cat $1 |wc -l
echo "Number of columns:"
head -n 1 $1 | tr -d ' ' | tr -d '\n' | wc -c
LENOVO@StalinPC MINGW64 ~/downloads/stalin/CSB-master/unix/data

$ bash netsize.sh Saavedra2013/n1.txt

Filename:
Saavedra2013/n1.txt
Number of rows:
97
Number of columns:
80
 NINGW64:/c/Users/LENOVO/downloads/stalin/CSB-master/unix/data
                                                                                                \times
                                              netsize_all.sh
                                                                                             Modified
  GNU nano 7.2
 !/bin/bash
FILES=../data/Saavedra2013/*.txt
for f in $FILES
          echo $f $myrow $mycol
done
```



```
MINGW64:/c/Users/LENOVO/downloads/stalin/CSB-master/unix/data
                                                                                         X
_ENOVO@StalinPC MINGW64 ~/downloads/stalin/CSB-master/unix/data
$ bash netsize_all.sh
../data/Saavedra2013/n1.txt 97 80
../data/Saavedra2013/n10.txt 14 20
../data/Saavedra2013/n11.txt 270 91
../data/Saavedra2013/n12.txt 7 72
../data/Saavedra2013/n13.txt 61 17
../data/Saavedra2013/n14.txt 35 15
../data/Saavedra2013/n15.txt 38 11
../data/Saavedra2013/n16.txt 118 24
../data/Saavedra2013/n17.txt 76 31
../data/Saavedra2013/n18.txt 13 14
../data/Saavedra2013/n19.txt 10 16
 ./data/Saavedra2013/n2.txt 62 41
./data/Saavedra2013/n20.txt 18 7
./data/Saavedra2013/n21.txt 19 45
./data/Saavedra2013/n22.txt 19 36
../data/Saavedra2013/n23.txt 179 26
../data/Saavedra2013/n24.txt 80 28
../data/Saavedra2013/n25.txt 17 16
../data/Saavedra2013/n26.txt 82 40
../data/Saavedra2013/n27.txt 27
./data/Saavedra2013/n28.txt 90 19
../data/Saavedra2013/n29.txt 61 25
../data/Saavedra2013/n3.txt 25 36
./data/Saavedra2013/n30.txt 8 19
 ./data/Saavedra2013/n31.txt 28 25
./data/Saavedra2013/n32.txt 45 21
./data/Saavedra2013/n33.txt 70 20
./data/Saavedra2013/n34.txt 79 25
../data/Saavedra2013/n35.txt 14 8
../data/Saavedra2013/n36.txt 40 169
../data/Saavedra2013/n37.txt 44 13
../data/Saavedra2013/n38.txt 51 99
../data/Saavedra2013/n39.txt 33 25
./data/Saavedra2013/n4.txt 101 11
../data/Saavedra2013/n40.txt 28 18
../data/Saavedra2013/n41.txt 12 10
./data/Saavedra2013/n42.txt 42 8
./data/Saavedra2013/n43.txt 55 29
./data/Saavedra2013/n44.txt 56 9
./data/Saavedra2013/n45.txt 36 61
./data/Saavedra2013/n46.txt 58 17
../data/Saavedra2013/n47.txt 139 41
 ./data/Saavedra2013/n48.txt 118 49
  /data/Saavedra2013/n49.txt 47 23
```



```
MINGW64:/c/Users/LENOVO/downloads/stalin/CSB-master/unix/data — X

GNU nano 7.2 explore.sh Modified

# $1 is the file name
# $2 is the column of interest

echo "Column name"

cut -d ',' -f $2 $1 | head -n 1

echo "Number of distinct values:"

cut -d ',' -f $2 $1 | tail -n +2 | sort | uniq | wc -l

echo "Minium value:"

cut -d ',' -f $2 $1 | tail -n +2 | sort -n | head -n 1

echo "Maxium value:"

cut -d ',' -f $2 $1 | tail -n +2 | sort -n | tail -n 1

MINGW64:/c/Users/LENOVO/downloads/stalin/CSB-master/unix/data

$ nano explore.sh

LENOVO@StalinPC MINGW64 ~/downloads/stalin/CSB-master/unix/data
```

```
MINGW64:/c/Users/LENOVO/downloads/stalin/CSB-master/unix/data

LENOVO@StalinPC MINGW64 ~/downloads/stalin/CSB-master/unix/data

nano explore.sh

LENOVO@StalinPC MINGW64 ~/downloads/stalin/CSB-master/unix/data

bash explore.sh Buzzard2015_data.csv 7

column name
biomass
Number of distinct values:

285

Minium value:

1.048466198

Maxium value:

14897.29471
```

```
LENOVO@StalinPC MINGW64 ~/downloads/stalin/CSB-master/unix/data
$ bash explore.sh Buzzard2015_data.csv 9
Column name
SLA
Number of distinct values:
144
Minium value:

Maxium value:
369.6
```