

MVLU COLLEGE.

PRACTICAL NO :- 12

AIM :- Combining datasets vertically (concatenation) using rbind() (R).

CODE :-

```
library(dplyr)
```

```
iris_df <- read.csv("C:/Users/itlab/Downloads/S100/Iris.csv")
```

```
flower_df <- read.csv("C:/Users/itlab/Downloads/S100/flower_dataset.csv")
```

```
print("--- Column Names Before Cleaning ---")
```

```
print(names(iris_df))
```

```
print(names(flower_df))
```

```
iris_clean <- iris_df[, c("Species", "SepalLengthCm")]
```

```
names(iris_clean) <- c("Species", "Height")
```

```
# --- Prepare FLOWER dataset ---
```

```
flower_clean <- flower_df[, c("species", "height_cm")]
```

```
names(flower_clean) <- c("Species", "Height")
```

```
# Convert heights to numeric
```

```
iris_clean$Height <- as.numeric(iris_clean$Height)
```

```
flower_clean$Height <- as.numeric(flower_clean$Height)
```

```
#
```

```
=====
```

```
=====
```

```
# 3. VERTICAL CONCATENATION USING rbind()
```

```
#
```

```
=====
```

```
=====
```

```
combined_data <- rbind(iris_clean, flower_clean)
```

```
# 4. OUTPUT
```

```
print("--- Combined Data Summary ---")
```

```
print(paste("Rows in Iris dataset:", nrow(iris_clean)))
```

```
print(paste("Rows in Flower dataset:", nrow(flower_clean)))
```

```
print(paste("Total Expected Rows:", nrow(iris_clean) + nrow(flower_clean)))
```

```
print(paste("Total Actual Rows:", nrow(combined_data)))
```

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```
print("--- Preview of Combined Data ---")
print(head(combined_data))
print(tail(combined_data))
```

