

SHETH L.U.J.AND SIR M.V. COLLEGE
PRACTICAL NO.13
Data Analysis with SAS / SPSS /R

AIM:- Identifying and handling duplicates using distinct() (R studio).

INPUT:-

R Script: Identifying and Handling Duplicates

Dataset: winequality-red.csv

Using distinct() from dplyr

library(dplyr)

1. READ YOUR WINE DATA

wine_df <- read.csv("D:/S079_VIBHUTI/ADV PYTHON FOR DATA
SCIENCE/winequality-red.csv")

print("--- 1. Original Wine Dataset ---")

print(head(wine_df))

2. IDENTIFYING DUPLICATES (Exact row duplicates)

duplicates_report <- wine_df %>%

group_by(across(everything())) %>% # group by ALL columns

count() %>% # count duplicates

filter(n > 1) # keep only duplicates

print("--- 2. Rows that are duplicated (Full duplicate report) ---")

print(duplicates_report)

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PRACTICAL NO.13
Data Analysis with SAS / SPSS /R

3. REMOVING EXACT DUPLICATE ROWS

```
clean_exact <- wine_df %>%  
  distinct()  # removes full duplicate rows
```

```
print("--- 3. Dataset After Removing Exact Duplicates ---")  
print(clean_exact)
```

4. HANDLING DUPLICATES BASED ON ONE COLUMN (Example: quality)

```
# Scenario: Keep ONLY ONE ROW per quality level  
# This is like your 'unique customers' example.
```

```
unique_quality <- wine_df %>%  
  distinct(quality, .keep_all = TRUE)
```

```
print("--- 4. Unique Quality Values (Only first appearance kept) ---")  
print(unique_quality)
```

OUTPUT:-

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PRACTICAL NO.13

Data Analysis with SAS / SPSS /R

```

Console Terminal Background Jobs
R - R 4.1.2 - ~/
> # R Script: Identifying and Handling Duplicates
> # Dataset: winequality-red.csv
> # using distinct() from dplyr

> library(dplyr)

> # 1. READ YOUR WINE DATA

> wine_df <- read.csv("D:/S079_VIBHUTI/ADV PYTHON FOR DATA SCIENCE/winequality-red.csv")

> print("--- 1. Original wine Dataset ---")
[1] "--- 1. Original wine Dataset ---"
> print(head(wine_df))
  fixed.acidity volatile.acidity citric.acid residual.sugar chlorides free.sulfur.dioxide
1          7.4           0.70         0.00           1.9      0.076             11
2          7.8           0.88         0.00           2.6      0.098             25
3          7.8           0.76         0.04           2.3      0.092             15
4         11.2           0.28         0.56           1.9      0.075             17
5          7.4           0.70         0.00           1.9      0.076             11
6          7.4           0.66         0.00           1.8      0.075             13
total.sulfur.dioxide density pH sulphates alcohol quality
1             34  0.9978 3.51         0.56      9.4         5
2             67  0.9968 3.20         0.68      9.8         5
3             54  0.9970 3.26         0.65      9.8         5
4             60  0.9980 3.16         0.58      9.8         6
5             34  0.9978 3.51         0.56      9.4         5
6             40  0.9978 3.51         0.56      9.4         5

> # 2. IDENTIFYING DUPLICATES (Exact row duplicates)

> duplicates_report <- wine_df %>%
+   group_by(across(everything())) %>% # group by ALL columns
+   count() %>% # count duplicates
+   filter(n > 1) # keep only duplicates

> print("--- 2. Rows that are duplicated (Full duplicate report) ---")
[1] "--- 2. Rows that are duplicated (Full duplicate report) ---"
> print(duplicates_report)
# A tibble: 220 x 13

```

```

Console Terminal Background Jobs
R - R 4.1.2 - ~/
# free.sulfur.dioxide, total.sulfur.dioxide, density, pH, sulphates, alcohol, quality [220]
fixed.acidity volatile.acidity citric.acid residual.sugar chlorides free.sulfur.dioxide
<dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
1          5.2           0.34         0           1.8      0.05             27
2          5.6           0.5           0.09         2.3      0.049             17
3          5.6           0.54         0.04         1.7      0.049             5
4          5.6           0.66         0           2.2      0.087             3
5          5.9           0.61         0.08         2.1      0.071             16
6          6           0.5           0           1.4      0.057             15
7          6           0.51         0           2.1      0.064             40
8          6.1           0.32         0.25         2.3      0.071             23
9          6.2           0.36         0.24         2.2      0.095             19
10         6.2           0.56         0.09         1.7      0.053             24
# i 210 more rows
# i 7 more variables: total.sulfur.dioxide <dbl>, density <dbl>, pH <dbl>, sulphates <dbl>,
# alcohol <dbl>, quality <int>, n <int>
# i use 'print(n = ...)' to see more rows

> # 3. REMOVING EXACT DUPLICATE ROWS

> clean_exact <- wine_df %>%
+   distinct() # removes full duplicate rows

> print("--- 3. Dataset After Removing Exact Duplicates ---")
[1] "--- 3. Dataset After Removing Exact Duplicates ---"
> print(clean_exact)
  fixed.acidity volatile.acidity citric.acid residual.sugar chlorides free.sulfur.dioxide
1          7.4           0.700         0.00           1.90      0.076             11
2          7.8           0.880         0.00           2.60      0.098             25
3          7.8           0.760         0.04           2.30      0.092             15
4         11.2           0.280         0.56           1.90      0.075             17
5          7.4           0.660         0.00           1.80      0.075             13
6          7.9           0.600         0.06           1.60      0.069             15
7          7.3           0.650         0.00           1.20      0.065             15
8          7.8           0.580         0.02           2.00      0.073             9
9          7.5           0.500         0.36           6.10      0.071             17
10         6.7           0.580         0.08           1.80      0.097             15
11         5.6           0.615         0.00           1.60      0.089             16
12         7.8           0.610         0.29           1.60      0.114             9
13         8.9           0.620         0.18           3.80      0.176             52
14         8.9           0.620         0.19           3.90      0.170             51
15         8.5           0.280         0.56           1.80      0.092             35

```

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PRACTICAL NO.13

Data Analysis with SAS / SPSS /R

Console	Terminal	Background Jobs
R - R 4.1.2 - ~/		
54	7.5	0.630
55	7.8	0.590
56	7.3	0.390
57	8.8	0.400
58	7.7	0.690
59	7.5	0.520
60	7.0	0.735
61	7.2	0.725
62	7.5	0.520
63	6.6	0.705
64	9.3	0.320
65	8.0	0.705
66	7.7	0.630
67	7.7	0.670
68	7.7	0.690
69	8.3	0.675
70	9.7	0.320
71	8.8	0.410
72	6.8	0.785
73	6.7	0.750
74	8.3	0.625
75	6.2	0.450
76	7.8	0.430
77	7.4	0.500
78	7.3	0.670
79	6.3	0.300
80	6.9	0.550
81	8.6	0.490
82	7.7	0.490
83	9.3	0.390
total.sulfur.dioxide density pH sulphates alcohol quality		
1	34	0.9978 3.51 0.56 9.4 5
2	67	0.9968 3.20 0.68 9.8 5
3	54	0.9970 3.26 0.65 9.8 5
4	60	0.9980 3.16 0.58 9.8 6
5	40	0.9978 3.51 0.56 9.4 5
6	59	0.9964 3.30 0.46 9.4 5
7	21	0.9946 3.39 0.47 10.0 7
8	18	0.9968 3.36 0.57 9.5 7
9	102	0.9978 3.35 0.80 10.5 5
10	65	0.9959 3.28 0.54 9.2 5
11	59	0.9943 3.58 0.52 9.9 5
12	29	0.9974 3.26 1.56 9.1 5

Console	Terminal	Background Jobs
R - R 4.1.2 - ~/		
45	12	0.9958 3.34 0.56 9.2 5
46	96	0.9954 3.32 0.58 9.2 5
47	23	0.9971 3.15 0.74 9.2 5
48	15	0.9956 3.40 0.63 9.4 6
49	14	0.9955 3.39 0.64 9.4 6
50	119	0.9970 3.20 0.56 9.4 5
51	73	0.9955 3.17 0.63 10.2 6
52	45	0.9978 3.34 0.53 9.5 5
53	10	0.9971 3.04 0.63 9.6 5
54	110	0.9983 3.26 0.77 9.4 5
55	54	0.9975 3.43 0.59 10.0 5
56	46	0.9962 3.41 0.54 9.4 6
57	52	0.9980 3.44 0.64 9.2 5
58	112	0.9968 3.21 0.71 9.3 5
59	35	0.9968 3.38 0.62 9.5 7
60	54	0.9966 3.39 0.57 9.8 5
61	11	0.9962 3.41 0.39 10.9 5
62	39	0.9968 3.42 0.58 9.6 5
63	15	0.9962 3.44 0.58 10.7 5
64	65	0.9969 3.28 0.79 10.7 5
65	19	0.9962 3.34 0.95 10.5 6
66	27	0.9967 3.32 0.54 9.5 6
67	96	0.9962 3.32 0.48 9.5 5
68	94	0.9961 3.31 0.48 9.5 5
69	43	0.9976 3.31 0.53 9.2 4
70	83	0.9984 3.28 0.82 9.6 5
71	42	0.9986 3.54 0.66 10.5 5
72	30	0.9966 3.52 0.55 10.7 6
73	80	0.9958 3.38 0.52 10.1 5
74	119	0.9972 3.16 1.12 9.1 4
75	15	0.9958 3.41 0.56 9.2 5
76	67	0.9974 3.13 1.28 9.4 5
77	73	0.9970 3.36 0.57 9.1 5
78	51	0.9969 3.16 1.14 9.4 5
79	61	0.9959 3.44 0.78 10.3 6
80	40	0.9961 3.41 0.59 10.1 5
81	136	0.9972 2.93 1.95 9.9 6
82	31	0.9966 3.39 0.64 9.6 5
83	125	0.9978 3.14 1.22 9.5 5
[reached 'max' / getoption("max.print") -- omitted 1276 rows]		
>		
>		
> # 4. HANDLING DUPLICATES BASED ON ONE COLUMN (Example: quality)		

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PRACTICAL NO.13
Data Analysis with SAS / SPSS /R

```

Console Terminal Background Jobs
R 4.1.2 ~ /
71      42 0.9986 3.54      0.66 10.5      5
72      30 0.9966 3.52      0.55 10.7      6
73      80 0.9958 3.38      0.52 10.1      5
74     119 0.9972 3.16      1.12  9.1      4
75      15 0.9958 3.41      0.56  9.2      5
76      67 0.9974 3.13      1.28  9.4      5
77      73 0.9970 3.36      0.57  9.1      5
78      51 0.9969 3.16      1.14  9.4      5
79      61 0.9959 3.44      0.78 10.3      6
80      40 0.9961 3.41      0.59 10.1      5
81     136 0.9972 2.93      1.95  9.9      6
82      31 0.9966 3.39      0.64  9.6      5
83     125 0.9978 3.14      1.22  9.5      5
[ reached 'max' / getOption("max.print") -- omitted 1276 rows ]
>
>
> # 4. HANDLING DUPLICATES BASED ON ONE COLUMN (Example: quality)
>
> # Scenario: Keep ONLY ONE ROW per quality level
> # This is like your 'unique customers' example.
>
> unique_quality <- wine_df %>%
+   distinct(quality, .keep_all = TRUE)
>
> print("--- 4. Unique Quality values (only first appearance kept) ---")
[1] "--- 4. Unique Quality values (only first appearance kept) ---"
> print(unique_quality)
  fixed.acidity volatile.acidity citric.acid residual.sugar chlorides free.sulfur.dioxide
1         7.4          0.70         0.00          1.9      0.076          11
2        11.2          0.28         0.56          1.9      0.075          17
3         7.3          0.65         0.00          1.2      0.065          15
4         7.4          0.59         0.08          4.4      0.086           6
5         7.9          0.35         0.46          3.6      0.078          15
6        11.6          0.58         0.66          2.2      0.074          10
 total.sulfur.dioxide density    pH sulphates alcohol quality
1          34 0.9978 3.51      0.56      9.4      5
2          60 0.9980 3.16      0.58      9.8      6
3          21 0.9946 3.39      0.47     10.0      7
4          29 0.9974 3.38      0.50      9.0      4
5          37 0.9973 3.35      0.86     12.8      8
6          47 1.0008 3.25      0.57      9.0      3
>

```