

# Class17

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## Class 17

```
library(tximport)
folders <- dir(pattern="SRR21568*")
samples <- sub("_quant", "", folders)
files <- file.path( folders, "abundance.h5" )
names(files) <- samples
```

```
txi.kallisto <- tximport(files, type = "kallisto", txOut = TRUE)
```

1 2 3 4

```
head(txi.kallisto$counts)
```

	SRR2156848	SRR2156849	SRR2156850	SRR2156851
ENST00000539570	0	0	0.00000	0
ENST00000576455	0	0	2.62037	0
ENST00000510508	0	0	0.00000	0
ENST00000474471	0	1	1.00000	0
ENST00000381700	0	0	0.00000	0
ENST00000445946	0	0	0.00000	0

```
colSums(txi.kallisto$counts)
```

SRR2156848	SRR2156849	SRR2156850	SRR2156851
2563611	2600800	2372309	2111474

```
sum(rowSums(txi.kallisto$counts)>0)
```

```
[1] 94561
```

```
to.keep <- rowSums(txi.kallisto$counts) > 0  
kset.nonzero <- txi.kallisto$counts[to.keep,]  
keep2 <- apply(kset.nonzero,1,sd)>0  
x <- kset.nonzero[keep2,]
```

Principle Component Analysis

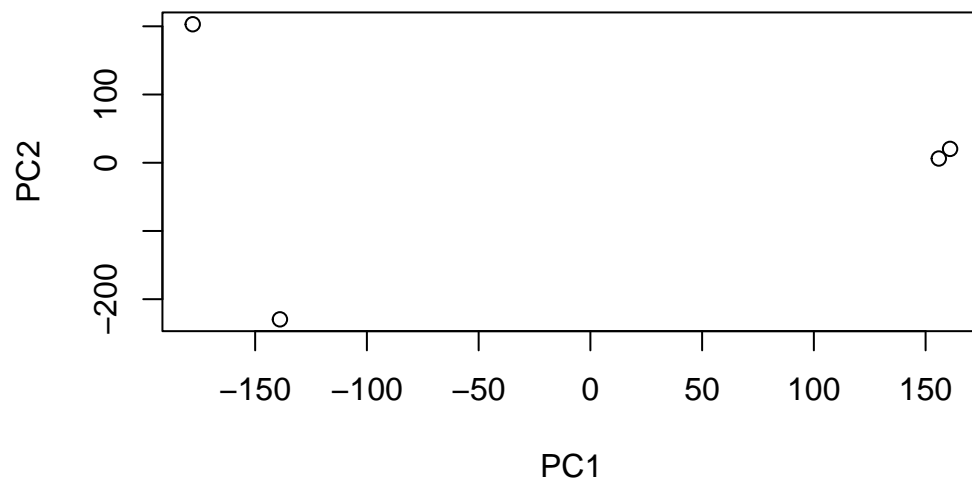
```
pca <- prcomp(t(x), scale=TRUE)  
summary(pca)
```

Importance of components:

	PC1	PC2	PC3	PC4
Standard deviation	183.6379	177.3605	171.3020	1e+00
Proportion of Variance	0.3568	0.3328	0.3104	1e-05
Cumulative Proportion	0.3568	0.6895	1.0000	1e+00

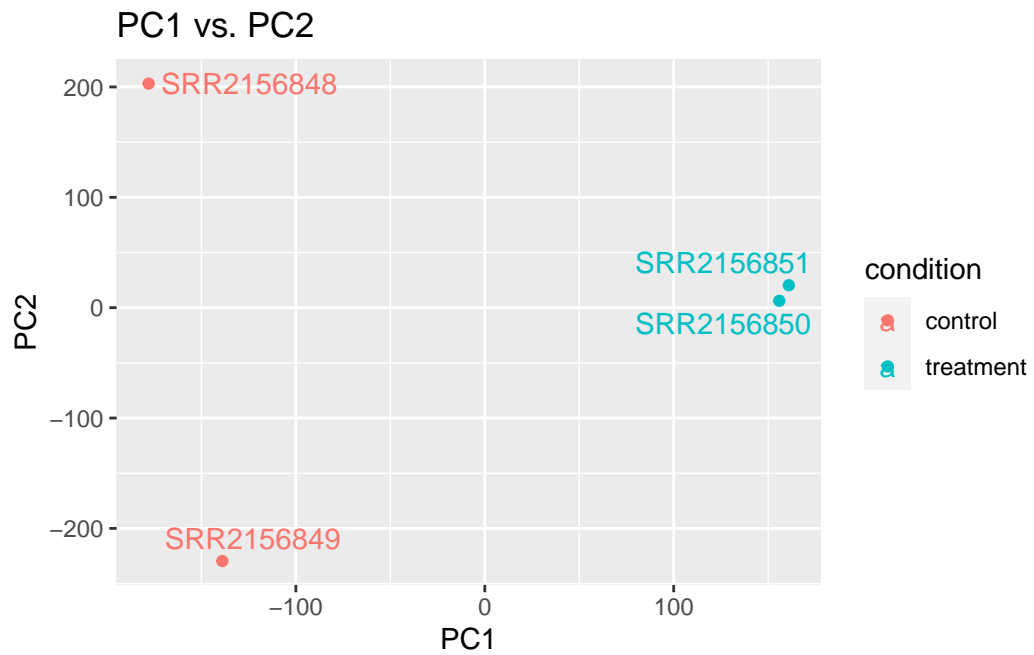
Plots

```
plot(pca$x[,1], pca$x[,2], xlab="PC1", ylab="PC2")
```



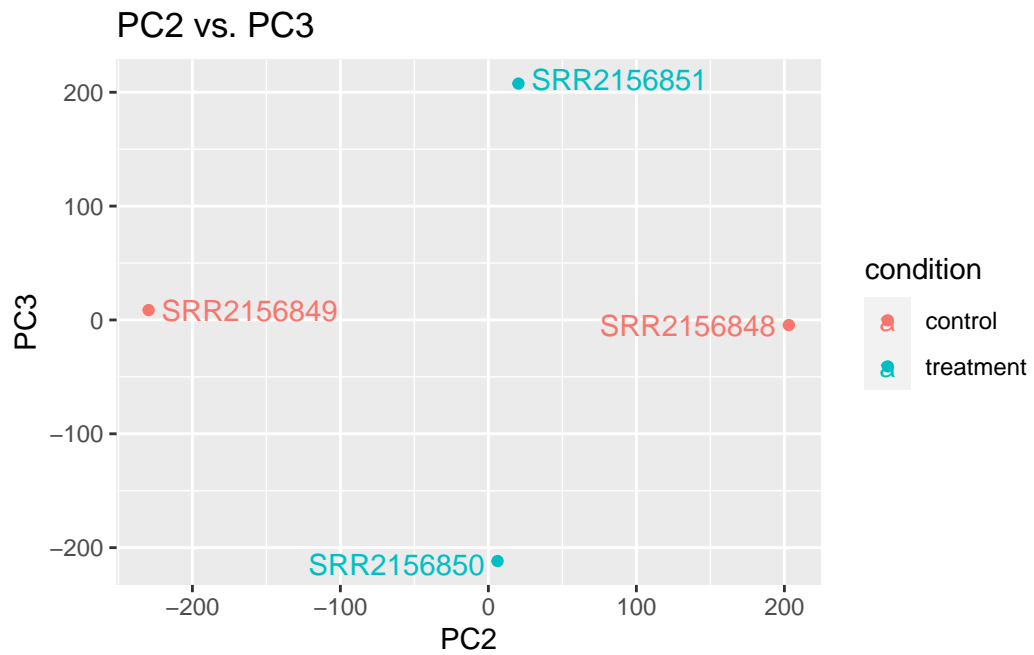
PC1 vs. PC2

```
library(ggplot2)
library(ggrepel)
pcares <- as.data.frame(pca$x)
condition = factor(c("control", "control", "treatment", "treatment"))
ggplot(pcares) + aes(x=PC1, y=PC2, col=condition) + geom_point() +
  geom_text_repel(label=rownames(pcares)) + labs(title= "PC1 vs. PC2")
```



PC2 vs. PC3

```
ggplot(pcares)+ aes(x=PC2,y=PC3, col=condition) + geom_point() +  
  geom_text_repel(label=rownames(pcares)) +labs(title= "PC2 vs. PC3")
```



PC1 vs. PC3

```
ggplot(pcares)+ aes(x=PC1,y=PC3, col=condition) + geom_point() +  
  geom_text_repel(label=rownames(pcares))+labs(title= "PC1 vs. PC3")
```

