

- `l[2]` return sub-list containing that element; `l[[2]]` return element

Data Frames

- `data.frame(col1,col2,...)`
- Access Methods: `df[1:2]`, `df[c('col1','col2')]`, `df$col1`
- Assign column names: `names(df) <- c('col1', 'col2', col3')`
- `attach(df)` - can directly access df variable `x` without need for `df$x`

Data Input

- CSV: `read.csv('/path/to/file.csv', header=T/F, sep=',')`
- Export delimited file: `write.table(data, 'file.txt', sep='\t')`

Viewing Data

- `ls()` list objects in the working environment
- `names(data)` list variables (e.g. col names of dataframe) in data
- `str(data)` list structure of data
- `dim(data)` dimensions of data
- `class(data)` list data types of an object
- `length(data)` number of elements or components in the object
- `head(data, n=2)` return the first 2 rows of data
- `tail(data, n=1)` return the last rows of data

Missing Data

- `x <- c(1,2,NA,3)`
- `mean(x)` returns NA, `mean(x, na.rm=T)` returns 2
- `complete.cases()` returns logical vector of rows without NA
- `df[!complete.cases(df)]` return complete rows
- `df <- na.omit(df)` create new dataset without the missing data

Manipulating Data

- `c(obj1, obj2, ...)` combine objects into a vector
- `cbind(obj1, obj2, ...)` nrow don't change, ncol++
- `rbind(obj1, obj2, ...)` ncol don't change, nrow++

Operators & Control Structure

- addition: `+`
- subtraction: `-`
- multiply: `*`
- division: `/`
- exponential: `^` / `**`
- modulus: `x %% y`
- int division: `%/%`
- test if T: `isTRUE()`
- comparators: `<`, `<=`, `>`, `>=`, `==`, `!=`, `==`
- `if(cond) 1 else 2`
- `for(var in seq) 1`
- `while(cond) 1`
- `switch(expr, default_case, case1=result1 ...)`
- `||`, `&&` element-wise logical operator
- `||`, `&&` single element logical operator
- `ifelse(cond, yes, no)` similar to a?b:c

Numeric Functions

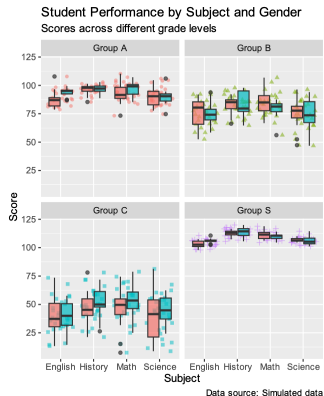
- absolute: `abs(x)`
- square root: `sqrt(x)`
- 3.4→4: `ceiling(x)`
- -3.4→-4: `floor(x)` round -∞
- -3.4→-3: `trunc(x)` round to 0
- `round(3.475, digits=2)` 3.48
- `signif(3.475, digits=2)` 3.5
- `cos()`, `sin()`, `tan()`, `acos()`...
- Natural log: `log(x)`
- Common log: `log10(x)`
- e^x : `exp(x)`
- `pretty(c(min, max), count)` evenly spaced numbers
- random variable: `rxxx(n, ...)`
- density function: `dxxx`
- cumulative: `pxxx` value → prob
- quantile: `qxxx` prob → value
- `xbinom(n,size,prob,lower.tail=T)` $T=P(X \leq x)$, $F=P(X > x)$
- `xnorm(_,m,sd)`
- `xpois(_,lambda)`
- `xunif(_,min,max)`

Plotting in R

- `plot(x)` plot scatterplot of x=index, y=x
- `plot(x, y)` plot scatterplot of x=x, y=y
 - `type`: p=point, l=lines, b=both
- `hist(x)` plot density histogram (distribution of x)
 - `breaks`: vector giving the breakpoints between histogram cells / function computing breakpoints / single value - number of cells / function computing num of cells

Plotting in ggplot2

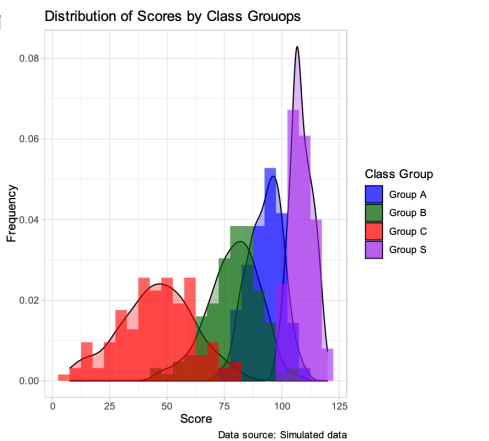
```
ggplot(data, aes(x=subject, y=score, fill=gender)) +
  geom_point(
    aes(shape = class, color = class),
    position = "jitter",
    alpha = 0.5
  ) +
  geom_boxplot(alpha = 0.7) +
  facet_wrap(~class, ncol = 2) +
  labs(x = "Subject",
       y = "Score",
       color = "class",
       fill = "Gender",
       title = "Student Performance by Subject and Gender",
       subtitle = "Scores across different grade levels",
       caption = "Data source: Simulated data")
```



```
ggplot(student_data, aes(x = study_hours, y = score)) +
  geom_point(aes(color = class), alpha = 0.7) +
  geom_smooth(method = "lm") +
  labs(x = "Study Hours",
       y = "Score",
       color = "Class Group",
       title = "Score vs. Study Hours",
       caption = "Data source: Simulated data") +
  scale_color_manual(values = c(
    "Group C" = "blue",
    "Group B" = "purple",
    "Group A" = "orange",
    "Group S" = "red"
  ))
```



```
ggplot(student_data, aes(x = score, fill = class)) +
  geom_density(alpha = 0.3) +
  geom_histogram(
    aes(y = ..density..),
    binwidth = 5,
    alpha = 0.6,
    position = "identity"
  ) +
  labs(x = "Score", y = "Frequency", fill = "Class Group",
       title = "Distribution of Scores by Class Groups",
       caption = "Data source: Simulated data") +
  theme_light() +
  scale_fill_manual(values = c(
    "Group C" = "red",
    "Group B" = "darkgreen",
    "Group A" = "blue",
    "Group S" = "purple"
  ))
```



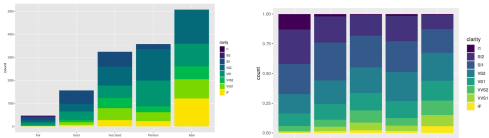
Coordinate Systems

```
g<- ggplot(...)
g + coord_flip() // Switch to Cartesian coordinate system
g + coord_polar() // Switch to polar coordinate system
```

General Template

```
ggplot(data = DATA) +
  GEOM_FUNCTION(
    mapping = aes(MAPPINGS),
    stat = STAT,
    position = POSITION
  ) +
  COORDINATE_FUNCTION() +
  FACET_FUNCTION() +
  LABEL_FUNCTION()
```

Position Arguments



- position: "identity"

For reproducibility: `set.seed(1234)`

Sequence generation: `seq_len(length) ≡ seq(1, length)`

`seq_along(obj) ≡ seq(1, length(obj))`

Regression

Sum of Squared Error (SSE): $S(w, b) = \sum_i e^2 = \sum_i (y_i - wx_i - b)^2$

$$\frac{\partial S}{\partial w} = 2 \sum_i (y_i - wx_i - b) * (-1) = 0, \quad \bar{x} = \frac{1}{n} \sum_i x_i \text{ and } \bar{y} = \frac{1}{n} \sum_i y_i$$

$$\frac{\partial S}{\partial b} = 2 \sum_i (y_i - wx_i - b) * (-x_i) = 0, \quad s_{xx} = \frac{1}{n-1} \sum_i (x_i - \bar{x})^2$$

$$\hat{w} = \frac{s_{xy}}{s_{xx}} \text{ and } \hat{b} = \bar{y} - \hat{w} \cdot \bar{x}, \quad s_{xy} = \frac{1}{n-1} \sum_i (x_i - \bar{x})(y_i - \bar{y})$$

R Linear Model lm()

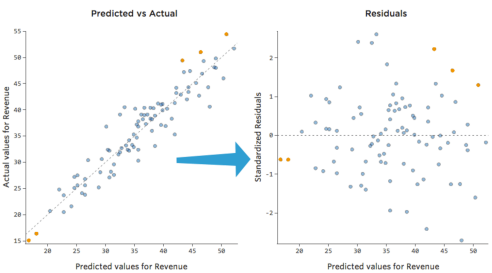
```
Univariate
x <- c(0, 2, 3)
y <- c(1, 1, 4)
dataset <- data.frame(x, y)
model <- lm(y ~ x, data = dataset)
coefficients(model)
predict(model, data.frame(x=2))

Multivariate
x1 <- c(1, 2, 3)
x2 <- c(4, 7, 8)
y <- c(9, 10, 11)
dataset <- data.frame(x1,x2,y)
model <- lm(y ~ x1 + x2, data = dataset)
predict(model, data.frame(x1=2, x2=3))
```

- `summary()`: comprehensive summary including coefficients, standard error, **R-squared**, F-statistic, t-value, and p-value
- `attributes()`: reveal various components stored in model, such as coefficients, residuals, fitted values, and the formula used in the call
- `coefficients()`: directly extract model's coeff. (intercept & slope)
- `predict()`: make predictions based on fitted model.

Residuals

$y_i = \hat{w}x_i + \hat{b} + \epsilon_i$, where ϵ_i is the left over term (or error) between ground truth and prediction. Formally, ϵ_i is called **residuals**



- **Homoscedasticity**: ϵ_i has constant variance for all i :
 - $\text{Var}(\epsilon_i | x_i) = \text{constant}^2$
- **Heteroscedasticity**: Intuitively, there is info inherent in the system not captured by the linear model
 - $\text{Var}(\epsilon_i | x_i) = f(x_i)$; where $f(x_i)$ is the not captured info

Measuring Model Performance

- **Observed**: $y = \{y_1, y_2, \dots, y_n\}$
- **Predicted**: $\hat{y} = \{\hat{y}_1, \hat{y}_2, \dots, \hat{y}_n\}$
- Total Sum of Squares (**TSS**): $\text{TSS} = \sum_i (y_i - \bar{y})^2$
- Explained Sum of Squares (**ESS**): $\text{ESS} = \sum_i (\hat{y}_i - \bar{y})^2$
- Residual Sum of Squares (**RSS**): $\text{RSS} = \sum_i (y_i - \hat{y}_i)^2$

TSS = ESS + RSS

- **RSS** is the "Unexplained Variability"

- **ESS** is the "Explained Variability"

$$R^2 = \frac{\text{ESS}}{\text{TSS}} = 1 - \frac{\text{RSS}}{\text{TSS}}$$

Time Series Forecasting

ARIMA Model

- **Definition**: Autoregressive Integrated Moving Average
 - Integration of autoregressive (AR) and moving average (MA) models
 - **AR model** forecast: linear combination of past values of variables
 - **MA model** forecast: linear combination of past forecast errors

$$y_t = \delta + \{\phi_1 y_{t-1} + \phi_2 y_{t-2} + \dots + \phi_p y_{t-p}\} + \{\theta_1 \epsilon_{t-1} + \theta_2 \epsilon_{t-2} + \dots + \theta_q \epsilon_{t-q}\} + \epsilon_t$$

$$\Rightarrow y_t = \delta + \sum_{i=1}^p \phi_i y_{t-i} + \sum_{j=1}^q \theta_j \epsilon_{t-j} + \epsilon_t$$

`install.packages('forecast')`

`library(forecast)`

`model <- auto.arima(AirPassengers)`

`plot(forecast(model, h=10*12))`

More on SVD

[SVD 分解] 性质: $A[m \times n] = U[m \times m] \Sigma[m \times n] (V^T)^{n \times n}$, $U^T U = V^T V = I$

1. 计算 $(A A^T)^{m \times m}$, 找到它的特征向量 \vec{u} , 它的特征值 λ 并开方 $\sigma = \sqrt{\lambda}$
2. 给 σ 降序排列得到序列 σ^* , 然后得到 $\Sigma = \text{diag}(\sigma^*)$, 其他地方补 0
3. 对所有 \vec{u} 按照对应的 σ^* 排序, 得到 U , 即: $U^{[m \times n]} = \begin{bmatrix} \vec{u}_1^T & \vec{u}_2^T & \dots & \vec{u}_n^T \\ |\vec{u}_1| & |\vec{u}_2| & \dots & |\vec{u}_n| \end{bmatrix}$
4. 根据 $A = U \Sigma V^T$, 得到 $V^T = \Sigma^* U^* A$, Σ^* 是左广义逆矩阵 ($\Sigma^* \Sigma = I$), 例如, $\Sigma = \begin{bmatrix} 2 & 0 \\ 0 & 3 \end{bmatrix} \Rightarrow \Sigma^* = \Sigma^{-1} = \frac{1}{\Sigma} = \begin{bmatrix} \frac{1}{2} & 0 \\ 0 & \frac{1}{3} \end{bmatrix}$, $\Sigma = \begin{bmatrix} 2 & 0 \\ 0 & 0 \end{bmatrix} \Rightarrow \Sigma^* = \begin{bmatrix} \frac{1}{2} & 0 \\ 0 & 0 \end{bmatrix}$ (pseudoinverse)

去除 $\sigma_k^2 = 0$ (甚至忽略较小的 σ_k^2) 对应的 U 、 V 的列, 实现数据压缩, 例如,

$$\begin{bmatrix} 1 & 1 \\ 0 & 0 \end{bmatrix} \begin{bmatrix} \frac{1}{\sqrt{2}} & 0 \\ 0 & \frac{1}{\sqrt{2}} \end{bmatrix} \begin{bmatrix} \frac{1}{\sqrt{2}} & \frac{1}{\sqrt{2}} \\ 0 & 0 \end{bmatrix}^T = \begin{bmatrix} \frac{1}{\sqrt{2}} & 0 \\ 0 & 0 \end{bmatrix} \begin{bmatrix} \frac{1}{\sqrt{2}} & \frac{1}{\sqrt{2}} \\ 0 & 0 \end{bmatrix}^T = \begin{bmatrix} \frac{1}{2} & \frac{1}{2} \\ 0 & 0 \end{bmatrix} \cdot \begin{bmatrix} \frac{1}{\sqrt{2}} & \frac{1}{\sqrt{2}} \\ 0 & 0 \end{bmatrix}^T$$
