# Normal Distribution Sampling

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```
library(tidyr) #the pipe (%>%) tool is extremely useful
library(MASS) # used for murnorm
```

#### $\mathbf{Q}\mathbf{1}$

Suppose Suppose  $X_1, X_2, Y_1, Y_2$  are mutually independent.

- $X_1$  and  $X_2$  are iid from  $N(\mu=0,\sigma^2=2^2)$   $Y_1$  and  $Y_2$  are iid from  $N(\mu=0,\sigma^2=1^2)$
- a) Calculate  $P(|X_1 X_2| > 2)$ 
  - Let  $X = X_1 X_2$  then  $P(X > 2) \sim N(0, 8)$
  - Transform  $X \to Z_x$  then  $P(Z_X > 2/\sqrt{8}) \sim N(0,1)$
  - Calculation of  $1 P(Z_X < 2/\sqrt{8})$  in R below:
- 1 pnorm(2/sqrt(8))

#### ## [1] 0.2397501

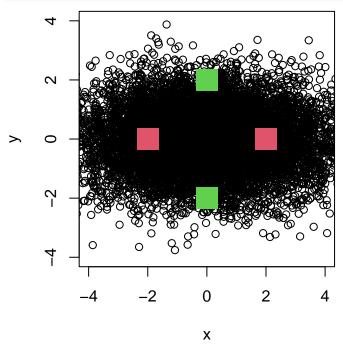
- b) Calculate  $P(|Y_1 Y_2| > 2)$ 
  - Let  $Y = Y_1 Y_2$  then  $P(Y > 2) \sim N(0, 2)$
  - Transform  $Y \to Z_y$  then  $P(Z_Y > 2/\sqrt{2}) \sim N(0,1)$
  - Calculation of  $1 P(Z_Y < 2/\sqrt{2})$  in R below:
- 1 pnorm(2/sqrt(2))

## [1] 0.0786496

### Q1 continued

c) Estimate the two probabilities using simulations. The code in the previous page generates 1000 random samples. Change the sample size from n=1000 to n=10000 and then estimate the two probabilities. To do that, you need to examine all pairs of data points and then calculate the proportion of pairs satisfying a certain condition.

```
set.seed(20230404)
n <- 10000 # number of samples
# bivariate normal random sample parameters
bivariate_mu \leftarrow c(0,0)
cov_matrix \leftarrow matrix(c(4,0,0,1),2,2)
# bivariate normal sample
sample <- mvrnorm(n=n, mu=bivariate_mu, Sigma=cov_matrix)</pre>
# Extract the X values, X \sim N(0, 4)
x <- sample[, 1]
# Extract the Y values, X \sim N(0, 1)
y <- sample[, 2]
# plot
par(pty="s") #to make sure the shape of figure is a square
sample %>%
  plot(xlab="x", ylab="y", xlim=c(-4,4), ylim=c(-4,4))
points(x=c(-2, 0, 0, 2), y=c(0, -2, 2, 0), pch=15,
       col=c(2,3,3,2),cex=3)
```



Calculate P(X > 2) and calculate P(Y > 2)

```
# Calculate the proportion of pairs satisfying X > 2
mean(x > 2)

## [1] 0.1589
# Calculate the proportion of pairs satisfying Y > 2
mean(y > 2)
```