

Coronavirus Liver and Blood Capillary Samples

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These samples are the headers added from three Gene Expression Omnibus studies at

- ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE89166
- ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE89160
- ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE100509

The first two studies are part of the same study that used human liver tumor samples in vitro to compare the effects of the coronavirus over time. The third study used human microvascular blood capillaries in vitro to study the effects of the coronavirus over time.

In the first two studies that used the liver tumor samples to examine the effects of the coronavirus in vitro, there were four groups inoculated or treated with the active coronavirus and four groups not inoculated with the active coronavirus, and two samples that were treated with heat inactivated coronavirus, and two samples that were treated with active coronavirus and IL-1alpha to see the gene expression changes over one hour's time.

In the the third study that used blood capillaries, there were five samples followed over a 0,12,24,36, and 48 hour time intervals in groups A,B,C,D, and E that compared the time interval values of screening for changes in microarray analysis with a mock group of the same.

This following data is the data of all genes in common between these three studies, cleaned to remove missing values and with the attached gene symbols from the GEO platform for the probe IDs.

```
both <- read.csv('both_clean_liver_capillary_CoV.csv', sep=',', header=TRUE,
                 na.strings=c('', ' '))

dim(both)

## [1] 21754    63

colnames(both)

## [1] "GENE_SYMBOL"
## [2] "LiverTumorSamples.GSM2359851_CoV1"
## [3] "LiverTumorSamples.GSM2359853_CoV2"
## [4] "LiverTumorSamples.GSM2359910_CoV3"
## [5] "LiverTumorSamples.GSM2359913_CoV4"
## [6] "LiverTumorSamples.GSM2359850_ctrl1"
## [7] "LiverTumorSamples.GSM2359852_ctrl2"
```

```
## [8] "LiverTumorSamples.GSM2359911_ctrl3"
## [9] "LiverTumorSamples.GSM2359914_ctrl4"
## [10] "LiverTumorSamples.GSM2359912_IL1"
## [11] "LiverTumorSamples.GSM2359917_IL2"
## [12] "LiverTumorSamples.GSM2359915_inactiveHeatCoV1"
## [13] "LiverTumorSamples.GSM2359916_inactiveHeatCoV2"
## [14] "capillarySamples.GSM2685693_MERS_CoV_0hr_A"
## [15] "capillarySamples.GSM2685694_MERS_CoV_0hr_B"
## [16] "capillarySamples.GSM2685695_MERS_CoV_0hr_C"
## [17] "capillarySamples.GSM2685696_MERS_CoV_0hr_D"
## [18] "capillarySamples.GSM2685697_MERS_CoV_0hr_E"
## [19] "capillarySamples.GSM2685698_ctrl_0hr_A"
## [20] "capillarySamples.GSM2685699_ctrl_0hr_B"
## [21] "capillarySamples.GSM2685700_ctrl_0hr_C"
## [22] "capillarySamples.GSM2685701_ctrl_0hr_D"
## [23] "capillarySamples.GSM2685702_ctrl_0hr_E"
## [24] "capillarySamples.GSM2685703_MERS_CoV_12hr_A"
## [25] "capillarySamples.GSM2685704_MERS_CoV_12hr_B"
## [26] "capillarySamples.GSM2685705_MERS_CoV_12hr_C"
## [27] "capillarySamples.GSM2685706_MERS_CoV_12hr_D"
## [28] "capillarySamples.GSM2685707_MERS_CoV_12hr_E"
## [29] "capillarySamples.GSM2685708_ctrl_12hr_A"
## [30] "capillarySamples.GSM2685709_ctrl_12hr_B"
## [31] "capillarySamples.GSM2685710_ctrl_12hr_C"
## [32] "capillarySamples.GSM2685711_ctrl_12hr_D"
## [33] "capillarySamples.GSM2685712_ctrl_12hr_E"
## [34] "capillarySamples.GSM2685713_MERS_CoV_24hr_A"
## [35] "capillarySamples.GSM2685714_MERS_CoV_24hr_B"
## [36] "capillarySamples.GSM2685715_MERS_CoV_24hr_C"
## [37] "capillarySamples.GSM2685716_MERS_CoV_24hr_D"
## [38] "capillarySamples.GSM2685717_MERS_CoV_24hr_E"
## [39] "capillarySamples.GSM2685718_ctrl_24hr_A"
## [40] "capillarySamples.GSM2685719_ctrl_24hr_B"
## [41] "capillarySamples.GSM2685720_ctrl_24hr_C"
## [42] "capillarySamples.GSM2685721_ctrl_24hr_D"
## [43] "capillarySamples.GSM2685722_ctrl_24hr_E"
## [44] "capillarySamples.GSM2685723_MERS_CoV_36hr_A"
## [45] "capillarySamples.GSM2685724_MERS_CoV_36hr_B"
## [46] "capillarySamples.GSM2685725_MERS_CoV_36hr_C"
## [47] "capillarySamples.GSM2685726_MERS_CoV_36hr_D"
## [48] "capillarySamples.GSM2685727_MERS_CoV_36hr_E"
## [49] "capillarySamples.GSM2685728_ctrl_36hr_A"
## [50] "capillarySamples.GSM2685729_ctrl_36hr_B"
## [51] "capillarySamples.GSM2685730_ctrl_36hr_C"
## [52] "capillarySamples.GSM2685731_ctrl_36hr_D"
## [53] "capillarySamples.GSM2685732_ctrl_36hr_E"
## [54] "capillarySamples.GSM2685733_MERS_CoV_48hr_A"
## [55] "capillarySamples.GSM2685734_MERS_CoV_48hr_B"
## [56] "capillarySamples.GSM2685735_MERS_CoV_48hr_C"
## [57] "capillarySamples.GSM2685736_MERS_CoV_48hr_D"
```

```
## [58] "capillarySamples.GSM2685737_MERS_CoV_48hr_E"
## [59] "capillarySamples.GSM2685738_ctrl_48hr_A"
## [60] "capillarySamples.GSM2685739_ctrl_48hr_B"
## [61] "capillarySamples.GSM2685740_ctrl_48hr_C"
## [62] "capillarySamples.GSM2685741_ctrl_48hr_D"
## [63] "capillarySamples.GSM2685742_ctrl_48hr_E"
```

Lets group the samples that are our columns with descriptive and GEO ID names into their respective groups, get the fold change between the controls from those groups, attach to the original data table, both, as a different names, then order by the genes that have the most fold change then the least fold change. Take the first 100 genes from both lists, combine into one table of 200 genes and the samples with their fold change values ordered, make into a transposed data frame so that the samples are the rows, the stats removed, and the 200 genes are the header columns to save as a machine learning ready file.

Liver tumor study control and CoV treated. Also, the IL-alpha treated and the inactive CoV treated tables are in this code block.

```
names <- both$GENE_SYMBOL

liverCtrl <- both[,c(6:9)]
row.names(liverCtrl) <- names

liverCoV <- both[,c(2:5)]
row.names(liverCoV) <- names

liverIL <- both[,10:11]
row.names(liverIL) <- names

liverIACoV <- both[,12:13]
row.names(liverIACoV) <- names
```

Get the row means of those liver samples groups each.

```
liverCtrl$CtrlMeanLvr <- rowMeans(liverCtrl)
liverCoV$CoVMeanLvr <- rowMeans(liverCoV)
liverIL$ILMeanLvr <- rowMeans(liverIL)
liverIACoV$IACoVMeanLvr <- rowMeans(liverIACoV)
```

Get the fold change values of those states as a ratio to the control group values.

```
fold1 <-
as.data.frame(cbind(liverCtrl$CtrlMeanLvr,liverCoV$CoVMeanLvr,liverIL$ILMeanL
```

```

vr,
      liverIACoV$IACoVMeanLvr))
row.names(fold1) <- names
colnames(fold1) <- c('CtrlMeanLvr', 'CoVMeanLvr', 'ILMeanLvr', 'IACoVMeanLvr')

fold1$FC_CoV <- fold1$CoVMeanLvr/fold1$CtrlMeanLvr
fold1$FC_IL <- fold1$ILMeanLvr/fold1$CtrlMeanLvr
fold1$FC_IACoV <- fold1$IACoVMeanLvr/fold1$CtrlMeanLvr

```

Most expressed in liver samples by fold change of the Coronavirus, inactive CoronaVirus, and the IL-alpha treated Coronavirus as tables.

```

mostCoV <- fold1[order(fold1$FC_CoV, decreasing = TRUE)[0:100],]
mostIL <- fold1[order(fold1$FC_IL, decreasing = TRUE)[0:100],]
mostIACoV <- fold1[order(fold1$FC_IACoV, decreasing = TRUE)[0:100],]

```

Least expressed in liver samples by fold change of the Coronavirus, inactive CoronaVirus, and the IL-alpha treated Coronavirus as tables.

```

leastCoV <- fold1[order(fold1$FC_CoV, decreasing = FALSE)[0:100],]
leastIL <- fold1[order(fold1$FC_IL, decreasing = FALSE)[0:100],]
leastIACoV <- fold1[order(fold1$FC_IACoV, decreasing = FALSE)[0:100],]

```

Gene Expressions with most changes in the liver samples.

```

changes <- rbind(mostCoV,mostIL,mostIACoV,leastCoV,leastIL,leastIACoV)
Changes <- changes[!duplicated(row.names(changes)),]
length(unique(row.names(Changes)))

## [1] 600

```

Get the magnitude of the fold change genes' row means.

```

Changes$MagnitudeFCs <- abs(rowMeans(Changes[,5:7]))

```

Combine this to the samples data for the liver tumor group.

```

Changes$Gene <- row.names(Changes)
combined1 <- merge(both, Changes, by.x='GENE_SYMBOL', by.y='Gene')

combined2 <- combined1[order(combined1$MagnitudeFCs, decreasing=TRUE),]

CombinedLiver <- combined2[c(0:100,354:453),]

```

Machine Learning data for liver samples with 200 genes in the group of most gene expression changes.

```

names1 <- CombinedLiver$GENE_SYMBOL
names2 <- colnames(CombinedLiver)
row.names(CombinedLiver) <- names1

Combo_lvr_ML <- as.data.frame(t(CombinedLiver))

```

```
colnames(Combo_lvr_ML) <- gsub('-', '_', colnames(Combo_lvr_ML))
Combo1 <- Combo_lvr_ML[c(2:63),] #remove stats of fold change values and gene symbol row
```

Lets add a class field called Class_Type to use machine learning on predicting class with these 200 genes and 62 mixed samples of capillary and liver tumor both inoculated with Coronavirus.

```
a <- rep('liver_CoV', 4)
b <- rep('liver_Ctrl', 4)
c <- rep('liver_CoV_IL', 2)
d <- rep('liver_IA_CoV', 2)
e <- rep('capillary_CoV_0hr', 5)
f <- rep('capillary_Ctrl_0hr', 5)
g <- rep('capillary_Cov_12hr', 5)
h <- rep('capillary_Ctrl_12hr', 5)
i <- rep('capillary_Cov_24hr', 5)
j <- rep('capillary_Ctrl_24hr', 5)
k <- rep('capillary_Cov_36hr', 5)
l <- rep('capillary_Ctrl_36hr', 5)
m <- rep('capillary_Cov_48hr', 5)
n <- rep('capillary_Ctrl_48hr', 5)
```

```
type <- as.data.frame(c(a,b,c,d,e,f,g,h,i,j,k,l,m,n))
colnames(type) <- 'Class_Type'
row.names(type) <- row.names(Combo1)
type
```

##	Class_Type
## LiverTumorSamples.GSM2359851_CoV1	liver_CoV
## LiverTumorSamples.GSM2359853_CoV2	liver_CoV
## LiverTumorSamples.GSM2359910_CoV3	liver_CoV
## LiverTumorSamples.GSM2359913_CoV4	liver_CoV
## LiverTumorSamples.GSM2359850_ctrl11	liver_Ctrl
## LiverTumorSamples.GSM2359852_ctrl12	liver_Ctrl
## LiverTumorSamples.GSM2359911_ctrl13	liver_Ctrl
## LiverTumorSamples.GSM2359914_ctrl14	liver_Ctrl
## LiverTumorSamples.GSM2359912_IL1	liver_CoV_IL
## LiverTumorSamples.GSM2359917_IL2	liver_CoV_IL
## LiverTumorSamples.GSM2359915_inactiveHeatCoV1	liver_IA_CoV
## LiverTumorSamples.GSM2359916_inactiveHeatCoV2	liver_IA_CoV
## capillarySamples.GSM2685693_MERS_CoV_0hr_A	capillary_CoV_0hr
## capillarySamples.GSM2685694_MERS_CoV_0hr_B	capillary_CoV_0hr
## capillarySamples.GSM2685695_MERS_CoV_0hr_C	capillary_CoV_0hr
## capillarySamples.GSM2685696_MERS_CoV_0hr_D	capillary_CoV_0hr
## capillarySamples.GSM2685697_MERS_CoV_0hr_E	capillary_CoV_0hr
## capillarySamples.GSM2685698_ctrl_0hr_A	capillary_Ctrl_0hr
## capillarySamples.GSM2685699_ctrl_0hr_B	capillary_Ctrl_0hr
## capillarySamples.GSM2685700_ctrl_0hr_C	capillary_Ctrl_0hr

```
## capillarySamples.GSM2685701_ctrl_0hr_D      capillary_Ctrl_0hr
## capillarySamples.GSM2685702_ctrl_0hr_E      capillary_Ctrl_0hr
## capillarySamples.GSM2685703_MERS_CoV_12hr_A  capillary_Cov_12hr
## capillarySamples.GSM2685704_MERS_CoV_12hr_B  capillary_Cov_12hr
## capillarySamples.GSM2685705_MERS_CoV_12hr_C  capillary_Cov_12hr
## capillarySamples.GSM2685706_MERS_CoV_12hr_D  capillary_Cov_12hr
## capillarySamples.GSM2685707_MERS_CoV_12hr_E  capillary_Cov_12hr
## capillarySamples.GSM2685708_ctrl_12hr_A      capillary_Ctrl_12hr
## capillarySamples.GSM2685709_ctrl_12hr_B      capillary_Ctrl_12hr
## capillarySamples.GSM2685710_ctrl_12hr_C      capillary_Ctrl_12hr
## capillarySamples.GSM2685711_ctrl_12hr_D      capillary_Ctrl_12hr
## capillarySamples.GSM2685712_ctrl_12hr_E      capillary_Ctrl_12hr
## capillarySamples.GSM2685713_MERS_CoV_24hr_A  capillary_Cov_24hr
## capillarySamples.GSM2685714_MERS_CoV_24hr_B  capillary_Cov_24hr
## capillarySamples.GSM2685715_MERS_CoV_24hr_C  capillary_Cov_24hr
## capillarySamples.GSM2685716_MERS_CoV_24hr_D  capillary_Cov_24hr
## capillarySamples.GSM2685717_MERS_CoV_24hr_E  capillary_Cov_24hr
## capillarySamples.GSM2685718_ctrl_24hr_A      capillary_Ctrl_24hr
## capillarySamples.GSM2685719_ctrl_24hr_B      capillary_Ctrl_24hr
## capillarySamples.GSM2685720_ctrl_24hr_C      capillary_Ctrl_24hr
## capillarySamples.GSM2685721_ctrl_24hr_D      capillary_Ctrl_24hr
## capillarySamples.GSM2685722_ctrl_24hr_E      capillary_Ctrl_24hr
## capillarySamples.GSM2685723_MERS_CoV_36hr_A  capillary_Cov_36hr
## capillarySamples.GSM2685724_MERS_CoV_36hr_B  capillary_Cov_36hr
## capillarySamples.GSM2685725_MERS_CoV_36hr_C  capillary_Cov_36hr
## capillarySamples.GSM2685726_MERS_CoV_36hr_D  capillary_Cov_36hr
## capillarySamples.GSM2685727_MERS_CoV_36hr_E  capillary_Cov_36hr
## capillarySamples.GSM2685728_ctrl_36hr_A      capillary_Ctrl_36hr
## capillarySamples.GSM2685729_ctrl_36hr_B      capillary_Ctrl_36hr
## capillarySamples.GSM2685730_ctrl_36hr_C      capillary_Ctrl_36hr
## capillarySamples.GSM2685731_ctrl_36hr_D      capillary_Ctrl_36hr
## capillarySamples.GSM2685732_ctrl_36hr_E      capillary_Ctrl_36hr
## capillarySamples.GSM2685733_MERS_CoV_48hr_A  capillary_Cov_48hr
## capillarySamples.GSM2685734_MERS_CoV_48hr_B  capillary_Cov_48hr
## capillarySamples.GSM2685735_MERS_CoV_48hr_C  capillary_Cov_48hr
## capillarySamples.GSM2685736_MERS_CoV_48hr_D  capillary_Cov_48hr
## capillarySamples.GSM2685737_MERS_CoV_48hr_E  capillary_Cov_48hr
## capillarySamples.GSM2685738_ctrl_48hr_A      capillary_Ctrl_48hr
## capillarySamples.GSM2685739_ctrl_48hr_B      capillary_Ctrl_48hr
## capillarySamples.GSM2685740_ctrl_48hr_C      capillary_Ctrl_48hr
## capillarySamples.GSM2685741_ctrl_48hr_D      capillary_Ctrl_48hr
## capillarySamples.GSM2685742_ctrl_48hr_E      capillary_Ctrl_48hr
```

```
Combo2 <- cbind(type, Combo1)
Combo2[1:10, 1:5]
```

##	Class_Type	NEURL3	DUSP1
## LiverTumorSamples.GSM2359851_CoV1	liver_CoV	1429.61750	8491.40875
## LiverTumorSamples.GSM2359853_CoV2	liver_CoV	190.21750	2219.85650
## LiverTumorSamples.GSM2359910_CoV3	liver_CoV	10.004148	11.494585

```
## LiverTumorSamples.GSM2359913_CoV4      liver_CoV      11.245589      12.898250
## LiverTumorSamples.GSM2359850_ctrl1      liver_Ctrl      34.57000      228.18775
## LiverTumorSamples.GSM2359852_ctrl12     liver_Ctrl      17.25750      216.08550
## LiverTumorSamples.GSM2359911_ctrl13     liver_Ctrl      3.708157      7.184185
## LiverTumorSamples.GSM2359914_ctrl14     liver_Ctrl      4.757780      7.113854
## LiverTumorSamples.GSM2359912_IL1       liver_CoV_IL      4.879242      9.576161
## LiverTumorSamples.GSM2359917_IL2       liver_CoV_IL      5.1138565     9.5527540
##                                           ATF3             PCLO
## LiverTumorSamples.GSM2359851_CoV1      3608.28250      17.74792
## LiverTumorSamples.GSM2359853_CoV2      974.76937      613.28583
## LiverTumorSamples.GSM2359910_CoV3       8.337322      3.355859
## LiverTumorSamples.GSM2359913_CoV4       9.441972      2.741117
## LiverTumorSamples.GSM2359850_ctrl1      108.52250      13.80667
## LiverTumorSamples.GSM2359852_ctrl12     97.05125      12.88750
## LiverTumorSamples.GSM2359911_ctrl13     5.373051      3.904719
## LiverTumorSamples.GSM2359914_ctrl14     5.513552      3.621765
## LiverTumorSamples.GSM2359912_IL1       6.500059      3.653289
## LiverTumorSamples.GSM2359917_IL2       6.6111744     3.7909157
```

Write this ML ready file to csv.

```
write.csv(Combo2, 'ML_ready_CoV_14_classes.csv', row.names=TRUE)
```

Make a separate ML ready file with a smaller set of classes to classify by liver or capillary and control or CoronaVirus

```
a <- rep('liver', 4)
b <- rep('liver', 4)
c <- rep('liver', 2)
d <- rep('liver', 2)
e <- rep('capillary', 5)
f <- rep('capillary', 5)
g <- rep('capillary', 5)
h <- rep('capillary', 5)
i <- rep('capillary', 5)
j <- rep('capillary', 5)
k <- rep('capillary', 5)
l <- rep('capillary', 5)
m <- rep('capillary', 5)
n <- rep('capillary', 5)

type <- as.data.frame(c(a,b,c,d,e,f,g,h,i,j,k,l,m,n))
colnames(type) <- 'Class_Type'
row.names(type) <- row.names(Combo1)

Combo3 <- cbind(type, Combo1)

Combo3[1:10, 1:5]

##                               Class_Type      NEURL3      DUSP1
## LiverTumorSamples.GSM2359851_CoV1      liver      1429.61750      8491.40875
```



```
## LiverTumorSamples.GSM2359853_CoV2      liver  190.21750  2219.85650
## LiverTumorSamples.GSM2359910_CoV3      liver  10.004148  11.494585
## LiverTumorSamples.GSM2359913_CoV4      liver  11.245589  12.898250
## LiverTumorSamples.GSM2359850_ctrl11    liver  34.57000  228.18775
## LiverTumorSamples.GSM2359852_ctrl12    liver  17.25750  216.08550
## LiverTumorSamples.GSM2359911_ctrl13    liver  3.708157   7.184185
## LiverTumorSamples.GSM2359914_ctrl14    liver  4.757780   7.113854
## LiverTumorSamples.GSM2359912_IL1      liver  4.879242   9.576161
## LiverTumorSamples.GSM2359917_IL2      liver  5.1138565  9.5527540
##                                     ATF3      PCLO
## LiverTumorSamples.GSM2359851_CoV1    3608.28250  17.74792
## LiverTumorSamples.GSM2359853_CoV2    974.76937  613.28583
## LiverTumorSamples.GSM2359910_CoV3     8.337322   3.355859
## LiverTumorSamples.GSM2359913_CoV4     9.441972   2.741117
## LiverTumorSamples.GSM2359850_ctrl11  108.52250  13.80667
## LiverTumorSamples.GSM2359852_ctrl12   97.05125  12.88750
## LiverTumorSamples.GSM2359911_ctrl13   5.373051   3.904719
## LiverTumorSamples.GSM2359914_ctrl14   5.513552   3.621765
## LiverTumorSamples.GSM2359912_IL1     6.500059   3.653289
## LiverTumorSamples.GSM2359917_IL2     6.6111744  3.7909157
```

```
write.csv(Combo3, 'ML_ready_CoV_2_classes.csv', row.names=TRUE)
```

```
a <- rep('CoV', 4)
b <- rep('Ctrl', 4)
c <- rep('CoV_IL', 2)
d <- rep('IA_CoV', 2)
e <- rep('CoV', 5)
f <- rep('Ctrl', 5)
g <- rep('Cov', 5)
h <- rep('Ctrl', 5)
i <- rep('Cov', 5)
j <- rep('Ctrl', 5)
k <- rep('Cov', 5)
l <- rep('Ctrl', 5)
m <- rep('Cov', 5)
n <- rep('Ctrl', 5)
```

```
type <- as.data.frame(c(a,b,c,d,e,f,g,h,i,j,k,l,m,n))
colnames(type) <- 'Class_Type'
row.names(type) <- row.names(Combo1)
```

```
Combo4 <- cbind(type, Combo1)
```

```
Combo4[1:10,1:5]
```

```
##                                     Class_Type      NEURL3      DUSP1
## LiverTumorSamples.GSM2359851_CoV1      CoV  1429.61750  8491.40875
## LiverTumorSamples.GSM2359853_CoV2      CoV  190.21750  2219.85650
## LiverTumorSamples.GSM2359910_CoV3      CoV  10.004148  11.494585
```



```
## LiverTumorSamples.GSM2359913_CoV4      CoV  11.245589  12.898250
## LiverTumorSamples.GSM2359850_ctrl1      Ctrl  34.57000  228.18775
## LiverTumorSamples.GSM2359852_ctrl12     Ctrl  17.25750  216.08550
## LiverTumorSamples.GSM2359911_ctrl13     Ctrl  3.708157   7.184185
## LiverTumorSamples.GSM2359914_ctrl14     Ctrl  4.757780   7.113854
## LiverTumorSamples.GSM2359912_IL1       CoV_IL  4.879242   9.576161
## LiverTumorSamples.GSM2359917_IL2       CoV_IL  5.1138565  9.5527540
##                                           ATF3      PCLO
## LiverTumorSamples.GSM2359851_CoV1  3608.28250  17.74792
## LiverTumorSamples.GSM2359853_CoV2   974.76937  613.28583
## LiverTumorSamples.GSM2359910_CoV3     8.337322   3.355859
## LiverTumorSamples.GSM2359913_CoV4     9.441972   2.741117
## LiverTumorSamples.GSM2359850_ctrl1    108.52250  13.80667
## LiverTumorSamples.GSM2359852_ctrl12   97.05125  12.88750
## LiverTumorSamples.GSM2359911_ctrl13    5.373051   3.904719
## LiverTumorSamples.GSM2359914_ctrl14    5.513552   3.621765
## LiverTumorSamples.GSM2359912_IL1      6.500059   3.653289
## LiverTumorSamples.GSM2359917_IL2     6.6111744  3.7909157
```

```
write.csv(Combo4, 'ML_ready_CoV_4_classes.csv', row.names=TRUE)
```

We didn't do any fold change or stat measures on the capillary samples, but we can plot them by using ggplot2 and group the sets by timed intervals for each group A through E and picking a handful of genes to compare over the 0,12,24,36, and 48 hour time intervals for the control group and the Coronavirus inoculated groups.

When the values are a ratio like this, it is easier to see the larger changes as in 9 compared to a low change like 0.0005, but this just means that compared to the control samples the inoculated Coronavirus had 9 times the gene expression values or had downregulated or suppressed gene expression values to 1/5000th the amount of the normal range of gene expression values respectively. ***

It makes sense to use some genes we already know have a higher magnitude of change, and we have a column for that in the CombinedLiver table called MagnitudeFCs that was already sorted from largest to smallest when made. We'll just select the first five of those genes to compare in these capillary samples over time.

```
mostChanged <- CombinedLiver[1:5,c(1,71)]
mostSuppressed <- CombinedLiver[196:200,c(1,71)]
row.names(mostChanged)

## [1] "NEURL3" "DUSP1" "ATF3" "PCLO" "LHB"

row.names(mostSuppressed)
```

```

## [1] "RASSF7"          "LOC100335030" "C2orf78"          "DEFB1"           "ZNF610"

capillary <- merge(mostChanged, CombinedLiver, by.x='GENE_SYMBOL',
by.y='GENE_SYMBOL')
capillary1 <- merge(mostSuppressed, CombinedLiver, by.x='GENE_SYMBOL',
by.y='GENE_SYMBOL')
capillaries <- rbind(capillary, capillary1)
Capillaries <- capillaries[,c(1,15:64)]
row.names(Capillaries) <- Capillaries$GENE_SYMBOL

Capillaries2 <- as.data.frame(t(Capillaries))
Capillaries2 <- Capillaries2[-1,]
row.names(Capillaries2) <-
gsub('capillarySamples.', '', row.names(Capillaries2))
row.names(Capillaries2) <- gsub('GSM[0-9][0-9][0-9][0-9][0-9][0-9][0-9]_', '', row.names(Capillaries2))
row.names(Capillaries2) <- gsub('MERS_', '', row.names(Capillaries2))

CoV <- grep('CoV', row.names(Capillaries2))
ctrl <- grep('ctrl', row.names(Capillaries2))

Capillaries2$Class <- 'CoV or ctrl'

Capillaries2[CoV,11] <- 'Coronavirus'
Capillaries2[ctrl,11] <- 'control'

A <- grep('_A', row.names(Capillaries2))
B <- grep('_B', row.names(Capillaries2))
C <- grep('_C', row.names(Capillaries2))
D <- grep('_D', row.names(Capillaries2))
E <- grep('_E', row.names(Capillaries2))

Capillaries2$Group <- 'group'

Capillaries2[A,12] <- 'A'
Capillaries2[B,12] <- 'B'
Capillaries2[C,12] <- 'C'
Capillaries2[D,12] <- 'D'
Capillaries2[E,12] <- 'E'

hr0 <- grep('0hr', row.names(Capillaries2))
hr12 <- grep('12hr', row.names(Capillaries2))
hr24 <- grep('24hr', row.names(Capillaries2))
hr36 <- grep('36hr', row.names(Capillaries2))
hr48 <- grep('48hr', row.names(Capillaries2))

Capillaries2$TimeInterval <- 'time'

Capillaries2[hr0,13] <- '0 hr'

```

```

Capillaries2[hr12,13] <- '12 hr'
Capillaries2[hr24,13] <- '24 hr'
Capillaries2[hr36,13] <- '36 hr'
Capillaries2[hr48,13] <- '48 hr'

```

Capillaries2

##		ATF3	DUSP1	LHB	NEURL3	PCLO	C2orf78
##	CoV_0hr_A	10.290996	13.901505	9.093671	7.498965	6.468670	5.936451
##	CoV_0hr_B	9.780412	13.852933	9.128886	7.446590	6.476326	6.023280
##	CoV_0hr_C	9.574148	13.740905	8.662565	7.495092	6.463702	5.909934
##	CoV_0hr_D	9.848204	13.863910	8.729845	7.364854	6.572579	5.986079
##	CoV_0hr_E	10.114265	13.968982	8.738079	7.436675	6.625125	5.862801
##	ctrl_0hr_A	10.173291	14.398733	8.776979	7.267773	6.410249	6.016610
##	ctrl_0hr_B	10.132629	14.237192	9.610252	7.015544	6.467982	5.931786
##	ctrl_0hr_C	10.308477	14.382699	9.557514	7.440873	6.478339	5.853411
##	ctrl_0hr_D	9.888505	14.404923	9.602914	7.102155	6.489396	5.855096
##	ctrl_0hr_E	9.892730	14.120833	9.577233	6.936543	6.417981	5.942510
##	CoV_12hr_A	9.819186	11.303627	10.795022	12.768140	6.256646	5.865277
##	CoV_12hr_B	10.011539	11.577456	11.006393	12.402078	6.327304	5.922540
##	CoV_12hr_C	9.783105	11.626722	11.000187	12.532051	6.322820	5.988353
##	CoV_12hr_D	9.849858	11.692725	10.938022	12.668202	6.261566	5.901327
##	CoV_12hr_E	9.617070	12.220997	10.553798	12.273302	6.333485	5.898912
##	ctrl_12hr_A	7.825226	11.488735	9.286872	6.829080	6.528833	5.954890
##	ctrl_12hr_B	7.872041	11.710908	8.845744	6.784278	6.532492	5.862801
##	ctrl_12hr_C	7.989155	11.468072	9.267328	6.908973	6.489589	5.973506
##	ctrl_12hr_D	8.028735	11.407001	9.820556	6.615987	6.412667	5.928711
##	ctrl_12hr_E	8.154875	11.042038	9.741747	6.833504	6.431864	5.862305
##	CoV_24hr_A	12.356029	14.922819	10.081784	13.713997	6.330955	5.981748
##	CoV_24hr_B	12.393863	14.762759	10.125434	13.516154	6.376025	5.938335
##	CoV_24hr_C	12.347139	15.286795	9.988253	13.718699	6.398037	6.047165
##	CoV_24hr_D	12.280955	15.019195	9.580647	13.576554	6.441270	6.099102
##	CoV_24hr_E	12.187980	14.989084	9.508540	13.545156	6.428447	6.044217
##	ctrl_24hr_A	7.914153	12.020454	8.926344	6.410178	6.709108	6.033401
##	ctrl_24hr_B	8.179651	11.901618	9.159004	6.560892	6.642224	5.909934
##	ctrl_24hr_C	8.062523	11.831523	9.275883	6.568645	6.626875	5.938570
##	ctrl_24hr_D	8.045721	11.806023	9.273915	6.631753	6.618130	5.997365
##	ctrl_24hr_E	8.204591	11.942950	9.125971	6.562419	6.697083	5.919444
##	CoV_36hr_A	12.318180	14.508819	9.861903	13.561144	6.548994	6.087523
##	CoV_36hr_B	12.173687	14.540567	9.802688	13.549071	6.644295	5.999396
##	CoV_36hr_C	12.163530	14.579001	9.872176	13.449660	6.515321	5.933150
##	CoV_36hr_D	12.142847	14.514722	9.982218	13.712803	6.671993	6.001650
##	CoV_36hr_E	12.297080	14.374295	10.201176	13.521456	6.372439	6.051522
##	ctrl_36hr_A	7.940682	11.742312	9.321322	6.730998	6.678982	6.220477
##	ctrl_36hr_B	7.783378	11.751105	9.056790	6.621403	6.643658	6.125131
##	ctrl_36hr_C	7.810544	11.798201	9.151894	6.822530	6.618794	6.092605
##	ctrl_36hr_D	7.731982	11.876269	9.024215	6.685524	6.629717	6.098892
##	ctrl_36hr_E	7.734450	11.888111	8.922796	6.630443	6.673160	6.028326
##	CoV_48hr_A	11.205349	13.523040	9.679922	12.333937	6.529574	6.049019
##	CoV_48hr_B	11.253111	13.511350	9.912659	12.481150	6.549794	5.980606

##	CoV_48hr_C	10.944238	13.316529	9.850259	12.260608	6.708351	6.203350	
##	CoV_48hr_D	10.983900	13.358680	9.899124	11.961065	6.599227	6.142515	
##	CoV_48hr_E	11.255318	13.367467	10.184716	12.498831	6.679909	6.017946	
##	ctrl_48hr_A	7.932607	12.174103	9.024090	6.757483	6.552106	6.016387	
##	ctrl_48hr_B	7.668012	12.449589	8.985262	6.882334	6.700790	5.961156	
##	ctrl_48hr_C	7.755264	12.365436	9.031925	6.844960	6.455577	6.003225	
##	ctrl_48hr_D	7.725891	12.415529	8.940948	6.872043	6.533961	5.986534	
##	ctrl_48hr_E	7.656200	12.455840	8.994880	6.868035	6.479109	6.091125	
##		DEFB1	LOC100335030	RASSF7	ZNF610			Class Group
##	CoV_0hr_A	7.622403	8.114815	6.822339	7.227249	Coronavirus		A
##	CoV_0hr_B	7.778352	8.270884	6.700971	7.428115	Coronavirus		B
##	CoV_0hr_C	7.352772	8.308822	6.966246	7.481717	Coronavirus		C
##	CoV_0hr_D	7.118591	8.246936	6.875601	7.426565	Coronavirus		D
##	CoV_0hr_E	7.179596	8.214737	6.952108	7.420473	Coronavirus		E
##	ctrl_0hr_A	7.331124	8.187333	6.964110	7.403619	control		A
##	ctrl_0hr_B	7.872561	8.288294	6.910892	7.464914	control		B
##	ctrl_0hr_C	7.584274	8.147007	6.994899	7.269174	control		C
##	ctrl_0hr_D	7.650353	8.185575	7.033153	7.359418	control		D
##	ctrl_0hr_E	7.847725	8.156999	7.056144	7.447995	control		E
##	CoV_12hr_A	8.059324	7.868801	6.594891	8.147592	Coronavirus		A
##	CoV_12hr_B	7.843962	7.735502	6.692292	8.178197	Coronavirus		B
##	CoV_12hr_C	7.985707	7.820541	6.586602	8.121462	Coronavirus		C
##	CoV_12hr_D	7.995350	7.717794	6.571151	8.224417	Coronavirus		D
##	CoV_12hr_E	7.418830	7.832236	6.801301	8.247149	Coronavirus		E
##	ctrl_12hr_A	8.009080	8.069778	6.880927	7.524340	control		A
##	ctrl_12hr_B	8.055858	7.986788	6.895230	7.537055	control		B
##	ctrl_12hr_C	7.952662	7.941232	6.903801	7.575149	control		C
##	ctrl_12hr_D	8.277781	8.181431	6.745836	7.641494	control		D
##	ctrl_12hr_E	8.235038	8.169354	6.554766	7.782781	control		E
##	CoV_24hr_A	7.571666	8.536174	6.747448	8.938377	Coronavirus		A
##	CoV_24hr_B	7.629778	8.509799	6.697222	8.733307	Coronavirus		B
##	CoV_24hr_C	7.260831	8.425157	6.692222	9.019246	Coronavirus		C
##	CoV_24hr_D	7.080690	8.584440	6.757083	9.012278	Coronavirus		D
##	CoV_24hr_E	7.058040	8.736338	6.698612	9.092390	Coronavirus		E
##	ctrl_24hr_A	7.563681	8.155811	6.789623	7.914991	control		A
##	ctrl_24hr_B	7.729397	8.138252	6.704019	7.747434	control		B
##	ctrl_24hr_C	7.946592	8.095640	6.643618	7.788730	control		C
##	ctrl_24hr_D	7.625182	8.078397	6.716886	7.876311	control		D
##	ctrl_24hr_E	7.486914	8.005603	6.680610	7.754600	control		E
##	CoV_36hr_A	8.078504	8.482066	6.721733	8.534288	Coronavirus		A
##	CoV_36hr_B	7.874253	8.583444	6.770072	8.659518	Coronavirus		B
##	CoV_36hr_C	7.801655	8.552423	6.862404	8.572155	Coronavirus		C
##	CoV_36hr_D	7.980233	8.606779	6.846841	8.654873	Coronavirus		D
##	CoV_36hr_E	7.610937	8.589262	6.822339	8.589262	Coronavirus		E
##	ctrl_36hr_A	7.737160	8.018402	6.849095	7.846848	control		A
##	ctrl_36hr_B	7.704646	8.042240	6.771987	7.839025	control		B
##	ctrl_36hr_C	7.652111	8.124825	6.842196	7.974192	control		C
##	ctrl_36hr_D	7.564977	8.045657	6.925022	7.847725	control		D
##	ctrl_36hr_E	7.592967	8.041310	6.908012	7.860321	control		E
##	CoV_48hr_A	7.328972	8.062943	6.505043	7.678934	Coronavirus		A

## CoV_48hr_B	7.354622	8.098275	6.544158	7.742997	Coronavirus	B
## CoV_48hr_C	7.478970	8.024786	6.488793	7.725653	Coronavirus	C
## CoV_48hr_D	7.584650	7.979318	6.643906	7.822518	Coronavirus	D
## CoV_48hr_E	7.345740	8.128902	6.493502	9.374216	Coronavirus	E
## ctrl_48hr_A	7.551584	8.234894	6.812821	7.882805	control	A
## ctrl_48hr_B	7.634577	8.221036	6.918835	8.076742	control	B
## ctrl_48hr_C	7.580736	8.234032	6.851086	8.033925	control	C
## ctrl_48hr_D	7.753263	8.230602	6.913527	7.983884	control	D
## ctrl_48hr_E	7.731435	8.270651	6.878047	7.998738	control	E
##	TimeInterval					
## CoV_0hr_A	0 hr					
## CoV_0hr_B	0 hr					
## CoV_0hr_C	0 hr					
## CoV_0hr_D	0 hr					
## CoV_0hr_E	0 hr					
## ctrl_0hr_A	0 hr					
## ctrl_0hr_B	0 hr					
## ctrl_0hr_C	0 hr					
## ctrl_0hr_D	0 hr					
## ctrl_0hr_E	0 hr					
## CoV_12hr_A	12 hr					
## CoV_12hr_B	12 hr					
## CoV_12hr_C	12 hr					
## CoV_12hr_D	12 hr					
## CoV_12hr_E	12 hr					
## ctrl_12hr_A	12 hr					
## ctrl_12hr_B	12 hr					
## ctrl_12hr_C	12 hr					
## ctrl_12hr_D	12 hr					
## ctrl_12hr_E	12 hr					
## CoV_24hr_A	24 hr					
## CoV_24hr_B	24 hr					
## CoV_24hr_C	24 hr					
## CoV_24hr_D	24 hr					
## CoV_24hr_E	24 hr					
## ctrl_24hr_A	24 hr					
## ctrl_24hr_B	24 hr					
## ctrl_24hr_C	24 hr					
## ctrl_24hr_D	24 hr					
## ctrl_24hr_E	24 hr					
## CoV_36hr_A	36 hr					
## CoV_36hr_B	36 hr					
## CoV_36hr_C	36 hr					
## CoV_36hr_D	36 hr					
## CoV_36hr_E	36 hr					
## ctrl_36hr_A	36 hr					
## ctrl_36hr_B	36 hr					
## ctrl_36hr_C	36 hr					
## ctrl_36hr_D	36 hr					
## ctrl_36hr_E	36 hr					

```
## CoV_48hr_A      48 hr
## CoV_48hr_B      48 hr
## CoV_48hr_C      48 hr
## CoV_48hr_D      48 hr
## CoV_48hr_E      48 hr
## ctrl_48hr_A      48 hr
## ctrl_48hr_B      48 hr
## ctrl_48hr_C      48 hr
## ctrl_48hr_D      48 hr
## ctrl_48hr_E      48 hr

write.csv(Capillaries2, 'FC_10_capillaries_CoV.csv', row.names=TRUE)
```

The above table has 10 genes as the columns with the added Class (Coronavirus or control), Group (A,B,C,D,E), and TimeInterval (0,12,24,36,48 hours) fields to filter by and plot.

Lets make these group tables for the corona virus and see how they compare over time.

```
library(dplyr)

A_group <- filter(Capillaries2, Group=='A' & Class == 'Coronavirus')
B_group <- filter(Capillaries2, Group=='B' & Class == 'Coronavirus')
C_group <- filter(Capillaries2, Group=='C' & Class == 'Coronavirus')
D_group <- filter(Capillaries2, Group=='D' & Class == 'Coronavirus')
E_group <- filter(Capillaries2, Group=='E' & Class == 'Coronavirus')
```

Lets use the tidyr package to put the 10 genes into one Gene field.

```
library(tidyr)
```

We will do this for the A_group table and ignore the Group and Class fields, because we made it only the A group of the Coronavirus class.

```
A_group2 <- A_group[,c(1,3,5,7,9,11:13)]
A_tidy <- gather(A_group2, 'Gene', 'GeneExpression', 1:5)

## Warning: attributes are not identical across measure variables;
## they will be dropped

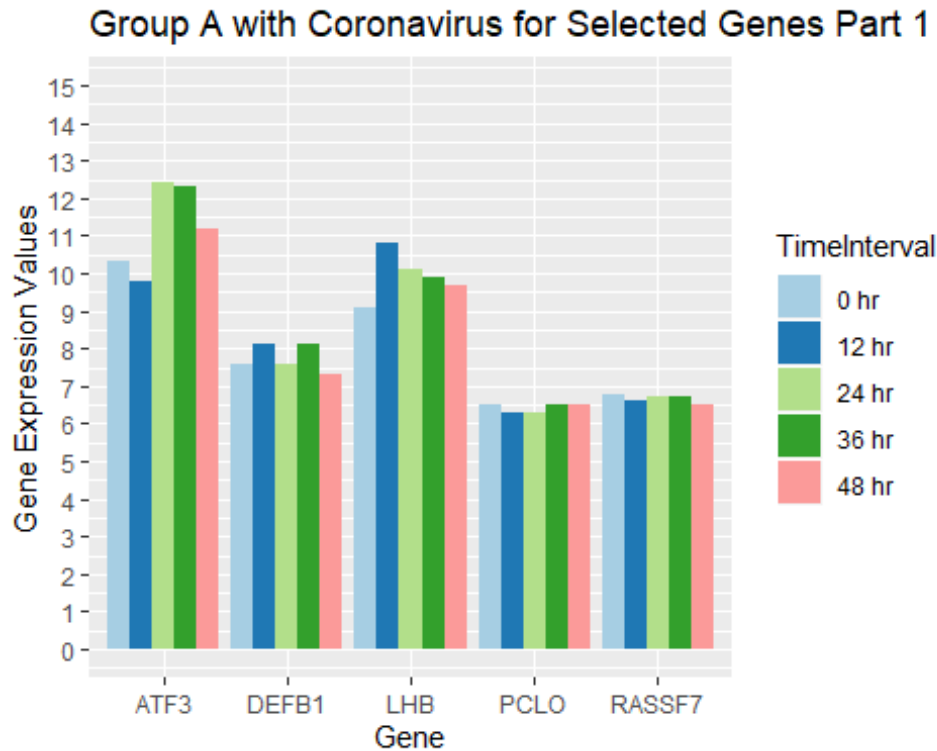
A_tidy$GeneExpression <- round(as.numeric(A_tidy$GeneExpression),1)
A_tidy$TimeInterval <- as.factor(A_tidy$TimeInterval)
A_tidy$Gene <- as.factor(A_tidy$Gene)
A_tidy
```

##	Class	Group	TimeInterval	Gene	GeneExpression
## 1	Coronavirus	A	0 hr	ATF3	10.3
## 2	Coronavirus	A	12 hr	ATF3	9.8
## 3	Coronavirus	A	24 hr	ATF3	12.4
## 4	Coronavirus	A	36 hr	ATF3	12.3
## 5	Coronavirus	A	48 hr	ATF3	11.2

## 6	Coronavirus	A	0 hr	LHB	9.1
## 7	Coronavirus	A	12 hr	LHB	10.8
## 8	Coronavirus	A	24 hr	LHB	10.1
## 9	Coronavirus	A	36 hr	LHB	9.9
## 10	Coronavirus	A	48 hr	LHB	9.7
## 11	Coronavirus	A	0 hr	PCLO	6.5
## 12	Coronavirus	A	12 hr	PCLO	6.3
## 13	Coronavirus	A	24 hr	PCLO	6.3
## 14	Coronavirus	A	36 hr	PCLO	6.5
## 15	Coronavirus	A	48 hr	PCLO	6.5
## 16	Coronavirus	A	0 hr	DEFB1	7.6
## 17	Coronavirus	A	12 hr	DEFB1	8.1
## 18	Coronavirus	A	24 hr	DEFB1	7.6
## 19	Coronavirus	A	36 hr	DEFB1	8.1
## 20	Coronavirus	A	48 hr	DEFB1	7.3
## 21	Coronavirus	A	0 hr	RASSF7	6.8
## 22	Coronavirus	A	12 hr	RASSF7	6.6
## 23	Coronavirus	A	24 hr	RASSF7	6.7
## 24	Coronavirus	A	36 hr	RASSF7	6.7
## 25	Coronavirus	A	48 hr	RASSF7	6.5

```
library(ggplot2)
```

```
ggplot(data = A_tidy, aes(x=Gene, y=GeneExpression, fill=TimeInterval)) +
  geom_bar(stat='identity', position=position_dodge())+
  scale_y_continuous(breaks = seq(0, 15, by=1), limits=c(0,15))+
  scale_fill_brewer(palette='Paired') +
  ggtitle('Group A with Coronavirus for Selected Genes Part 1')+
  xlab('Gene')+
  ylab('Gene Expression Values')
```

The genes above for Part 1 of the group A samples of coronavirus in blood capillaries show some variation in gene expression values for some of these genes that had the most change in the liver tumor samples. Starting at the initial hour up to 48 hours after being inoculated in vitro, there is an increase then decrease for ATF3 and LHB genes, while a decrease then increase close to initial value with PCLO and slightly with RASSF7. For DEFB1, it has a cyclical increase, decrease, increase, then decrease to stabilize closer to the initial gene expression value.

Now let's find the other five genes in the group A set of ten genes found to have the most change in the liver tumor samples, and examined here in the blood capillary samples.

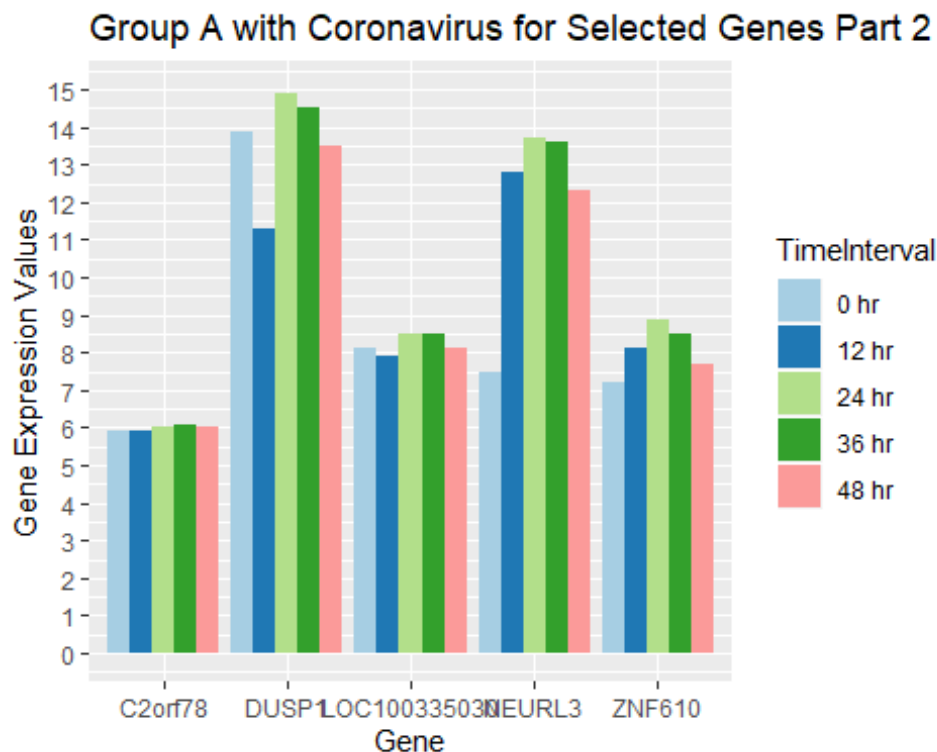
```
A_group3 <- A_group[,c(2,4,6,8,10,11:13)]
A_tidy1 <- gather(A_group3, 'Gene', 'GeneExpression', 1:5)

## Warning: attributes are not identical across measure variables;
## they will be dropped

A_tidy1$GeneExpression <- round(as.numeric(A_tidy1$GeneExpression), 1)
A_tidy1$TimeInterval <- as.factor(A_tidy1$TimeInterval)
A_tidy1$Gene <- as.factor(A_tidy1$Gene)
A_tidy1
```

##	Class	Group	TimeInterval	Gene	GeneExpression
## 1	Coronavirus	A	0 hr	DUSP1	13.9
## 2	Coronavirus	A	12 hr	DUSP1	11.3
## 3	Coronavirus	A	24 hr	DUSP1	14.9
## 4	Coronavirus	A	36 hr	DUSP1	14.5
## 5	Coronavirus	A	48 hr	DUSP1	13.5
## 6	Coronavirus	A	0 hr	NEURL3	7.5
## 7	Coronavirus	A	12 hr	NEURL3	12.8
## 8	Coronavirus	A	24 hr	NEURL3	13.7
## 9	Coronavirus	A	36 hr	NEURL3	13.6
## 10	Coronavirus	A	48 hr	NEURL3	12.3
## 11	Coronavirus	A	0 hr	C2orf78	5.9
## 12	Coronavirus	A	12 hr	C2orf78	5.9
## 13	Coronavirus	A	24 hr	C2orf78	6.0
## 14	Coronavirus	A	36 hr	C2orf78	6.1
## 15	Coronavirus	A	48 hr	C2orf78	6.0
## 16	Coronavirus	A	0 hr	LOC100335030	8.1
## 17	Coronavirus	A	12 hr	LOC100335030	7.9
## 18	Coronavirus	A	24 hr	LOC100335030	8.5
## 19	Coronavirus	A	36 hr	LOC100335030	8.5
## 20	Coronavirus	A	48 hr	LOC100335030	8.1
## 21	Coronavirus	A	0 hr	ZNF610	7.2
## 22	Coronavirus	A	12 hr	ZNF610	8.1
## 23	Coronavirus	A	24 hr	ZNF610	8.9
## 24	Coronavirus	A	36 hr	ZNF610	8.5
## 25	Coronavirus	A	48 hr	ZNF610	7.7

```
ggplot(data = A_tidy1, aes(x=Gene, y=GeneExpression, fill=TimeInterval)) +
  geom_bar(stat='identity', position=position_dodge()) +
  scale_y_continuous(breaks = seq(0, 15, by=1), limits=c(0,15)) +
  scale_fill_brewer(palette='Paired') +
  ggtitle('Group A with Coronavirus for Selected Genes Part 2') +
  xlab('Gene') +
  ylab('Gene Expression Values')
```



The above genes in part 2 of the group A Coronavirus samples over 48 hours, shows that the gene expression values increase up to 24 hours then decrease to 48 hours for most of the genes above.

Lets look back at the platforms and the features removed. The sequence field is an interesting field because it can show copy number variants of genes by the genes that are duplicates at other probes from the samples.

```
Platform13497 <- read.csv('GPL13497-9755-forSequenceFeature-GSE100509.csv',
  sep=',',
  na.strings=c('', ' '), header=TRUE)
Platform16699 <- read.csv('GPL16699-forSequenceFeatureGSE89166_GSE89160.csv',
  sep=',',
  na.strings=c('', ' '), header=TRUE)
```

The features in Platform13497 and the first five listed ID values for this platform:

```
colnames(Platform13497)

## [1] "ID"                "SPOT_ID"           "CONTROL_TYPE"
## [4] "REFSEQ"            "GB_ACC"            "GENE"
## [7] "GENE_SYMBOL"       "GENE_NAME"         "UNIGENE_ID"
```

```
## [10] "ENSEMBL_ID"      "TIGR_ID"          "ACCESSION_STRING"
## [13] "CHROMOSOMAL_LOCATION" "CYTOBAND"         "DESCRIPTION"
## [16] "GO_ID"           "SEQUENCE"
```

```
head(Platform13497$SPOT_ID)[1:5]
```

```
## [1] (+)E1A_r60_1      (+)E1A_r60_3      (+)E1A_r60_a104 (+)E1A_r60_a107
## [5] (+)E1A_r60_a135
## 34184 Levels: (-)3xSLv1 (+)E1A_r60_1 (+)E1A_r60_3 ... NA
```

The features in Platform16699 and the first five listed ID values of that platform:

```
colnames(Platform16699)
```

```
## [1] "ID"          "COL"          "ROW"
## [4] "NAME"        "SPOT_ID"      "CONTROL_TYPE"
## [7] "REFSEQ"      "GB_ACC"       "LOCUSLINK_ID"
## [10] "GENE_SYMBOL" "GENE_NAME"    "UNIGENE_ID"
## [13] "ENSEMBL_ID"  "ACCESSION_STRING" "CHROMOSOMAL_LOCATION"
## [16] "CYTOBAND"    "DESCRIPTION"   "GO_ID"
## [19] "SEQUENCE"
```

```
head(Platform16699$SPOT_ID)[1:5]
```

```
## [1] CONTROL      CONTROL      CONTROL      A_23_P117082 A_33_P3246448
## 50491 Levels: A_19_P00315452 A_19_P00315459 A_19_P00315482 ...
tc|THC2788944
```

Lets keep only the ID (Platform16699) or SPOT_ID (Platform13497), GENE_SYMBOL, DESCRIPTION, and SEQUENCE features of both platforms.

```
P16699 <- Platform16699[,c(1,10,17,19)]
P13497 <- Platform13497[,c(2,7,15,17)]
```

Lets also remove the incomplete cases in both platforms.

```
work16699 <- P16699[complete.cases(P16699),]
work13497 <- P13497[complete.cases(P13497),]
```

Now merge these data sets to their corresponding samples by SPOT_ID. First read in the samples data for each platform and series.

```
GSE89166_89160 <- read.csv('GSE89166_GSE89160.csv', sep=',', na.strings=c('', ' '),
                          header=TRUE)
GSE100509 <- read.csv('GSE100509.csv', sep=',', header=TRUE,
                      na.strings=c('', ' '))
```

Now merge the series data sets to their respective platforms of gene informational meta features.

```
Series100509 <- merge(work13497, GSE100509, by.x='SPOT_ID', by.y='ID_REF')
Series89166_89160 <- merge(work16699, GSE89166_89160, by.x='ID', by.y='ID_REF')
```

Rename the columns of the Series100509 to the 5 groups for each of CoV and Ctrl over 0,12,24,36, and 48 hours.

```
colnames(Series100509)
```

```
## [1] "SPOT_ID" "GENE_SYMBOL"
## [3] "DESCRIPTION" "SEQUENCE"
## [5] "GSM2685693_MERS_CoV_0hr" "GSM2685694_MERS_CoV_0hr"
## [7] "GSM2685695_MERS_CoV_0hr" "GSM2685696_MERS_CoV_0hr"
## [9] "GSM2685697_MERS_CoV_0hr" "GSM2685698_ctrl_0hr"
## [11] "GSM2685699_ctrl_0hr" "GSM2685700_ctrl_0hr"
## [13] "GSM2685701_ctrl_0hr" "GSM2685702_ctrl_0hr"
## [15] "GSM2685703_MERS_CoV_12hr" "GSM2685704_MERS_CoV_12hr"
## [17] "GSM2685705_MERS_CoV_12hr" "GSM2685706_MERS_CoV_12hr"
## [19] "GSM2685707_MERS_CoV_12hr" "GSM2685708_ctrl_12hr"
## [21] "GSM2685709_ctrl_12hr" "GSM2685710_ctrl_12hr"
## [23] "GSM2685711_ctrl_12hr" "GSM2685712_ctrl_12hr"
## [25] "GSM2685713_MERS_CoV_24hr" "GSM2685714_MERS_CoV_24hr"
## [27] "GSM2685715_MERS_CoV_24hr" "GSM2685716_MERS_CoV_24hr"
## [29] "GSM2685717_MERS_CoV_24hr" "GSM2685718_ctrl_24hr"
## [31] "GSM2685719_ctrl_24hr" "GSM2685720_ctrl_24hr"
## [33] "GSM2685721_ctrl_24hr" "GSM2685722_ctrl_24hr"
## [35] "GSM2685723_MERS_CoV_36hr" "GSM2685724_MERS_CoV_36hr"
## [37] "GSM2685725_MERS_CoV_36hr" "GSM2685726_MERS_CoV_36hr"
## [39] "GSM2685727_MERS_CoV_36hr" "GSM2685728_ctrl_36hr"
## [41] "GSM2685729_ctrl_36hr" "GSM2685730_ctrl_36hr"
## [43] "GSM2685731_ctrl_36hr" "GSM2685732_ctrl_36hr"
## [45] "GSM2685733_MERS_CoV_48hr" "GSM2685734_MERS_CoV_48hr"
## [47] "GSM2685735_MERS_CoV_48hr" "GSM2685736_MERS_CoV_48hr"
## [49] "GSM2685737_MERS_CoV_48hr" "GSM2685738_ctrl_48hr"
## [51] "GSM2685739_ctrl_48hr" "GSM2685740_ctrl_48hr"
## [53] "GSM2685741_ctrl_48hr" "GSM2685742_ctrl_48hr"
```

```
group <- rep(1:5,10)
```

```
Group <- gsub('1','Group_A',group)
```

```
Group <- gsub('2','Group_B',Group)
```

```
Group <- gsub('3','Group_C',Group)
```

```
Group <- gsub('4','Group_D',Group)
```

```
Group <- gsub('5','Group_E',Group)
```

```
names <- colnames(Series100509)[5:54]
```

```
Names <- paste(names,Group,sep='_')
```

```
newNames <- gsub('_MERS','', Names)
```

```
newNames
```

```
## [1] "GSM2685693_CoV_0hr_Group_A" "GSM2685694_CoV_0hr_Group_B"
## [3] "GSM2685695_CoV_0hr_Group_C" "GSM2685696_CoV_0hr_Group_D"
## [5] "GSM2685697_CoV_0hr_Group_E" "GSM2685698_ctrl_0hr_Group_A"
```

```
## [7] "GSM2685699_ctrl_0hr_Group_B" "GSM2685700_ctrl_0hr_Group_C"
## [9] "GSM2685701_ctrl_0hr_Group_D" "GSM2685702_ctrl_0hr_Group_E"
## [11] "GSM2685703_CoV_12hr_Group_A" "GSM2685704_CoV_12hr_Group_B"
## [13] "GSM2685705_CoV_12hr_Group_C" "GSM2685706_CoV_12hr_Group_D"
## [15] "GSM2685707_CoV_12hr_Group_E" "GSM2685708_ctrl_12hr_Group_A"
## [17] "GSM2685709_ctrl_12hr_Group_B" "GSM2685710_ctrl_12hr_Group_C"
## [19] "GSM2685711_ctrl_12hr_Group_D" "GSM2685712_ctrl_12hr_Group_E"
## [21] "GSM2685713_CoV_24hr_Group_A" "GSM2685714_CoV_24hr_Group_B"
## [23] "GSM2685715_CoV_24hr_Group_C" "GSM2685716_CoV_24hr_Group_D"
## [25] "GSM2685717_CoV_24hr_Group_E" "GSM2685718_ctrl_24hr_Group_A"
## [27] "GSM2685719_ctrl_24hr_Group_B" "GSM2685720_ctrl_24hr_Group_C"
## [29] "GSM2685721_ctrl_24hr_Group_D" "GSM2685722_ctrl_24hr_Group_E"
## [31] "GSM2685723_CoV_36hr_Group_A" "GSM2685724_CoV_36hr_Group_B"
## [33] "GSM2685725_CoV_36hr_Group_C" "GSM2685726_CoV_36hr_Group_D"
## [35] "GSM2685727_CoV_36hr_Group_E" "GSM2685728_ctrl_36hr_Group_A"
## [37] "GSM2685729_ctrl_36hr_Group_B" "GSM2685730_ctrl_36hr_Group_C"
## [39] "GSM2685731_ctrl_36hr_Group_D" "GSM2685732_ctrl_36hr_Group_E"
## [41] "GSM2685733_CoV_48hr_Group_A" "GSM2685734_CoV_48hr_Group_B"
## [43] "GSM2685735_CoV_48hr_Group_C" "GSM2685736_CoV_48hr_Group_D"
## [45] "GSM2685737_CoV_48hr_Group_E" "GSM2685738_ctrl_48hr_Group_A"
## [47] "GSM2685739_ctrl_48hr_Group_B" "GSM2685740_ctrl_48hr_Group_C"
## [49] "GSM2685741_ctrl_48hr_Group_D" "GSM2685742_ctrl_48hr_Group_E"
```

Change the column names in Series100509 to the new column names identifying which group the samples is from in A:E.

```
colnames(Series100509)[5:54] <- newNames
colnames(Series100509)
```

```
## [1] "SPOT_ID" "GENE_SYMBOL"
## [3] "DESCRIPTION" "SEQUENCE"
## [5] "GSM2685693_CoV_0hr_Group_A" "GSM2685694_CoV_0hr_Group_B"
## [7] "GSM2685695_CoV_0hr_Group_C" "GSM2685696_CoV_0hr_Group_D"
## [9] "GSM2685697_CoV_0hr_Group_E" "GSM2685698_ctrl_0hr_Group_A"
## [11] "GSM2685699_ctrl_0hr_Group_B" "GSM2685700_ctrl_0hr_Group_C"
## [13] "GSM2685701_ctrl_0hr_Group_D" "GSM2685702_ctrl_0hr_Group_E"
## [15] "GSM2685703_CoV_12hr_Group_A" "GSM2685704_CoV_12hr_Group_B"
## [17] "GSM2685705_CoV_12hr_Group_C" "GSM2685706_CoV_12hr_Group_D"
## [19] "GSM2685707_CoV_12hr_Group_E" "GSM2685708_ctrl_12hr_Group_A"
## [21] "GSM2685709_ctrl_12hr_Group_B" "GSM2685710_ctrl_12hr_Group_C"
## [23] "GSM2685711_ctrl_12hr_Group_D" "GSM2685712_ctrl_12hr_Group_E"
## [25] "GSM2685713_CoV_24hr_Group_A" "GSM2685714_CoV_24hr_Group_B"
## [27] "GSM2685715_CoV_24hr_Group_C" "GSM2685716_CoV_24hr_Group_D"
## [29] "GSM2685717_CoV_24hr_Group_E" "GSM2685718_ctrl_24hr_Group_A"
## [31] "GSM2685719_ctrl_24hr_Group_B" "GSM2685720_ctrl_24hr_Group_C"
## [33] "GSM2685721_ctrl_24hr_Group_D" "GSM2685722_ctrl_24hr_Group_E"
## [35] "GSM2685723_CoV_36hr_Group_A" "GSM2685724_CoV_36hr_Group_B"
## [37] "GSM2685725_CoV_36hr_Group_C" "GSM2685726_CoV_36hr_Group_D"
## [39] "GSM2685727_CoV_36hr_Group_E" "GSM2685728_ctrl_36hr_Group_A"
## [41] "GSM2685729_ctrl_36hr_Group_B" "GSM2685730_ctrl_36hr_Group_C"
```

```
## [43] "GSM2685731_ctrl_36hr_Group_D" "GSM2685732_ctrl_36hr_Group_E"
## [45] "GSM2685733_CoV_48hr_Group_A"  "GSM2685734_CoV_48hr_Group_B"
## [47] "GSM2685735_CoV_48hr_Group_C"  "GSM2685736_CoV_48hr_Group_D"
## [49] "GSM2685737_CoV_48hr_Group_E"  "GSM2685738_ctrl_48hr_Group_A"
## [51] "GSM2685739_ctrl_48hr_Group_B"  "GSM2685740_ctrl_48hr_Group_C"
## [53] "GSM2685741_ctrl_48hr_Group_D"  "GSM2685742_ctrl_48hr_Group_E"
```

Remove the ID and SPOT_ID fields of the probe labels that won't be needed for the analysis.

```
Series89 <- Series89166_89160[, -1]
Series100 <- Series100509[, -c(1,3,4)]
```

Now combine the series together by genes in common.

```
ComboLiverCapillarySequences <- merge(Series89, Series100, by.x='GENE_SYMBOL',
                                       by.y='GENE_SYMBOL')
```

There should be different genotypes or copy number variations in the SEQUENCE feature column, which will identify which nucleotide jumps, is deleted, rearranged in the gene expressions. This time around let's group by SEQUENCE to see how many genotypes there are in all the genes. This could give more information in the analysis of any genotypes of genes could be more susceptible to pathogenesis of CoV or immunity to it. ***

```
library(dplyr)
```

This next portion of this analysis shows the top five genes in the data, the number of genotypes or copy number variations of the nucleotide sequences are in each gene, and the gene name. This could be useful to determine how well the genotypes within these five genes that were expressed more than the other 65k genes were when analyzing the effects of CoV, inactive CoV, CoV treated with an interleukin, and the control samples.

```
SeqGroup <- ComboLiverCapillarySequences %>% group_by(GENE_SYMBOL) %>%
count(n=n())
SeqGroup <- SeqGroup[order(SeqGroup$n, decreasing=TRUE), -3]

SeqGroup5 <- SeqGroup[1:5,]

genes5sequences <- merge(SeqGroup5, ComboLiverCapillarySequences,
                        by.x='GENE_SYMBOL', by.y='GENE_SYMBOL')

genotypes5 <- genes5sequences %>% group_by(SEQUENCE) %>% count(n=n)

genotypes5$n <- as.factor(genotypes5$n)
SeqGroup5$n <- as.factor(SeqGroup5$n)

genotypes5_1 <- merge(genotypes5, SeqGroup5, by.x='n', by.y='n')
colnames(genotypes5_1)[c(1,3)] <- c('geneCount', 'genotypeCount')
head(genotypes5_1)
```



```
##      geneCount                               SEQUENCE
## 1      255  AAACCTTACTCCAGAGCTCCTTGTGCATCTGACCAGCACCATCGACAGAATAAACACAGAA
## 2      255  AACAGAGTCCTCAGGGAAGAAAATCGAAGACTTCAGGCTCAACTGAGTCATGTTTCCAGA
## 3      255  GCACCTGTGTTCTTTGAGTTACATCATGAATGTGGTGATTTCCCAGATACCATCTCAGG
## 4      255  ATGGGGTGCTCTGGGGAAAATATTGGAGGGTCATCCATTCCACATTAAGAGCAAGTTGT
## 5      255  AAGTGCTTGGAATACTTGGGTGAATGTTACCAGACTCCTTCTCTCTCAGCTTACAGCCT
## 6      255  AATCTACGAGGCACTTTATGGCAATTCCAAGAAGGGGCTGAAAGGTATGTGTTCTTCTCC
##      genotypeCount GENE_SYMBOL
## 1      15      PDE4DIP
## 2      15      PDE4DIP
## 3      15      PDE4DIP
## 4      15      PDE4DIP
## 5      15      PDE4DIP
## 6      15      PDE4DIP
```

Write the files above out to csv.

```
write.csv(ComboLiverCapillarySequences,
'SequencesBothCleaned.csv',row.names=FALSE)
write.csv(genotypes5_1, 'Genotypes_5_1.csv', row.names=FALSE)
```

```
ComboCNV <- ComboLiverCapillarySequences[,-c(1,2)]
copynumbers <- merge(genotypes5_1,ComboCNV,
                     by.x='SEQUENCE', by.y='SEQUENCE')
CNV <- copynumbers[!duplicated(copynumbers$SEQUENCE),]
write.csv(CNV, 'copyNumbers.csv', row.names=FALSE)
```

CNV

```
##                               SEQUENCE geneCount
## 1  AAACCTTACTCCAGAGCTCCTTGTGCATCTGACCAGCACCATCGACAGAATAAACACAGAA      255
## 16 AACAGAGTCCTCAGGGAAGAAAATCGAAGACTTCAGGCTCAACTGAGTCATGTTTCCAGA      255
## 31 AAGGATTTGCTTATAAGGGTTCCTGCTTTACAAAATTATTCCAGGTTTTATGTGTCAGG       70
## 41 AAGGATTTGGTTGTAAGGGCTCCCGCTTTACAGAATTATTCCAGGGTTTTATGTGTCAGG       70
## 51 AAGTGCTTGGAATACTTGGGTGAATGTTACCAGACTCCTTCTCTCTCAGCTTACAGCCT      255
## 66 AATCTACGAGGCACTTTATGGCAATTCCAAGAAGGGGCTGAAAGGTATGTGTTCTTCTCC      255
## 81 AATGATCAGAAAAAGAAAGAAGCCAAAGAGAAAGGTACCTGGGTTCAACTGAAGTGCCAG       81
## 90 ACAAGTGTTACCATGGCAAACTGGAAGAGTCTACAATGTTACCCAGCATGCTGTTGGCA       81
## 99 ACCCTACAGCTCTTCCAAATATGCCACTGACCTTTTGAGTGTGGCTTTGAACAGGAACTT       70
## 109 ACGCTGTTGATGCCGGCAATATTGCTACTTTCGCTTTTTTGCAAATGCATTCACTTTGACA       70
## 209 AGGTCAAGTTTATACATCTTAATTATGGTGGAATTCCTATGTAGAGTCTAAAAAGCCAGG       70
## 219 ATATGTAAAGGGCCATTCTTAAGTTCTCTCCTTAACTTAATGCTGTCAAGTGTTAGATG       90
## 279 ATCCGGGACATGATCCGCAGGGCCCAGAGTTACCGAGTCCTCACTACTTTTCTTCCAGAC       90
## 285 ATCTTGTCATGGCAAATGCTGGACCCAACACAAATGGTTCCCAGTTTTTTCATCTGCACT       70
## 385 ATGGAGATTACAACCTGCTTGTTGGAGACGATCCTGTGTGATCAGCTTCATAAAAGGGCTT       90
## 391 ATGGATGACACAAACAGCGAGCAGCAGTTTAGAGTCTTCAGAGACTTCGACTTCCTAGAT       90
## 397 ATGGGGTGCTCTGGGGAAAATATTGGAGGGTCATCCATTCCACATTAAGAGCAAGTTGT      255
## 412 ATTTGCCTTGTTTATTGTGAGCATGGAATACTTCTGGAAGCTTTTCTAGTAGATTTTTTCT       70
## 422 CAAAAAATGGACTAGAAGAGAAGCTGGCTGAGGAGCTGAGATCAGCCTCGTGGCCTGGGT      255
## 437 CAAAAACTGGGAAGATGTACCAGGAGACCAGGTCAAGCCCGACCAATACACTGAGGCCCT      255
## 452 CAGGACAAGTTCCTTCTAAAATCAATCCTTTGCTTTGGCCTGGAAGACCGAACATATTAG      255
```

```

## 467 CCTTTCTGACAGCTATGGAATACCGTTAGCCAAGGTCCACTTGGCCCAGCACTAAGAAAA 255
## 482 CTACCAAAGCTGTGTATTTTTTCATTTCTTCATGGCACTGTGCTGTTAATTTCTGTTAACA 255
## 497 CTATGGAATACGATTAGCCAAGGTCCACTTGGCCCAGCACTAAGAAGAAGATGCATAGTT 255
## 512 CTGTTCTTTTGGCGACGTATATGCGAATCTATAAGAAAGGTGATATTGTAGACATCAAGG 81
## 521 GAACACAAAGAGAAAAGAGGAGGGGCACCCGATATATGTTCTTTAGGCCTTTTAGAAAAACA 81
## 530 GAAGCTGCTGCTATCAGAAGCCACTGTCTTTGCTCAGGCGAACGAGCTGGAGAAATACAG 255
## 545 GAGGTTTCTTCGTTTTTGAAAAGTAGAGGTATAAAATAACCTTTTTTGCTGGATACTGTC 90
## 551 GCACCTGTGTTCTTTGAGTTCACATCATGAATGTGGTGATTTCCCAGATACCATCTCAGG 255
## 566 GGAAGGAGTCTGAGCTGCTGGAACCTATTCCTATGAATTCATGGCATAATAGGTGTTAA 81
## 575 GGCCACATATATGCTAATCTATAAGAAAGGTGATATTGCAGACATCAAGGGAAGGGGTAC 81
## 584 GTGGACAACCTCGAATAAGTTTTGACTTGTGTTTTATCTTAACCACCAGATCATTCTTCTG 70
## 594 GTGTTGAGTCAGTGGGGACCAGCCTGCCCACAGGAGAAAATTTAAAGACACAAAAACAAA 255
## 609 GTTACCCAGCATGCTGTTGGCATTGTTGTAAACAAACAAGTTAAAGGCAAGATTCTTGCC 81
## 618 GTTGTAACAAACAAGTTAAGGGCAAGATTCTTGCCAAGAGAATTAATGTGCGTATTGAG 81
## 627 TCCCAGATACCATTTCAGGCTTAACCTAGCACATCCTATTTCTTTCTTCTATGATATCC 255
## 642 TGGTATAAAAGGCAAAATGGCATTGAGGATGAATCACATGAATACAGACCAAGAACAAGC 90
## 648 TTAAGCACTCTAAGAGCCGAGATAGCTTCCTGAAATGTGTGAAGGAAAATGATCAGAAAA 81
## 657 TTCCATCACCTTTCTTGAAAATATATCTTCAGCTTTGGGTAGGAGGAATCTTGGTGTAT 255
## 672 TTGAGGAAGTCTGTATAGGGTGACCCATTGTGTCTCAGTACCACGTTTTCTTTCTTTTT 70
## 682 TTTGGCGTGCAAGGAATATGAGCAAGGCAGAAGCTGTCTGTGCTGCTCTGCTGGCCTCTCA 70
## 692 TTTTAAAGAAGGCAAGCACTAACTTAGAGGAACCATGCAAGAAGCGCAGCCACCAGAAGT 255

```

```

##      genotypeCount GENE_SYMBOL GSM2359851_CoV1 GSM2359853_CoV2
GSM2359910_CoV3

```

## 1	15	PDE4DIP	27.80000	12.33750
3.696339				
## 16	15	PDE4DIP	14.38750	20.88750
2.637107				
## 31	5	HSD17B7	5145.79000	5373.31250
12.672146				
## 41	5	HSD17B7	11934.96500	10287.10500
13.095230				
## 51	15	PDE4DIP	32.27375	16.51000
3.847676				
## 66	15	PDE4DIP	16.67250	23.47500
3.444825				
## 81	9	RPL21	13038.30250	11010.62750
13.528036				
## 90	9	RPL21	30119.53750	30572.70750
14.712802				
## 99	5	HSD17B7	371.82000	322.18250
8.134598				
## 109	50	HSD17B7	213.67500	195.09500
7.177490				
## 209	5	HSD17B7	268.28000	198.08250
7.012814				
## 219	60	FRY	271.65250	357.92000
8.202159				
## 279	6	FRY	58.96375	84.71000
5.671111				
## 285	50	HSD17B7	17904.94250	19128.75750

13.734397				
## 385	6	FRY	9.06500	20.70250
3.829277				
## 391	6	FRY	32.45750	23.73000
4.310822				
## 397	15	PDE4DIP	75.22875	90.13250
5.752750				
## 412	5	HSD17B7	15.40250	19.88250
4.420457				
## 422	15	PDE4DIP	10.65000	12.76750
2.504837				
## 437	15	PDE4DIP	91.41000	103.70750
6.408463				
## 452	15	PDE4DIP	237.00000	63.49250
7.746742				
## 467	15	PDE4DIP	30.81875	39.75750
4.488047				
## 482	15	PDE4DIP	125.58750	134.66250
5.723797				
## 497	15	PDE4DIP	29.16750	11.06750
3.676894				
## 512	9	RPL21	5280.45750	3875.71750
12.254249				
## 521	9	RPL21	6381.22250	6865.95250
12.451731				
## 530	15	PDE4DIP	19.30000	36.25375
3.989351				
## 545	6	FRY	22.37500	36.61625
5.296403				
## 551	15	PDE4DIP	104.62875	85.94500
5.821977				
## 566	9	RPL21	46392.50000	45883.44000
15.416743				
## 575	9	RPL21	3746.36500	2738.23500
12.059480				
## 584	5	HSD17B7	43078.28000	35250.26250
15.063675				
## 594	15	PDE4DIP	18.22125	34.20625
5.068949				
## 609	9	RPL21	24563.01000	27131.15750
14.494375				
## 618	9	RPL21	6638.93000	6876.98250
12.970757				
## 627	15	PDE4DIP	63.95375	61.22750
5.621959				
## 642	6	FRY	32.43250	19.90500
4.781334				
## 648	9	RPL21	28633.13750	35054.02500
14.760531				
## 657	15	PDE4DIP	70.56500	45.82500

5.832804				
## 672	5	HSD17B7	23.60250	12.80875
3.931491				
## 682	5	HSD17B7	119.60500	123.33000
6.507425				
## 692	15	PDE4DIP	18.62250	18.30500
3.088424				
##	GSM2359913_CoV4	GSM2359850_ctrl11	GSM2359852_ctrl12	GSM2359911_ctrl13
## 1	3.790725	16.5200	17.73000	3.578243
## 16	3.344586	11.3625	18.11250	3.551566
## 31	11.573693	9277.9550	8067.56500	12.808809
## 41	11.838288	17429.8675	16112.38250	13.205094
## 51	4.423909	18.1100	13.44125	4.284877
## 66	4.177188	18.1550	17.48000	3.766645
## 81	13.571143	13483.4400	9879.27000	13.472757
## 90	14.614834	34445.3250	28827.18500	14.702680
## 99	8.181267	154.2550	220.73750	6.715907
## 109	7.729111	121.7325	172.01750	6.576652
## 209	6.513342	553.5950	301.95250	7.412307
## 219	8.706800	97.3250	202.96750	7.429588
## 279	6.315514	31.9675	48.28000	4.905673
## 285	12.522699	21680.6475	27867.61750	13.917649
## 385	3.968947	12.5900	13.29000	4.006208
## 391	3.833337	13.9325	16.31500	3.619992
## 397	6.626211	78.0075	68.24750	6.160618
## 412	4.407728	19.2275	30.75000	5.107143
## 422	4.583563	8.9625	14.71000	4.094687
## 437	6.654450	138.3750	138.58000	6.012364
## 452	7.233624	16.8675	11.08500	4.309826
## 467	4.560705	16.7325	30.56750	3.631049
## 482	6.186636	118.2613	77.22250	5.631851
## 497	3.528587	13.5300	15.50500	4.189106
## 512	12.136696	7492.1175	4175.44000	12.504104
## 521	12.137630	8573.3300	6704.05500	12.609506
## 530	3.853302	23.2675	22.38500	4.161641
## 545	5.767478	37.2500	51.34750	4.881077
## 551	5.744729	58.5300	35.14000	5.162767
## 566	15.387552	47613.6350	44025.17500	15.357757
## 575	11.973380	5260.6125	3136.01000	12.295686
## 584	14.145177	49908.4750	42313.35750	15.161325
## 594	4.875253	31.3550	26.55875	5.163443
## 609	14.396291	25395.2325	27531.33250	14.490896
## 618	12.907781	9555.7550	7781.74500	13.138377
## 627	5.613410	33.8425	60.97500	4.944435
## 642	5.524324	20.2175	18.00625	4.146513
## 648	14.719929	30448.7650	30448.76500	14.784022
## 657	5.556593	33.6350	46.85125	4.272972
## 672	3.564095	17.3825	28.25250	4.269237
## 682	6.579500	67.2175	90.32750	5.815040
## 692	3.260962	29.7900	11.46000	3.011269

##	GSM2359914_ctrl14	GSM2359912_IL1	GSM2359917_IL2
GSM2359915_inactiveHeatCoV1			
## 1	3.624038	3.140928	3.670410
3.546095			
## 16	4.151561	2.785070	2.552815
4.287654			
## 31	12.783884	13.140587	12.767002
13.136716			
## 41	12.949140	13.434763	13.042793
13.454729			
## 51	3.420708	3.601485	3.039668
3.450441			
## 66	4.254487	4.957138	3.629882
3.956468			
## 81	13.614149	13.633448	13.630940
13.430600			
## 90	14.702680	14.750075	14.688661
14.633296			
## 99	6.808397	6.558839	6.755108
6.888168			
## 109	6.391192	6.560465	6.498674
6.666895			
## 209	7.243844	7.590200	7.424797
7.242360			
## 219	6.633365	7.263232	7.037296
7.826073			
## 279	4.654331	5.951235	5.315131
5.197742			
## 285	13.530332	13.984778	13.664711
14.101873			
## 385	4.078738	3.198799	3.613910
4.292263			
## 391	2.635114	4.035037	3.496328
5.051240			
## 397	5.843770	6.207442	6.010504
6.287897			
## 412	4.780039	5.136244	3.083101
4.441393			
## 422	3.395185	3.520729	4.372599
3.739993			
## 437	6.430275	5.891864	6.410106
6.434970			
## 452	2.912074	3.819033	4.434493
2.699398			
## 467	3.428464	3.873689	4.308965
3.544474			
## 482	6.086149	5.750065	5.901224
5.722239			
## 497	3.798390	3.882634	3.369441
4.407039			

## 512	12.605234	12.523691	12.559814
12.367516			
## 521	12.586550	12.674812	12.614916
12.498591			
## 530	3.876763	3.252636	3.868612
3.849681			
## 545	5.088791	5.049082	5.006710
5.221105			
## 551	5.219856	4.821215	5.311027
5.561438			
## 566	15.331644	15.416743	15.309413
15.230848			
## 575	12.471203	12.359393	12.492727
12.373389			
## 584	14.933096	15.247829	14.906973
15.262285			
## 594	6.126075	4.908142	5.656664
5.879604			
## 609	14.451526	14.423394	14.512645
14.388731			
## 618	13.191864	13.159157	13.149836
13.099425			
## 627	5.193310	5.223439	5.285965
4.153434			
## 642	4.469801	4.652000	4.351019
4.885096			
## 648	14.876129	14.859144	14.836318
14.767892			
## 657	5.027764	5.223439	4.305323
4.621451			
## 672	5.066429	3.447057	4.335341
3.690485			
## 682	5.849889	5.674321	5.635349
6.016112			
## 692	3.249961	3.208008	3.379703
3.524824			
##	GSM2359916_inactiveHeatCoV2	GSM2685693_CoV_0hr_Group_A	
## 1	3.478594	6.046074	
## 16	3.088424	6.844269	
## 31	12.926635	13.743911	
## 41	13.390188	13.354586	
## 51	3.302000	6.558291	
## 66	3.372408	6.123684	
## 81	13.564872	15.471859	
## 90	14.805995	13.952961	
## 99	6.090212	7.760966	
## 109	6.421314	7.760966	
## 209	7.501435	9.039621	
## 219	7.244399	5.959534	
## 279	5.555414	11.322402	

## 285	13.863342	9.039621
## 385	2.960501	11.322402
## 391	2.435691	10.206249
## 397	6.065265	6.558291
## 412	3.771957	7.443340
## 422	3.962144	6.642102
## 437	6.320798	6.642102
## 452	3.481031	6.141494
## 467	2.245824	7.494292
## 482	5.751764	6.642102
## 497	3.064070	6.558291
## 512	12.608269	14.216561
## 521	12.717626	14.379203
## 530	4.276882	6.141494
## 545	5.310430	10.206249
## 551	4.847402	6.642102
## 566	15.381582	12.710011
## 575	12.473387	14.379203
## 584	15.137849	9.039621
## 594	5.326159	7.494292
## 609	14.530418	13.952961
## 618	13.191864	13.952961
## 627	4.898910	6.141494
## 642	4.637404	11.322402
## 648	14.915921	15.209228
## 657	5.464345	6.141494
## 672	3.477792	6.585250
## 682	5.770293	7.760966
## 692	3.860184	6.558291
##	GSM2685694_CoV_0hr_Group_B	GSM2685695_CoV_0hr_Group_C
## 1	5.934566	5.909934
## 16	6.971307	7.045377
## 31	14.112591	13.731221
## 41	13.772142	13.242660
## 51	6.538601	6.586001
## 66	6.109128	6.141085
## 81	15.541398	15.753597
## 90	13.758973	13.253953
## 99	7.679814	7.857275
## 109	7.679814	7.857275
## 209	9.012739	8.926670
## 219	5.955123	6.360556
## 279	11.419858	11.426284
## 285	9.012739	8.926670
## 385	11.419858	11.426284
## 391	10.558187	10.338125
## 397	6.538601	6.586001
## 412	7.360626	7.568523
## 422	6.488311	6.650799
## 437	6.488311	6.650799

## 452	6.146797	6.177145
## 467	7.657015	7.460948
## 482	6.488311	6.650799
## 497	6.538601	6.586001
## 512	14.150090	13.748373
## 521	14.347845	14.203153
## 530	6.146797	6.177145
## 545	10.558187	10.338125
## 551	6.488311	6.650799
## 566	12.681303	12.567839
## 575	14.347845	14.203153
## 584	9.012739	8.926670
## 594	7.657015	7.460948
## 609	13.758973	13.253953
## 618	13.758973	13.253953
## 627	6.180332	6.177145
## 642	11.419858	11.426284
## 648	15.162455	15.081966
## 657	6.146797	6.177145
## 672	6.552001	6.801947
## 682	7.679814	7.857275
## 692	6.538601	6.586001
##	GSM2685696_CoV_0hr_Group_D	GSM2685697_CoV_0hr_Group_E
## 1	5.928474	6.049563
## 16	7.000706	7.075463
## 31	13.879932	13.711439
## 41	13.423286	13.239789
## 51	6.563639	6.563182
## 66	6.181922	6.206088
## 81	15.743156	15.721350
## 90	13.545156	13.621073
## 99	7.798678	7.866392
## 109	7.798678	7.866392
## 209	8.949465	8.842747
## 219	5.958143	6.037140
## 279	11.359374	11.441523
## 285	8.949465	8.842747
## 385	11.359374	11.441523
## 391	10.330821	10.326638
## 397	6.563639	6.563182
## 412	7.419171	7.472914
## 422	6.697222	6.627382
## 437	6.697222	6.627382
## 452	6.137030	6.250091
## 467	7.350873	7.432630
## 482	6.697222	6.627382
## 497	6.563639	6.563182
## 512	14.150090	14.144609
## 521	14.358264	14.321305
## 530	6.137030	6.250091

## 545	10.330821	10.326638
## 551	6.697222	6.627382
## 566	12.706625	12.736276
## 575	14.358264	14.321305
## 584	8.949465	8.842747
## 594	7.350873	7.432630
## 609	13.545156	13.621073
## 618	13.545156	13.621073
## 627	6.088795	6.182121
## 642	11.359374	11.441523
## 648	15.220710	15.204887
## 657	6.137030	6.250091
## 672	6.764389	6.805821
## 682	7.798678	7.866392
## 692	6.563639	6.563182
##	GSM2685698_ctrl_0hr_Group_A	GSM2685699_ctrl_0hr_Group_B
## 1	5.968496	5.917775
## 16	7.021435	6.877443
## 31	13.679609	13.690249
## 41	13.189063	13.052863
## 51	6.640295	6.454374
## 66	6.130249	6.183511
## 81	15.721350	15.550718
## 90	13.582989	13.866949
## 99	7.742188	7.684730
## 109	7.742188	7.684730
## 209	8.900764	8.808501
## 219	5.887260	5.873911
## 279	11.517387	11.472704
## 285	8.900764	8.808501
## 385	11.517387	11.472704
## 391	10.480783	10.308663
## 397	6.640295	6.454374
## 412	7.415712	7.155158
## 422	6.564249	6.732084
## 437	6.564249	6.732084
## 452	6.236379	6.226813
## 467	7.430458	7.661405
## 482	6.564249	6.732084
## 497	6.640295	6.454374
## 512	14.113341	14.273581
## 521	14.361286	14.343081
## 530	6.236379	6.226813
## 545	10.480783	10.308663
## 551	6.564249	6.732084
## 566	12.748984	12.847926
## 575	14.361286	14.343081
## 584	8.900764	8.808501
## 594	7.430458	7.661405
## 609	13.582989	13.866949

## 618	13.582989	13.866949
## 627	6.142107	6.213495
## 642	11.517387	11.472704
## 648	15.239825	15.250367
## 657	6.236379	6.226813
## 672	6.694174	6.657960
## 682	7.742188	7.684730
## 692	6.640295	6.454374
##	GSM2685700_ctrl_0hr_Group_C	GSM2685701_ctrl_0hr_Group_D
## 1	5.900602	5.973161
## 16	6.843578	7.033153
## 31	13.961256	14.155264
## 41	13.405484	13.621073
## 51	6.529181	6.591771
## 66	6.192774	6.209796
## 81	15.560146	15.569952
## 90	13.842350	13.774116
## 99	7.532076	7.701530
## 109	7.532076	7.701530
## 209	8.868502	8.890951
## 219	5.946961	6.347886
## 279	11.495049	11.440057
## 285	8.868502	8.890951
## 385	11.495049	11.440057
## 391	10.169702	10.282719
## 397	6.529181	6.591771
## 412	7.145501	7.226388
## 422	6.588402	6.522684
## 437	6.588402	6.522684
## 452	6.124304	6.158761
## 467	7.544183	7.562423
## 482	6.588402	6.522684
## 497	6.529181	6.591771
## 512	14.298329	14.214335
## 521	14.338710	14.413252
## 530	6.124304	6.158761
## 545	10.169702	10.282719
## 551	6.588402	6.522684
## 566	12.755767	12.755081
## 575	14.338710	14.413252
## 584	8.868502	8.890951
## 594	7.544183	7.562423
## 609	13.842350	13.774116
## 618	13.842350	13.774116
## 627	6.192774	6.258751
## 642	11.495049	11.440057
## 648	15.315381	15.310287
## 657	6.124304	6.158761
## 672	6.664146	6.663506
## 682	7.532076	7.701530

## 692	6.529181	6.591771
##	GSM2685702_ctrl_0hr_Group_E	GSM2685703_CoV_12hr_Group_A
## 1	5.965960	6.034281
## 16	6.998620	6.550617
## 31	13.806404	12.563610
## 41	13.289162	12.293758
## 51	6.439847	6.361784
## 66	6.250186	6.078798
## 81	15.486730	15.791166
## 90	13.715296	13.793138
## 99	7.715531	7.140323
## 109	7.715531	7.140323
## 209	8.866401	8.389528
## 219	5.979920	5.940803
## 279	11.482451	9.984313
## 285	8.866401	8.389528
## 385	11.482451	9.984313
## 391	10.311346	11.123686
## 397	6.439847	6.361784
## 412	7.227923	6.437373
## 422	6.730183	6.617164
## 437	6.730183	6.617164
## 452	6.122752	6.218734
## 467	7.728887	7.959343
## 482	6.730183	6.617164
## 497	6.439847	6.361784
## 512	14.176559	14.214335
## 521	14.272338	14.427768
## 530	6.122752	6.218734
## 545	10.311346	11.123686
## 551	6.730183	6.617164
## 566	12.780676	12.710683
## 575	14.272338	14.427768
## 584	8.866401	8.389528
## 594	7.728887	7.959343
## 609	13.715296	13.793138
## 618	13.715296	13.793138
## 627	6.200249	6.120370
## 642	11.482451	9.984313
## 648	15.204887	15.384111
## 657	6.122752	6.218734
## 672	6.672430	6.539512
## 682	7.715531	7.140323
## 692	6.439847	6.361784
##	GSM2685704_CoV_12hr_Group_B	GSM2685705_CoV_12hr_Group_C
## 1	5.986079	5.976717
## 16	6.692292	7.076586
## 31	12.931461	12.725504
## 41	12.506969	12.484960
## 51	6.351778	6.239437

## 66	5.986079	6.136415
## 81	15.989759	15.811264
## 90	13.756643	13.821900
## 99	7.179646	7.139658
## 109	7.179646	7.139658
## 209	8.009192	7.905606
## 219	5.996914	5.926815
## 279	10.224721	9.970963
## 285	8.009192	7.905606
## 385	10.224721	9.970963
## 391	11.205396	11.354632
## 397	6.351778	6.239437
## 412	6.385745	6.401408
## 422	6.501226	6.364588
## 437	6.501226	6.364588
## 452	6.117671	6.081783
## 467	8.018902	7.924227
## 482	6.501226	6.364588
## 497	6.351778	6.239437
## 512	14.194950	14.194950
## 521	14.444955	14.410594
## 530	6.117671	6.081783
## 545	11.205396	11.354632
## 551	6.501226	6.364588
## 566	12.707371	12.707371
## 575	14.444955	14.410594
## 584	8.009192	7.905606
## 594	8.018902	7.924227
## 609	13.756643	13.821900
## 618	13.756643	13.821900
## 627	6.093028	6.136415
## 642	10.224721	9.970963
## 648	15.418299	15.310287
## 657	6.117671	6.081783
## 672	6.578775	6.465496
## 682	7.179646	7.139658
## 692	6.351778	6.239437
##	GSM2685706_CoV_12hr_Group_D	GSM2685707_CoV_12hr_Group_E
## 1	5.856839	5.976717
## 16	6.551540	6.801301
## 31	12.735467	13.112807
## 41	12.449308	12.457226
## 51	6.289630	6.313741
## 66	6.123684	6.140472
## 81	15.943282	16.123661
## 90	13.931108	13.637411
## 99	7.096637	7.207673
## 109	7.096637	7.207673
## 209	7.834892	8.052411
## 219	5.914191	5.929894

## 279	9.965923	9.961133
## 285	7.834892	8.052411
## 385	9.965923	9.961133
## 391	11.237178	12.011102
## 397	6.289630	6.313741
## 412	6.341379	6.788131
## 422	6.471339	6.454703
## 437	6.471339	6.454703
## 452	6.183709	6.237144
## 467	8.058512	7.772063
## 482	6.471339	6.454703
## 497	6.289630	6.313741
## 512	14.266920	14.212112
## 521	14.501147	14.459021
## 530	6.183709	6.237144
## 545	11.237178	12.011102
## 551	6.471339	6.454703
## 566	12.828171	12.789918
## 575	14.501147	14.459021
## 584	7.834892	8.052411
## 594	8.058512	7.772063
## 609	13.931108	13.637411
## 618	13.931108	13.637411
## 627	6.143740	6.182915
## 642	9.965923	9.961133
## 648	15.427256	15.815665
## 657	6.183709	6.237144
## 672	6.511225	6.687345
## 682	7.096637	7.207673
## 692	6.289630	6.313741
##	GSM2685708_ctrl_12hr_Group_A	GSM2685709_ctrl_12hr_Group_B
## 1	6.101625	6.054130
## 16	7.304274	7.267867
## 31	13.469030	13.714536
## 41	13.045695	12.976714
## 51	6.556913	6.581489
## 66	6.319955	6.301538
## 81	15.617498	15.684402
## 90	13.461204	13.242660
## 99	7.879929	7.815682
## 109	7.879929	7.815682
## 209	9.474104	9.384559
## 219	6.375403	5.919921
## 279	11.347625	11.309167
## 285	9.474104	9.384559
## 385	11.347625	11.309167
## 391	11.198476	11.681931
## 397	6.556913	6.581489
## 412	7.353212	7.733486
## 422	6.677654	6.524879

## 437	6.677654	6.524879
## 452	6.556913	6.658103
## 467	8.266159	8.119336
## 482	6.677654	6.524879
## 497	6.556913	6.581489
## 512	13.906088	13.848895
## 521	14.079932	14.111086
## 530	6.556913	6.658103
## 545	11.198476	11.681931
## 551	6.677654	6.524879
## 566	12.471825	12.287010
## 575	14.079932	14.111086
## 584	9.474104	9.384559
## 594	8.266159	8.119336
## 609	13.461204	13.242660
## 618	13.461204	13.242660
## 627	6.239819	6.132205
## 642	11.347625	11.309167
## 648	15.152809	15.320354
## 657	6.556913	6.658103
## 672	6.844834	6.907652
## 682	7.879929	7.815682
## 692	6.556913	6.581489
##	GSM2685710_ctrl_12hr_Group_C	GSM2685711_ctrl_12hr_Group_D
## 1	6.056733	6.073452
## 16	7.233682	7.044011
## 31	13.529718	13.383898
## 41	12.970346	13.184021
## 51	6.587802	6.601072
## 66	6.278182	6.040386
## 81	15.651198	15.595299
## 90	13.642893	13.710017
## 99	7.818273	7.596400
## 109	7.818273	7.596400
## 209	9.389666	9.595884
## 219	5.837683	5.928711
## 279	11.282345	11.381308
## 285	9.389666	9.595884
## 385	11.282345	11.381308
## 391	11.098940	11.147201
## 397	6.587802	6.601072
## 412	7.306484	7.133544
## 422	6.604914	7.049685
## 437	6.604914	7.049685
## 452	6.579152	6.367561
## 467	8.186986	8.135535
## 482	6.604914	7.049685
## 497	6.587802	6.601072
## 512	14.033361	14.051589
## 521	14.260434	14.250394

## 530	6.579152	6.367561
## 545	11.098940	11.147201
## 551	6.604914	7.049685
## 566	12.569342	12.463613
## 575	14.260434	14.250394
## 584	9.389666	9.595884
## 594	8.186986	8.135535
## 609	13.642893	13.710017
## 618	13.642893	13.710017
## 627	6.298242	6.040386
## 642	11.282345	11.381308
## 648	15.271108	15.134904
## 657	6.579152	6.367561
## 672	6.834263	6.810573
## 682	7.818273	7.596400
## 692	6.587802	6.601072
##	GSM2685712_ctrl_12hr_Group_E	GSM2685713_CoV_24hr_Group_A
## 1	5.943683	5.852162
## 16	7.003569	6.871181
## 31	13.331293	11.921824
## 41	13.078701	11.600564
## 51	6.452728	6.771657
## 66	6.066732	6.237144
## 81	15.753597	15.727376
## 90	13.871693	13.644290
## 99	7.567679	7.437216
## 109	7.567679	7.437216
## 209	9.422309	7.967724
## 219	6.042577	5.981748
## 279	11.504963	8.847859
## 285	9.422309	7.967724
## 385	11.504963	8.847859
## 391	11.269751	11.056136
## 397	6.452728	6.771657
## 412	7.036521	6.781927
## 422	7.036124	6.689429
## 437	7.036124	6.689429
## 452	6.269044	6.237144
## 467	8.242917	7.835208
## 482	7.036124	6.689429
## 497	6.452728	6.771657
## 512	14.164148	14.073373
## 521	14.394424	14.290879
## 530	6.269044	6.237144
## 545	11.269751	11.056136
## 551	7.036124	6.689429
## 566	12.603593	12.505019
## 575	14.394424	14.290879
## 584	9.422309	7.967724
## 594	8.242917	7.835208

## 609	13.871693	13.644290
## 618	13.871693	13.644290
## 627	6.145983	7.148629
## 642	11.504963	8.847859
## 648	15.191398	15.334556
## 657	6.269044	6.237144
## 672	6.781657	6.595712
## 682	7.567679	7.437216
## 692	6.452728	6.771657
##	GSM2685714_CoV_24hr_Group_B	GSM2685715_CoV_24hr_Group_C
## 1	5.994476	5.975571
## 16	6.762131	7.027862
## 31	11.927784	11.939520
## 41	11.648884	11.386459
## 51	6.554766	6.962202
## 66	6.217184	6.279483
## 81	15.684402	15.989759
## 90	13.645558	13.465904
## 99	7.360648	7.359945
## 109	7.360648	7.359945
## 209	7.845939	7.896977
## 219	5.840833	5.837683
## 279	8.801857	8.940537
## 285	7.845939	7.896977
## 385	8.801857	8.940537
## 391	11.164641	11.167653
## 397	6.554766	6.962202
## 412	6.727406	6.858181
## 422	6.582995	6.700971
## 437	6.582995	6.700971
## 452	6.207846	6.556300
## 467	7.793875	7.589638
## 482	6.582995	6.700971
## 497	6.554766	6.962202
## 512	14.033361	13.983045
## 521	14.200832	14.115127
## 530	6.207846	6.556300
## 545	11.164641	11.167653
## 551	6.582995	6.700971
## 566	12.564091	12.663837
## 575	14.200832	14.115127
## 584	7.845939	7.896977
## 594	7.793875	7.589638
## 609	13.645558	13.465904
## 618	13.645558	13.465904
## 627	7.006153	7.128624
## 642	8.801857	8.940537
## 648	15.310287	15.537292
## 657	6.207846	6.556300
## 672	6.582995	6.743885

## 682	7.360648	7.359945
## 692	6.554766	6.962202
##	GSM2685716_CoV_24hr_Group_D	GSM2685717_CoV_24hr_Group_E
## 1	6.113093	6.081570
## 16	7.096057	7.274542
## 31	12.099577	12.317984
## 41	11.575374	11.876350
## 51	6.900674	6.827556
## 66	6.339209	6.214855
## 81	15.950377	15.955840
## 90	13.553828	13.536745
## 99	7.465240	7.550122
## 109	7.465240	7.550122
## 209	7.999472	8.024288
## 219	5.982204	6.329555
## 279	8.960830	8.883563
## 285	7.999472	8.024288
## 385	8.960830	8.883563
## 391	11.197719	11.185996
## 397	6.900674	6.827556
## 412	7.019623	7.078983
## 422	6.728551	6.627090
## 437	6.728551	6.627090
## 452	6.601369	6.523155
## 467	7.550776	7.513327
## 482	6.728551	6.627090
## 497	6.900674	6.827556
## 512	14.147362	14.130915
## 521	14.069355	14.099160
## 530	6.601369	6.523155
## 545	11.197719	11.185996
## 551	6.728551	6.627090
## 566	12.709173	12.825505
## 575	14.069355	14.099160
## 584	7.999472	8.024288
## 594	7.550776	7.513327
## 609	13.553828	13.536745
## 618	13.553828	13.536745
## 627	7.188175	7.254043
## 642	8.960830	8.883563
## 648	15.556080	15.605467
## 657	6.601369	6.523155
## 672	6.738571	6.763061
## 682	7.465240	7.550122
## 692	6.900674	6.827556
##	GSM2685718_ctrl_24hr_Group_A	GSM2685719_ctrl_24hr_Group_B
## 1	5.971028	6.006996
## 16	7.236172	7.108756
## 31	13.648472	13.539470
## 41	13.110465	13.124500

## 51	6.716954	6.591846
## 66	6.136825	6.274459
## 81	15.908781	15.698081
## 90	13.711263	13.752605
## 99	7.841700	7.824587
## 109	7.841700	7.824587
## 209	8.907840	8.912668
## 219	5.822295	5.874648
## 279	11.936456	11.941699
## 285	8.907840	8.912668
## 385	11.936456	11.941699
## 391	11.162259	11.112189
## 397	6.716954	6.591846
## 412	7.566727	7.559356
## 422	6.689709	6.704019
## 437	6.689709	6.704019
## 452	6.533938	6.482192
## 467	7.825446	7.853444
## 482	6.689709	6.704019
## 497	6.716954	6.591846
## 512	14.166197	14.210078
## 521	14.203153	14.316351
## 530	6.533938	6.482192
## 545	11.162259	11.112189
## 551	6.689709	6.704019
## 566	12.718928	12.526797
## 575	14.203153	14.316351
## 584	8.907840	8.912668
## 594	7.825446	7.853444
## 609	13.711263	13.752605
## 618	13.711263	13.752605
## 627	6.181525	6.087523
## 642	11.936456	11.941699
## 648	15.507334	15.519886
## 657	6.533938	6.482192
## 672	6.944874	6.829460
## 682	7.841700	7.824587
## 692	6.716954	6.591846
##	GSM2685720_ctrl_24hr_Group_C	GSM2685721_ctrl_24hr_Group_D
## 1	6.054347	6.016164
## 16	7.072893	7.111257
## 31	13.625575	13.478230
## 41	13.292808	13.190435
## 51	6.568645	6.576811
## 66	6.116008	6.084124
## 81	15.569952	15.591344
## 90	13.777280	13.791536
## 99	7.783632	7.805239
## 109	7.783632	7.805239
## 209	9.091769	8.998199

## 219	6.383866	5.961156
## 279	12.027264	11.940062
## 285	9.091769	8.998199
## 385	12.027264	11.940062
## 391	11.039873	11.061724
## 397	6.568645	6.576811
## 412	7.551625	7.509755
## 422	6.740191	6.801430
## 437	6.740191	6.801430
## 452	6.414388	6.366862
## 467	7.939793	7.793160
## 482	6.740191	6.801430
## 497	6.568645	6.576811
## 512	14.245321	14.178808
## 521	14.216561	14.062458
## 530	6.414388	6.366862
## 545	11.039873	11.061724
## 551	6.740191	6.801430
## 566	12.521572	12.518281
## 575	14.216561	14.062458
## 584	9.091769	8.998199
## 594	7.939793	7.793160
## 609	13.777280	13.791536
## 618	13.777280	13.791536
## 627	6.116008	6.100574
## 642	12.027264	11.940062
## 648	15.239825	15.262706
## 657	6.414388	6.366862
## 672	6.996312	6.968890
## 682	7.783632	7.805239
## 692	6.568645	6.576811
##	GSM2685722_ctrl_24hr_Group_E	GSM2685723_CoV_36hr_Group_A
## 1	6.065871	6.037753
## 16	7.153700	7.290151
## 31	13.511865	12.155727
## 41	13.198260	11.842594
## 51	6.596607	7.031230
## 66	6.189750	6.508058
## 81	15.575092	15.928588
## 90	13.818910	13.565265
## 99	7.700074	7.075356
## 109	7.700074	7.075356
## 209	9.018607	8.415451
## 219	5.946259	6.354954
## 279	11.842310	9.547340
## 285	9.018607	8.415451
## 385	11.842310	9.547340
## 391	11.162237	10.390335
## 397	6.596607	7.031230
## 412	7.483871	6.848334

## 422	6.601666	6.813655
## 437	6.601666	6.813655
## 452	6.306285	6.793453
## 467	7.728207	8.067333
## 482	6.601666	6.813655
## 497	6.596607	7.031230
## 512	14.167395	14.075771
## 521	14.060533	14.212112
## 530	6.306285	6.793453
## 545	11.162237	10.390335
## 551	6.601666	6.813655
## 566	12.560602	12.502460
## 575	14.060533	14.212112
## 584	9.018607	8.415451
## 594	7.728207	8.067333
## 609	13.818910	13.565265
## 618	13.818910	13.565265
## 627	6.158761	7.258195
## 642	11.842310	9.547340
## 648	15.217024	15.330807
## 657	6.306285	6.793453
## 672	6.891355	6.508058
## 682	7.700074	7.075356
## 692	6.596607	7.031230
##	GSM2685724_CoV_36hr_Group_B	GSM2685725_CoV_36hr_Group_C
## 1	6.075165	6.035602
## 16	7.038979	7.256969
## 31	12.263597	12.424504
## 41	11.984688	12.033150
## 51	6.920563	7.063761
## 66	6.419116	6.501226
## 81	15.963635	15.983105
## 90	13.673362	13.633552
## 99	7.076960	7.301533
## 109	7.076960	7.301533
## 209	8.349640	8.680838
## 219	5.813953	6.057816
## 279	9.470814	9.856731
## 285	8.349640	8.680838
## 385	9.470814	9.856731
## 391	10.657755	10.350293
## 397	6.920563	7.063761
## 412	6.838491	6.904409
## 422	6.648788	6.671737
## 437	6.648788	6.671737
## 452	6.564707	6.796636
## 467	8.033099	8.012799
## 482	6.648788	6.671737
## 497	6.920563	7.063761
## 512	14.151924	14.171229

## 521	14.264767	14.295878
## 530	6.564707	6.796636
## 545	10.657755	10.350293
## 551	6.648788	6.671737
## 566	12.621199	12.500993
## 575	14.264767	14.295878
## 584	8.349640	8.680838
## 594	8.033099	8.012799
## 609	13.673362	13.633552
## 618	13.673362	13.633552
## 627	7.091198	6.988153
## 642	9.470814	9.856731
## 648	15.410491	15.437017
## 657	6.564707	6.796636
## 672	6.540443	6.609427
## 682	7.076960	7.301533
## 692	6.920563	7.063761
##	GSM2685726_CoV_36hr_Group_D	GSM2685727_CoV_36hr_Group_E
## 1	6.040386	6.130249
## 16	7.247512	7.275706
## 31	12.440354	12.351661
## 41	12.082020	11.793496
## 51	6.929104	7.136585
## 66	6.419622	6.452728
## 81	15.943282	15.823487
## 90	13.520099	13.440834
## 99	7.613149	7.419926
## 109	7.613149	7.419926
## 209	8.576810	8.465203
## 219	5.894553	6.025496
## 279	9.696370	9.585969
## 285	8.576810	8.465203
## 385	9.696370	9.585969
## 391	10.378994	10.490391
## 397	6.929104	7.136585
## 412	6.833708	6.954109
## 422	6.626798	6.763061
## 437	6.626798	6.763061
## 452	6.671737	6.726782
## 467	8.166043	8.051814
## 482	6.626798	6.763061
## 497	6.929104	7.136585
## 512	14.097307	13.856158
## 521	14.230680	14.194950
## 530	6.671737	6.726782
## 545	10.378994	10.490391
## 551	6.626798	6.763061
## 566	12.440331	12.523054
## 575	14.230680	14.194950
## 584	8.576810	8.465203

## 594	8.166043	8.051814
## 609	13.520099	13.440834
## 618	13.520099	13.440834
## 627	7.194883	7.223682
## 642	9.696370	9.585969
## 648	15.315381	15.372981
## 657	6.671737	6.726782
## 672	6.615693	6.350894
## 682	7.613149	7.419926
## 692	6.929104	7.136585
##	GSM2685728_ctrl_36hr_Group_A	GSM2685729_ctrl_36hr_Group_B
## 1	6.026382	6.192380
## 16	7.224261	7.247512
## 31	13.717610	13.768831
## 41	13.315157	13.328181
## 51	6.797545	6.722416
## 66	6.220477	6.297905
## 81	15.705959	15.786439
## 90	13.530735	13.575280
## 99	7.815874	7.751624
## 109	7.815874	7.751624
## 209	8.793558	8.893911
## 219	6.125711	5.970798
## 279	12.382962	12.326271
## 285	8.793558	8.893911
## 385	12.382962	12.326271
## 391	11.308817	11.352996
## 397	6.797545	6.722416
## 412	7.509502	7.556387
## 422	6.478637	6.545394
## 437	6.478637	6.545394
## 452	6.437685	6.461104
## 467	8.036592	7.952603
## 482	6.478637	6.545394
## 497	6.797545	6.722416
## 512	14.084852	14.200832
## 521	14.176559	14.341048
## 530	6.437685	6.461104
## 545	11.308817	11.352996
## 551	6.478637	6.545394
## 566	12.611342	12.621199
## 575	14.176559	14.341048
## 584	8.793558	8.893911
## 594	8.036592	7.952603
## 609	13.530735	13.575280
## 618	13.530735	13.575280
## 627	6.235422	6.186089
## 642	12.382962	12.326271
## 648	15.213389	15.217024
## 657	6.437685	6.461104

## 672	6.815439	6.832745
## 682	7.815874	7.751624
## 692	6.797545	6.722416
##	GSM2685730_ctrl_36hr_Group_C	GSM2685731_ctrl_36hr_Group_D
## 1	6.083912	6.013710
## 16	7.135662	7.204644
## 31	13.867985	13.830899
## 41	13.379796	13.341531
## 51	6.836527	6.763127
## 66	6.236570	6.174351
## 81	15.807433	15.781413
## 90	13.536745	13.535802
## 99	7.830082	7.789513
## 109	7.830082	7.789513
## 209	8.814025	8.812357
## 219	5.881211	5.946493
## 279	12.428664	12.362715
## 285	8.814025	8.812357
## 385	12.428664	12.362715
## 391	11.305894	11.262485
## 397	6.836527	6.763127
## 412	7.571243	7.522806
## 422	6.544776	6.608540
## 437	6.544776	6.608540
## 452	6.321218	6.506314
## 467	7.977058	7.960183
## 482	6.544776	6.608540
## 497	6.836527	6.763127
## 512	14.082690	14.103762
## 521	14.264767	14.302531
## 530	6.321218	6.506314
## 545	11.305894	11.262485
## 551	6.544776	6.608540
## 566	12.570815	12.623629
## 575	14.264767	14.302531
## 584	8.814025	8.812357
## 594	7.977058	7.960183
## 609	13.536745	13.535802
## 618	13.536745	13.535802
## 627	6.282822	6.118087
## 642	12.428664	12.362715
## 648	15.330807	15.320354
## 657	6.321218	6.506314
## 672	6.870133	6.829460
## 682	7.830082	7.789513
## 692	6.836527	6.763127
##	GSM2685732_ctrl_36hr_Group_E	GSM2685733_CoV_48hr_Group_A
## 1	5.996010	6.245535
## 16	7.221895	7.588102
## 31	13.814027	13.145100

## 41	13.283499	12.611342
## 51	6.748119	7.418830
## 66	6.206479	6.421979
## 81	15.863114	15.770868
## 90	13.505737	13.308149
## 99	7.876741	7.615284
## 109	7.876741	7.615284
## 209	8.907780	8.896563
## 219	5.986761	5.952795
## 279	12.360358	11.238626
## 285	8.907780	8.896563
## 385	12.360358	11.238626
## 391	11.386529	9.754081
## 397	6.748119	7.418830
## 412	7.549440	7.106147
## 422	6.522449	6.655672
## 437	6.522449	6.655672
## 452	6.511225	6.737557
## 467	7.931425	8.121772
## 482	6.522449	6.655672
## 497	6.748119	7.418830
## 512	14.155982	13.910041
## 521	14.330268	14.161250
## 530	6.511225	6.737557
## 545	11.386529	9.754081
## 551	6.522449	6.655672
## 566	12.655145	12.470038
## 575	14.330268	14.161250
## 584	8.907780	8.896563
## 594	7.931425	8.121772
## 609	13.505737	13.308149
## 618	13.505737	13.308149
## 627	6.206479	6.885510
## 642	12.360358	11.238626
## 648	15.405375	15.169967
## 657	6.511225	6.737557
## 672	6.856325	6.516730
## 682	7.876741	7.615284
## 692	6.748119	7.418830
##	GSM2685734_CoV_48hr_Group_B	GSM2685735_CoV_48hr_Group_C
## 1	6.171151	6.220090
## 16	7.624817	7.440707
## 31	12.930745	13.184153
## 41	12.477358	12.738455
## 51	7.609460	7.428115
## 66	6.675679	6.538291
## 81	15.811264	15.943282
## 90	13.268812	12.955313
## 99	7.377139	7.650819
## 109	7.377139	7.650819

## 209	8.807600	8.831738
## 219	5.916103	5.924204
## 279	11.011476	11.453377
## 285	8.807600	8.831738
## 385	11.011476	11.453377
## 391	9.813139	9.731903
## 397	7.609460	7.428115
## 412	7.060422	7.233038
## 422	6.824505	6.832745
## 437	6.824505	6.832745
## 452	6.675679	6.895957
## 467	8.274663	8.323442
## 482	6.824505	6.832745
## 497	7.609460	7.428115
## 512	13.891601	13.584385
## 521	14.260434	14.084852
## 530	6.675679	6.895957
## 545	9.813139	9.731903
## 551	6.824505	6.832745
## 566	12.385422	12.066142
## 575	14.260434	14.084852
## 584	8.807600	8.831738
## 594	8.274663	8.323442
## 609	13.268812	12.955313
## 618	13.268812	12.955313
## 627	7.301030	6.822849
## 642	11.011476	11.453377
## 648	15.204887	15.148994
## 657	6.675679	6.895957
## 672	6.489574	6.563411
## 682	7.377139	7.650819
## 692	7.609460	7.428115
##	GSM2685736_CoV_48hr_Group_D	GSM2685737_CoV_48hr_Group_E
## 1	6.193563	6.168183
## 16	7.487355	7.538375
## 31	13.418502	13.056731
## 41	12.884399	12.555566
## 51	7.451626	7.479294
## 66	6.562724	6.449183
## 81	15.943282	15.770868
## 90	12.889978	13.048290
## 99	7.822518	7.494732
## 109	7.822518	7.494732
## 209	8.962595	8.838498
## 219	5.724779	5.929894
## 279	11.408702	11.297811
## 285	8.962595	8.838498
## 385	11.408702	11.297811
## 391	9.693743	9.734166
## 397	7.451626	7.479294

## 412	7.263597	7.147329
## 422	7.032002	6.806906
## 437	7.032002	6.806906
## 452	6.819596	6.878967
## 467	8.479178	8.383752
## 482	7.032002	6.806906
## 497	7.451626	7.479294
## 512	13.594991	13.642893
## 521	14.281527	14.103762
## 530	6.819596	6.878967
## 545	9.693743	9.734166
## 551	7.032002	6.806906
## 566	12.290165	12.237340
## 575	14.281527	14.103762
## 584	8.962595	8.838498
## 594	8.479178	8.383752
## 609	12.889978	13.048290
## 618	12.889978	13.048290
## 627	7.001776	6.854082
## 642	11.408702	11.297811
## 648	15.181885	15.077245
## 657	6.819596	6.878967
## 672	6.604914	6.551078
## 682	7.822518	7.494732
## 692	7.451626	7.479294
##	GSM2685738_ctrl_48hr_Group_A	GSM2685739_ctrl_48hr_Group_B
## 1	6.113510	6.118294
## 16	7.351580	7.255082
## 31	13.479785	13.809171
## 41	12.913809	13.272047
## 51	6.884900	6.755348
## 66	6.132513	6.191788
## 81	15.840441	15.823487
## 90	13.642893	13.424247
## 99	7.822295	7.826527
## 109	7.822295	7.826527
## 209	8.661194	8.709415
## 219	5.805983	6.059503
## 279	12.087007	12.095376
## 285	8.661194	8.709415
## 385	12.087007	12.095376
## 391	11.252360	11.380947
## 397	6.884900	6.755348
## 412	7.482716	7.507635
## 422	6.651517	6.676103
## 437	6.651517	6.676103
## 452	6.429698	6.527384
## 467	7.880480	7.940615
## 482	6.651517	6.676103
## 497	6.884900	6.755348

## 512	14.218930	14.054635
## 521	14.335814	14.307798
## 530	6.429698	6.527384
## 545	11.252360	11.380947
## 551	6.651517	6.676103
## 566	12.669945	12.641231
## 575	14.335814	14.307798
## 584	8.661194	8.709415
## 594	7.880480	7.940615
## 609	13.642893	13.424247
## 618	13.642893	13.424247
## 627	6.229315	6.288708
## 642	12.087007	12.095376
## 648	15.478388	15.289801
## 657	6.429698	6.527384
## 672	6.768088	6.790796
## 682	7.822295	7.826527
## 692	6.884900	6.755348
##	GSM2685740_ctrl_48hr_Group_C	GSM2685741_ctrl_48hr_Group_D
## 1	6.178938	6.181525
## 16	7.147917	7.127182
## 31	13.768155	13.649614
## 41	13.290203	13.167372
## 51	6.925081	6.742202
## 66	6.108500	6.154514
## 81	15.770868	15.786439
## 90	13.695565	13.390078
## 99	7.672054	7.855158
## 109	7.672054	7.855158
## 209	8.469137	8.748683
## 219	5.850421	6.065332
## 279	12.205447	12.108098
## 285	8.469137	8.748683
## 385	12.205447	12.108098
## 391	11.440833	11.395297
## 397	6.925081	6.742202
## 412	7.413660	7.568577
## 422	6.617604	6.551694
## 437	6.617604	6.551694
## 452	6.314647	6.477990
## 467	7.844684	7.929474
## 482	6.617604	6.551694
## 497	6.925081	6.742202
## 512	14.253092	13.973822
## 521	14.345264	14.221843
## 530	6.314647	6.477990
## 545	11.440833	11.395297
## 551	6.617604	6.551694
## 566	12.795429	12.503914
## 575	14.345264	14.221843

## 584	8.469137	8.748683
## 594	7.844684	7.929474
## 609	13.695565	13.390078
## 618	13.695565	13.390078
## 627	6.225850	6.136620
## 642	12.205447	12.108098
## 648	15.471859	15.279920
## 657	6.314647	6.477990
## 672	6.867602	6.882945
## 682	7.672054	7.855158
## 692	6.925081	6.742202
## GSM2685742_ctrl_48hr_Group_E		
## 1	6.050870	
## 16	7.342277	
## 31	13.621482	
## 41	13.121641	
## 51	6.787927	
## 66	6.120162	
## 81	15.815665	
## 90	13.471967	
## 99	7.714570	
## 109	7.714570	
## 209	8.781281	
## 219	5.903015	
## 279	12.159429	
## 285	8.781281	
## 385	12.159429	
## 391	11.449781	
## 397	6.787927	
## 412	7.398872	
## 422	6.586001	
## 437	6.586001	
## 452	6.458481	
## 467	7.953244	
## 482	6.586001	
## 497	6.787927	
## 512	14.119631	
## 521	14.228603	
## 530	6.458481	
## 545	11.449781	
## 551	6.586001	
## 566	12.656605	
## 575	14.228603	
## 584	8.781281	
## 594	7.953244	
## 609	13.471967	
## 618	13.471967	
## 627	6.282451	
## 642	12.159429	
## 648	15.275427	

## 657	6.458481
## 672	6.783099
## 682	7.714570
## 692	6.787927

The above table shows the five genes that had the highest count and then the genotypes within those five genes by number or count of those genotypes of DNA copy number variations in the SEQUENCE.

It would be interesting to analyze those sequences of copy number variations within the top 5 genes expressed the most number of times in this data. Comparing networks of genes associated with processes in the body like immune response, pathogenesis of disease onset, networks of human processes in the body associated with cancer or subsequent diseases like autoimmune, celiac disease, hemochromatosis, anemia, etc. To see how well any of those genotypes fair. We could also detect patterns in the fold change values between genotypes of the genes expressed in comparing the capillary samples all within 1 hour after being inoculated with CoV, inactive CoV, CoV and an interleukin alpha, or control group. Then compare the liver tumor samples of each group A through E that was monitored after being inoculated with CoV at 0,12,24,36, and 48 hours side by side with the control groups of groups A through E.
