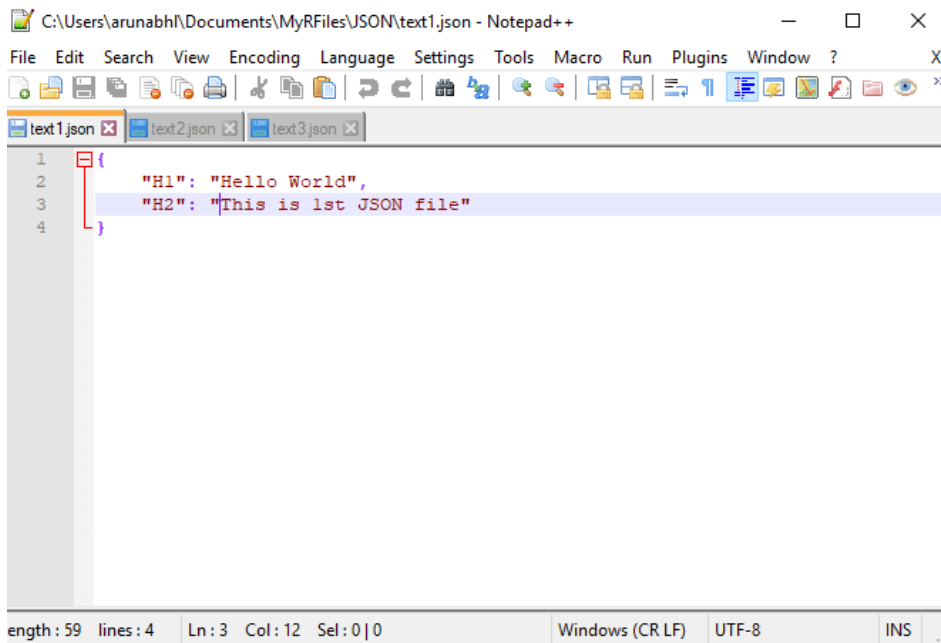


Problem Statement

1. Read multiple json files into a working directory for further converting into a dataset. I have files text1, text2, text3 in the directory json.

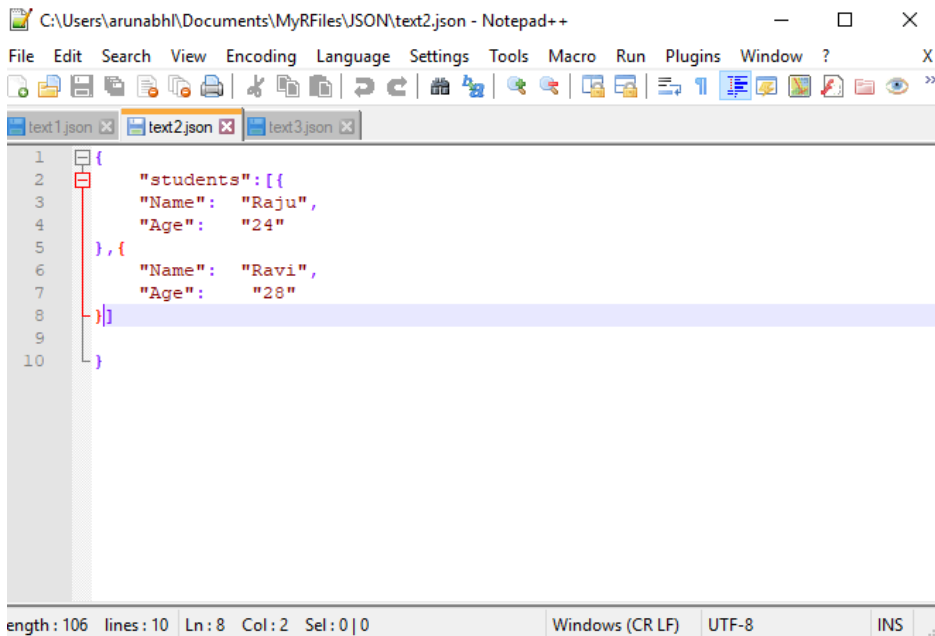
Answer: Created 3 json files under ("C:/Users/arunabh/MyFiles/JSON") folder



The screenshot shows the Notepad++ application with the file 'text1.json' open. The file contains a JSON object with two key-value pairs. The status bar at the bottom indicates the file is 59 characters long, 4 lines, and the cursor is at line 3, column 12.

```
1 {  
2   "H1": "Hello World",  
3   "H2": "This is 1st JSON file"  
4 }
```

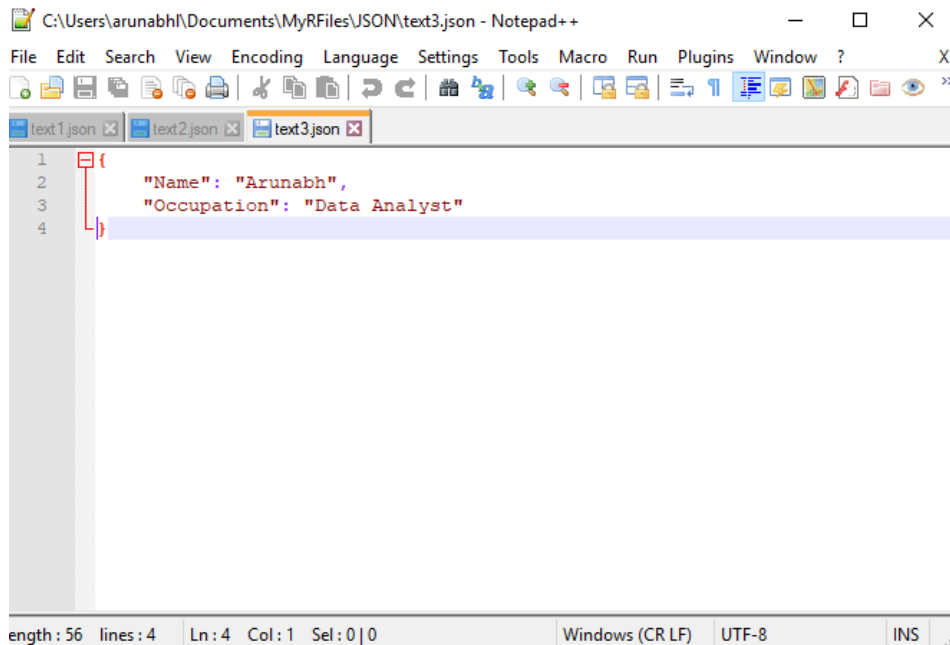
length : 59 lines : 4 Ln : 3 Col : 12 Sel : 0 | 0 Windows (CR LF) UTF-8 INS



The screenshot shows the Notepad++ application with the file 'text2.json' open. The file contains a JSON object with a 'students' array containing two student objects. The status bar at the bottom indicates the file is 106 characters long, 10 lines, and the cursor is at line 8, column 2.

```
1 {  
2   "students": [{  
3     "Name": "Raju",  
4     "Age": "24"  
5   }, {  
6     "Name": "Ravi",  
7     "Age": "28"  
8   }]  
9 }  
10 }
```

length : 106 lines : 10 Ln : 8 Col : 2 Sel : 0 | 0 Windows (CR LF) UTF-8 INS



Step 1- Install *rjson package* and then *library(rjson)*

```
install.packages("rjson")
```

```
library(rjson)
```

Step 2- set the working directory where the JSON files are located.

```
setwd("C:/Users/arunabh/Documents/MyRFiles/JSON")
```

Step 3- Use the command-

```
x <-list.files(pattern="*.json")
```

Here the list of all files under the working directory with the pattern or extension as **.json** is declared on a variable 'x'.

Output -

```
> x <-list.files(pattern="*.json")
> x
[1] "text1.json" "text2.json" "text3.json"
> |
```

Step 4 - Use lapply()

```
l<- lapply(x,function(x) fromJSON(file= x))
```

Output -

```
> l<- lapply(x,function(x) fromJSON(file= x))
> l
[[1]]
[[1]]$H1
[1] "Hello World"

[[1]]$H2
[1] "This is 1st JSON file"

[[2]]
[[2]]$students
[[2]]$students[[1]]
[[2]]$students[[1]]$Name
[1] "Raju"

[[2]]$students[[1]]$Age
[1] "24"

[[2]]$students[[2]]
[[2]]$students[[2]]$Name
[1] "Ravi"

[[2]]$students[[2]]$Age
[1] "28"

[[3]]
[[3]]$Name
[1] "Arunabh"
```

Here, the lapply functions is used on the variable 'x' and the output of all the jSoN files are as "list".

Step 5- Converting the list to dataframe in below script

```
df5<-as.data.frame(do.call("cbind", l))
```

Output-

```
> df5<- as.data.frame(do.call("cbind", l))
> df5
      V1      V2      V3
H1      Hello World Raju, 24, Ravi, 28      Arunabh
H2 This is 1st JSON file Raju, 24, Ravi, 28 Data Analyst
> |
```

2. Parse the following JSON into a data frame

```
js<-'{  
  "name": null, "release_date_local": null, "title": "3 (2011)",  
  "opening_weekend_take": 1234, "year": 2011,  
  "release_date_wide": "2011-09-16", "gross": 59954  
'
```

ANSWER:

We need to convert the JSON into a list and then to a data frame.

Step 1: Install rjson package and then pass the command library(rjson)

```
install.packages("rjson")  
library(rjson)
```

Step 2: Assign the json into a variable

```
js_ans<- '{  
  "name": null, "release_date_local": null, "title": "3 (2011)",  
  "opening_weekend_take": 1234, "year": 2011,  
  "release_date_wide": "2011-09-16", "gross": 59954  
'
```

Output:

```
> js_ans<- '{  
+ "name": null, "release_date_local": null, "title": "3 (2011)",  
+ "opening_weekend_take": 1234, "year": 2011,  
+ "release_date_wide": "2011-09-16", "gross": 59954  
+ }'  
> js_ans  
[1] "{\n\"name\": null, \"release_date_local\": null, \"title\": \"3 (2011)\",  
\n\"opening_weekend_take\": 1234, \"year\": 2011,\n\"release_date_wide\": \"2011-09-16\", \"gross\": 59954\n}"
```

Step 3: Converting json string into list

```
lst<- fromJSON(js_ans, simplify = T)
```

```
lst
```

```
> #Convert JSON to list
> lst<- fromJSON(js_ans, simplify = T)
> lst
$name
NULL

$release_date_local
NULL

$title
[1] "3 (2011)"

$opening_weekend_take
[1] 1234

$year
[1] 2011

$release_date_wide
[1] "2011-09-16"

$gross
[1] 59954
```

Step4: Replace all NULL values to NA

```
lst<- replace(lst, lst=="NULL", NA)
```

```
lst
```

```
> #Replace All NULL values to NA
> lst<- replace(lst, lst=="NULL", NA)
> lst
$name
[1] NA

$release_date_local
[1] NA

$title
[1] "3 (2011)"

$opening_weekend_take
[1] 1234

$year
[1] 2011

$release_date_wide
[1] "2011-09-16"

$gross
[1] 59954
```

Step 5: converts list to dataframe in column wise (**FINAL OUTPUT**)

```
df<-as.data.frame(do.call("cbind", lst))
```

```
df
```

```
> #Converting into data frame column wise
> df<- as.data.frame(do.call("cbind", lst))
> df
  name release_date_local  title opening_weekend_take year
1 <NA>                  <NA> 3 (2011)                1234 2011
  release_date_wide gross
1      2011-09-16 59954
```

3. Write a script for variable binning using R.

Answer- Variable Binning in R is the method of converting a numerical variable of a dataset into a categorical variable. To do so we use the cut() command, which divides the range of a variable (say x) into intervals.

Example - I have imported a csv file called "LungCapData" consisting of different variables like "Age", "Height", "Smoke", "Gender" and "Caesarean". Considering the "Height" variable we are going to divide this variable into different ranges.

Step1: Importing the File

```
LungCapData<- read.csv(file.choose(), header = T, sep = "\t")
```

```
LungCapData
```

Step2: #Creating a new Variable to save Height

```
CatHeight<- LungCapData$Height
```

```
CatHeight[1:10]
```

Output:

```
CatHeight[1:10]
[1] 62.1 74.7 69.7 71.0 56.9 58.7 63.3 70.4 70.5 59.2
```

Step3: #Dividing the height categories A<=50, B=50-55, C=55-60, D=60-65, E=65-70, F=70+

#Also adding a column to the dataset

```
LungCapData$bins<- cut(CatHeight , breaks = c(0,50,55,60,65,70,100), labels =  
c("A","B","C","D","E","F"))
```

LungCapData

```
> LungCapData$bins<- cut(CatHeight , breaks = c(0,50,55,60,65,70,100), labels =  
= c("A","B","C","D","E","F"))  
> LungCapData
```

	LungCap	Age	Height	Smoke	Gender	Caesarean	bins
1	6.475	6	62.1	no	male	no	D
2	10.125	18	74.7	yes	female	no	F
3	9.550	16	69.7	no	female	yes	E
4	11.125	14	71.0	no	male	no	F
5	4.800	5	56.9	no	male	no	C

Explanations:

breaks() is used to break as per as the ranges required

labels =c() gives label names to the CatHeight bins.

LungCapData\$bins is creating a new column consisting of the bins labels information.

Please find the **R-script** attached to see the **Output** in R named "R Bins By Variable".