

# RNA Seq DEG Analysis

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```
library(tidyverse)
library(DESeq2)
library(dplyr)
```

Create count matrix

```
# List all rsem count files
files <- list.files(path = "results/counts/rsem/", pattern = "*.genes.results",
                     full.names = T)
head(files)
```

```
## [1] "results/counts/rsem/DRR608910.genes.results"
## [2] "results/counts/rsem/DRR608911.genes.results"
## [3] "results/counts/rsem/DRR608912.genes.results"
## [4] "results/counts/rsem/DRR608913.genes.results"
## [5] "results/counts/rsem/DRR608914.genes.results"
## [6] "results/counts/rsem/DRR608915.genes.results"
```

```
# Extract sample names from filenames
sample_names <- gsub("\\.genes\\.results", "", basename(files))
head(sample_names)
```

```
## [1] "DRR608910" "DRR608911" "DRR608912" "DRR608913" "DRR608914" "DRR608915"
```

```
# Extract gene_id and expected count
count_list <- lapply(files, function(f){
  df <- read.delim(f, stringsAsFactors = F)
  df[, c("gene_id", "expected_count")]
})
```

```
#count_list
```

```
#str(count_list)
```

```
count_matrix <- Reduce(function(x, y) merge(x, y, by = "gene_id"), count_list)
head(count_matrix)
```

```
##      gene_id expected_count.x expected_count.y expected_count.x expected_count.y
```

```

## 1      A1BG        3.00        2.00        3.13        5.07
## 2 A1BG-AS1     7.63        4.87       12.46       38.54
## 3      A1CF        0.00        0.00        0.00        0.00
## 4      A2M       7157.31      6252.89      1872.42      5358.19
## 5 A2M-AS1      13.69        3.11        8.58       13.81
## 6      A2ML1     50819.00     43442.00     60477.00     42791.00
##   expected_count.x expected_count.y expected_count.x expected_count.y
## 1             1.00          8.48         10.26         7.00
## 2            23.31         16.60         3.71        13.99
## 3             0.00          0.00          0.00          0.00
## 4            7791.72        14466.00        1842.00        8931.50
## 5             8.28          20.00          5.00        25.50
## 6            29474.00        16737.00        32336.00        36409.00
##   expected_count.x expected_count.y expected_count.x expected_count.y
## 1             11.00         34.11         17.00        51.00
## 2            20.29         35.31         23.49        19.18
## 3             0.00          0.00          0.00          0.00
## 4            12060.55        5121.00        4124.00        1927.35
## 5            13.45          15.00          11.00         7.65
## 6            4687.00         150.00         2757.00        8513.00
##   expected_count.x expected_count.y expected_count.x expected_count.y
## 1             67.33         44.44         17.0          17.28
## 2            47.05         17.13          0.0        26.97
## 3             0.00          0.00          0.0          0.00
## 4            2503.00         644.20        3165.6        2406.00
## 5             6.00          28.80          17.4        30.00
## 6            1316.00         754.00         344.0        3815.00

```

```

colnames(count_matrix) <- c("gene_id", sample_names)

head(count_matrix)

```

```

##   gene_id DRR608910 DRR608911 DRR608912 DRR608913 DRR608914 DRR608915
## 1      A1BG      3.00      2.00      3.13      5.07      1.00      8.48
## 2 A1BG-AS1     7.63      4.87     12.46     38.54     23.31     16.60
## 3      A1CF      0.00      0.00      0.00      0.00      0.00      0.00
## 4      A2M     7157.31     6252.89     1872.42     5358.19     7791.72    14466.00
## 5 A2M-AS1     13.69      3.11      8.58     13.81      8.28     20.00
## 6      A2ML1    50819.00    43442.00    60477.00    42791.00    29474.00    16737.00
##   DRR608916 DRR608917 DRR608918 DRR608919 DRR608920 DRR608921 DRR608922
## 1      10.26      7.00     11.00     34.11     17.00      51.00     67.33
## 2      3.71     13.99     20.29     35.31     23.49     19.18     47.05
## 3      0.00      0.00      0.00      0.00      0.00      0.00      0.00
## 4     1842.00    8931.50   12060.55    5121.00    4124.00    1927.35   2503.00
## 5      5.00     25.50     13.45     15.00     11.00      7.65      6.00
## 6    32336.00   36409.00    4687.00    150.00    2757.00    8513.00   1316.00
##   DRR608923 DRR608924 DRR608925
## 1      44.44      17.0     17.28
## 2      17.13      0.0     26.97
## 3      0.00      0.0     0.00
## 4     644.20     3165.6    2406.00
## 5     28.80      17.4     30.00
## 6     754.00     344.0    3815.00

```

```

# Add metadata with the conditions of each sample
metadata <- data.frame(
  row.names = sample_names,
  condition = c("normal", "normal", "normal", "normal", "normal", "normal", "normal",
    "met_canc", "met_canc", "met_canc", "met_canc", "met_canc", "met_canc", "met_canc",
    "met_canc")
)

metadata

```

```

##           condition
## DRR608910      normal
## DRR608911      normal
## DRR608912      normal
## DRR608913      normal
## DRR608914      normal
## DRR608915      normal
## DRR608916      normal
## DRR608917      normal
## DRR608918    met_canc
## DRR608919    met_canc
## DRR608920    met_canc
## DRR608921    met_canc
## DRR608922    met_canc
## DRR608923    met_canc
## DRR608924    met_canc
## DRR608925    met_canc

```

### Create DESeq object

```

dds <- DESeqDataSetFromMatrix(
  countData = round(as.matrix(count_matrix[, -1])),
  colData = metadata,
  design = ~ condition
)

head(dds)

```

```

## class: DESeqDataSet
## dim: 6 16
## metadata(1): version
## assays(1): counts
## rownames: NULL
## rowData names(0):
## colnames(16): DRR608910 DRR608911 ... DRR608924 DRR608925
## colData names(1): condition

dim(dds)

```

```

## [1] 50037    16

```

```

colnames(dds)

## [1] "DRR608910" "DRR608911" "DRR608912" "DRR608913" "DRR608914" "DRR608915"
## [7] "DRR608916" "DRR608917" "DRR608918" "DRR608919" "DRR608920" "DRR608921"
## [13] "DRR608922" "DRR608923" "DRR608924" "DRR608925"

# Test whether there are incorrect metadata sample assignments during submissions to NCBI SRA

# Estimate size factors
dds <- estimateSizeFactors(dds)
sizeFactors(dds)

## DRR608910 DRR608911 DRR608912 DRR608913 DRR608914 DRR608915 DRR608916 DRR608917
## 0.9634680 0.7605656 0.6679232 0.8841481 0.9894707 1.0890224 0.7960631 0.8602211
## DRR608918 DRR608919 DRR608920 DRR608921 DRR608922 DRR608923 DRR608924 DRR608925
## 1.2825801 1.1538479 0.9634154 1.1519050 1.2546769 1.3179475 1.1808321 1.1651657

# counts(dds)
log.norm <- normTransform(dds)
log.norm

## class: DESeqTransform
## dim: 50037 16
## metadata(1): version
## assays(1): ''
## rownames: NULL
## rowData names(0):
## colnames(16): DRR608910 DRR608911 ... DRR608924 DRR608925
## colData names(2): condition sizeFactor

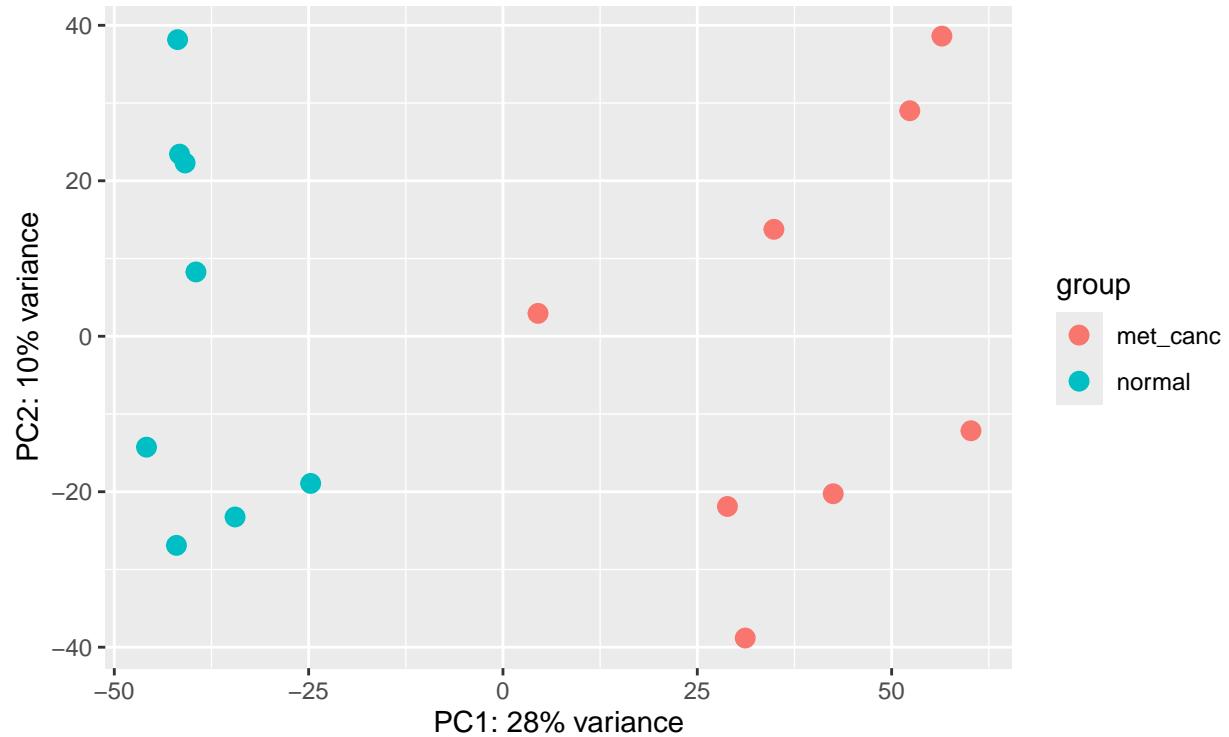
log.norm.counts <- log2(counts(dds, normalized = T) + 1)
head(log.norm.counts)

##          DRR608910 DRR608911 DRR608912 DRR608913 DRR608914 DRR608915 DRR608916
## [1,]  2.040455  1.859819  2.457209  2.734474  1.007656  3.061092  3.761479
## [2,]  3.217748  2.921066  4.245354  5.495384  4.599600  4.054009  2.590896
## [3,]  0.000000  0.000000  0.000000  0.000000  0.000000  0.000000  0.000000
## [4,] 12.859025 13.005364 11.453125 12.565357 12.943232 13.697453 11.176726
## [5,]  3.957064  2.305805  3.855452  4.073345  3.183507  4.275387  2.864119
## [6,] 15.686799 15.801684 16.466361 15.562690 14.862475 13.907814 15.309934
##          DRR608917 DRR608918 DRR608919 DRR608920 DRR608921 DRR608922 DRR608923
## [1,]  3.191790  3.259493  4.929158  4.220760  5.500626  5.765540  5.103718
## [2,]  4.110604  4.052552  4.969624  4.636532  4.128823  5.265281  3.796893
## [3,]  0.000000  0.000000  0.000000  0.000000  0.000000  0.000000  0.000000
## [4,] 13.342127 13.199166 12.116082 12.063936 10.708981 10.962850  8.935573
## [5,]  4.964620  3.477136  3.807353  3.634328  2.990050  2.531595  4.523807
## [6,] 15.369262 11.835795  7.033421 11.483158 12.851625 10.036003  9.162647
##          DRR608924 DRR608925
## [1,]  3.944542  3.962567
## [2,]  0.000000  4.595305
## [3,]  0.000000  0.000000

```

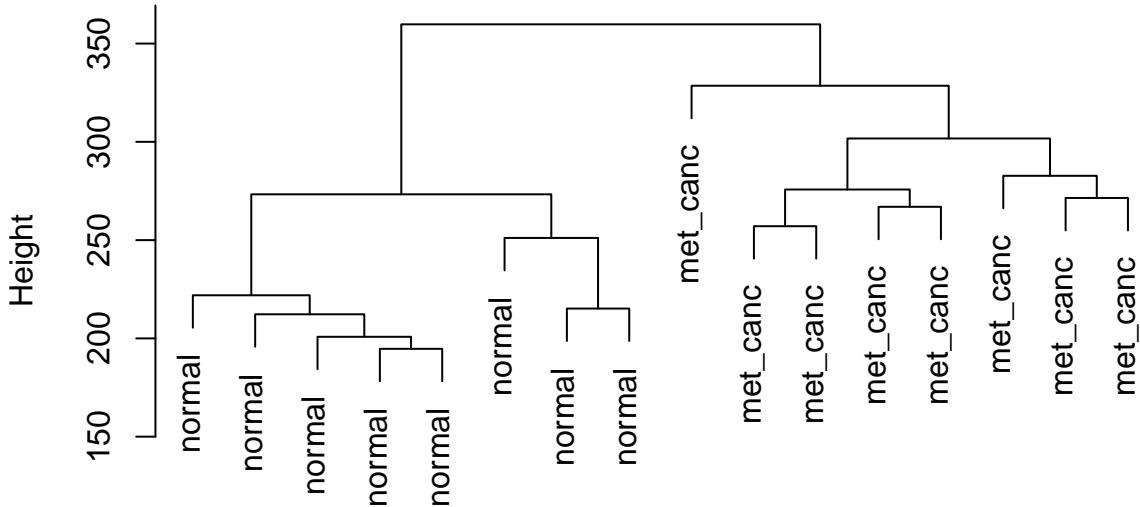
```
## [4,] 11.389180 11.012584  
## [5,] 3.944542  4.741327  
## [6,] 8.191405  11.677373
```

```
# Carryout plot PCA and hclust to verify that the samples group into their assigned conditions  
plotPCA(log.norm, intgroup = "condition")
```



```
plot(hclust(dist(t(log.norm.counts))), labels = colData(dds)$condition)
```

## Cluster Dendrogram



```
dist(t(log.norm.counts))
hclust (*, "complete")
```

```
# Add gene_ids to the rownames of dds

# count_matrix$gene_id
rownames(dds) <- count_matrix$gene_id

# rownames(dds)

# Rearrange the levels such that the comparison is done against the non-cancerous
# subjects
levels(dds$condition)

## [1] "met_canc" "normal"

dds$condition <- relevel(dds$condition, "normal")
levels(dds$condition)

## [1] "normal"    "met_canc"

# Filter out genes that have very low counts in at least 2 samples which don't contribute any significant information
filtered <- rowSums(counts(dds) >= 10) > 2
dds <- dds[filtered, ]

# dds
```

```

# Run DESeq
dds <- DESeq(dds)

# Create a dataframe with the gene symbol, p values, adjusted p values and logFC of each DEG
res_lfc <- results(dds, lfcThreshold = 1)
summary(res_lfc)

## 
## out of 20668 with nonzero total read count
## adjusted p-value < 0.1
## LFC > 1.00 (up) : 813, 3.9%
## LFC < -1.00 (down) : 624, 3%
## outliers [1] : 0, 0%
## low counts [2] : 0, 0%
## (mean count < 0)
## [1] see 'cooksCutoff' argument of ?results
## [2] see 'independentFiltering' argument of ?results

str(res_lfc)

## Formal class 'DESeqResults' [package "DESeq2"] with 7 slots
## ..@ priorInfo      : list()
## ..@ rownames       : chr [1:20668] "A1BG" "A1BG-AS1" "A2M" "A2M-AS1" ...
## ..@ nrows          : int 20668
## ..@ elementType    : chr "ANY"
## ..@ elementMetadata:Formal class 'DFrame' [package "S4Vectors"] with 6 slots
## ... ..@ rownames     : NULL
## ... ..@ nrows        : int 6
## ... ..@ elementType   : chr "ANY"
## ... ..@ elementMetadata: NULL
## ... ..@ metadata      : list()
## ... ..@ listData      :List of 2
## ... ...$ type        : chr [1:6] "intermediate" "results" "results" "results" ...
## ... ...$ description: chr [1:6] "mean of normalized counts for all samples" "log2 fold change"
## ..@ metadata        :List of 6
## ...$ filterThreshold: Named num 0.109
## ... ..- attr(*, "names")= chr "0%"
## ...$ filterTheta     : num 0
## ...$ filterNumRej    :'data.frame': 50 obs. of 2 variables:
## ... ...$ theta : num [1:50] 0 0.0194 0.0388 0.0582 0.0776 ...
## ... ...$ numRej: num [1:50] 1437 1402 1375 1335 1311 ...
## ...$ lo.fit         :List of 2
## ... ...$ x: num [1:50] 0 0.0194 0.0388 0.0582 0.0776 ...
## ... ...$ y: num [1:50] 1436 1404 1372 1341 1309 ...
## ...$ alpha          : num 0.1
## ...$ lfcThreshold   : num 1
## ..@ listData        :List of 6
## ...$ baseMean        : num [1:20668] 16.3 18.6 5337.1 13.7 24758.2 ...
## ...$ log2FoldChange: num [1:20668] 2.2686 0.218 -1.1109 -0.0352 -4.2933 ...
## ...$ lfcSE           : num [1:20668] 0.514 0.58 0.47 0.461 0.591 ...
## ...$ stat            : num [1:20668] 2.468 0 -0.236 0 -5.568 ...
## ...$ pvalue          : num [1:20668] 1.36e-02 1.00 8.13e-01 1.00 2.57e-08 ...
## ...$ padj            : num [1:20668] 1.70e-01 1.00 1.00 1.00 2.89e-06 ...

```

```

res_lfc_padj_0.05 <- res_lfc[res_lfc$padj < 0.05, ]
str(res_lfc_padj_0.05)

## Formal class 'DESeqResults' [package "DESeq2"] with 7 slots
## ..@ priorInfo      : list()
## ..@ rownames       : chr [1:1192] "A2ML1" "ABCA13" "ABCC2" "ABCG4" ...
## ..@ nrows         : int 1192
## ..@ elementType    : chr "ANY"
## ..@ elementMetadata:Formal class 'DFrame' [package "S4Vectors"] with 6 slots
## ... ..@ rownames     : NULL
## ... ..@ nrows        : int 6
## ... ..@ elementType   : chr "ANY"
## ... ..@ elementMetadata: NULL
## ... ..@ metadata      : list()
## ... ..@ listData     :List of 2
## ... ...$ type        : chr [1:6] "intermediate" "results" "results" "results" ...
## ... ...$ description: chr [1:6] "mean of normalized counts for all samples" "log2 fold change"
## ..@ metadata       :List of 6
## ... $ filterThreshold: Named num 0.109
## ... ..- attr(*, "names")= chr "0%"
## ... $ filterTheta    : num 0
## ... $ filterNumRej   :'data.frame': 50 obs. of 2 variables:
## ... ...$ theta : num [1:50] 0 0.0194 0.0388 0.0582 0.0776 ...
## ... ...$ numRej: num [1:50] 1437 1402 1375 1335 1311 ...
## ... $ lo.fit       :List of 2
## ... ...$ x: num [1:50] 0 0.0194 0.0388 0.0582 0.0776 ...
## ... ...$ y: num [1:50] 1436 1404 1372 1341 1309 ...
## ... $ alpha        : num 0.1
## ... $ lfcThreshold  : num 1
## ..@ listData      :List of 6
## ... $ baseMean      : num [1:1192] 24758 373 132 29 6872 ...
## ... $ log2FoldChange: num [1:1192] -4.29 3.35 4.39 -2.72 -2.43 ...
## ... $ lfcSE         : num [1:1192] 0.591 0.677 0.822 0.5 0.342 ...
## ... $ stat          : num [1:1192] -5.57 3.46 4.13 -3.45 -4.18 ...
## ... $ pvalue        : num [1:1192] 2.57e-08 5.32e-04 3.68e-05 5.56e-04 2.93e-05 ...
## ... $ padj          : num [1:1192] 2.89e-06 1.25e-02 1.38e-03 1.30e-02 1.13e-03 ...

```

```

deg_list <- rownames(res_lfc_padj_0.05)

degs <- data.frame(
  gene_id = rownames(res_lfc_padj_0.05),
  P.Value = res_lfc_padj_0.05$pvalue,
  adj.P.Val = res_lfc_padj_0.05$padj,
  logFC = res_lfc_padj_0.05$log2FoldChange
)
degs

```

	gene_id	P.Value	adj.P.Val	logFC
## 1	A2ML1	2.573314e-08	2.890503e-06	-4.293321
## 2	ABCA13	5.316538e-04	1.247244e-02	3.345934
## 3	ABCC2	3.676964e-05	1.384253e-03	4.392891

## 4	ABCG4	5.561698e-04	1.297395e-02	-2.724376
## 5	ABLIM1	2.930047e-05	1.129817e-03	-2.430274
## 6	ABLIM3	7.328133e-05	2.470764e-03	-3.245304
## 7	ACER1	1.425965e-08	1.723499e-06	-5.248480
## 8	ACKR1	4.453428e-05	1.614797e-03	-2.736071
## 9	ACKR3	2.758081e-03	4.851406e-02	2.305740
## 10	ACOT11	2.096467e-03	3.882596e-02	-2.159534
## 11	ACOX1	1.412286e-04	4.256241e-03	-1.731111
## 12	ACOX2	1.617922e-05	6.908930e-04	-2.729130
## 13	ACOX3	6.520376e-06	3.352317e-04	-2.288334
## 14	ACP3	2.678986e-04	7.184753e-03	-2.778791
## 15	ACTG2	4.499720e-14	2.268298e-11	-4.835173
## 16	ACVR1C	1.389269e-05	6.180592e-04	3.812397
## 17	ADAM12	5.332085e-05	1.887047e-03	3.635348
## 18	ADAM23	6.673881e-08	6.728575e-06	3.126805
## 19	ADAM32_1	2.425635e-03	4.359393e-02	3.392516
## 20	ADAM33	1.302545e-04	4.000149e-03	-2.710397
## 21	ADAMTS12	1.536425e-07	1.362868e-05	4.218968
## 22	ADAMTS14	1.863847e-04	5.328075e-03	2.976114
## 23	ADAMTS7	4.132060e-04	1.021548e-02	2.388743
## 24	ADAMTS8	3.909642e-07	2.938344e-05	-4.018085
## 25	ADAMTS9-AS1	1.300500e-14	7.679641e-12	-6.334097
## 26	ADAMTS9-AS2	5.473167e-04	1.279631e-02	-3.608809
## 27	ADCY5	6.178884e-07	4.271076e-05	-2.934183
## 28	ADD2	3.880832e-05	1.455699e-03	3.187843
## 29	ADGRF2P	1.471048e-03	2.920616e-02	-2.664184
## 30	ADGRG6	2.802716e-03	4.909029e-02	-1.836851
## 31	ADH1B	5.770564e-13	2.208630e-10	-5.267329
## 32	AFDN	2.331455e-04	6.412398e-03	-2.066169
## 33	AFF3	1.069216e-09	1.753853e-07	-3.039541
## 34	AGFG2	2.544316e-08	2.873548e-06	-2.589640
## 35	AGTR1	2.876195e-03	4.987014e-02	-2.970563
## 36	AIF1L	5.478900e-05	1.922545e-03	-2.870940
## 37	AKR1B1	1.570847e-03	3.059801e-02	2.135055
## 38	AKR1C1	4.519830e-04	1.100304e-02	3.726399
## 39	AKR1C2	5.637977e-05	1.965020e-03	3.385866
## 40	AKR1C3	6.955118e-04	1.559093e-02	3.769443
## 41	ALDH9A1	4.083514e-04	1.014400e-02	-1.673958
## 42	ALG1L1P	4.285828e-10	7.770131e-08	6.114394
## 43	ALOX12	1.158416e-08	1.442054e-06	-4.441886
## 44	ALOX12P2	1.217839e-08	1.498232e-06	3.406411
## 45	ALOX15	1.142238e-09	1.830061e-07	6.651260
## 46	ALX1	6.187678e-06	3.205186e-04	5.859546
## 47	AMOTL2	5.682106e-04	1.319526e-02	-2.116939
## 48	ANGPTL1	3.320303e-10	6.182345e-08	-3.884158
## 49	ANKRD19P	6.541929e-04	1.474467e-02	2.808824
## 50	ANKRD20A11P	3.557681e-04	9.044299e-03	-4.221272
## 51	ANKRD20A5P	1.247145e-04	3.852913e-03	-3.993332
## 52	ANKRD20A9P	1.403292e-04	4.246449e-03	-4.154720
## 53	ANKRD20A9P_1	1.403292e-04	4.246449e-03	-4.154720
## 54	ANKRD22	1.012590e-03	2.131182e-02	-2.053582
## 55	ANKRD34B	1.632805e-04	4.807238e-03	5.244232
## 56	ANO1	4.714316e-04	1.135612e-02	3.208133
## 57	ANXA10	1.577296e-03	3.066750e-02	5.729620

## 58	ANXA9	2.183325e-04	6.073346e-03	-3.105041
## 59	AOC3	1.518325e-03	2.994346e-02	-2.395572
## 60	AOX1	3.493113e-11	8.493607e-09	-4.180983
## 61	APLN	1.894643e-03	3.582660e-02	2.670513
## 62	APOC1	1.128272e-05	5.244467e-04	3.831572
## 63	AQP5	1.182394e-03	2.438021e-02	-3.573483
## 64	AQP7P1	4.849868e-05	1.737356e-03	-4.143373
## 65	ARHGAP10	1.182649e-10	2.468988e-08	-2.823413
## 66	ARHGAP27	3.900443e-04	9.771438e-03	-2.180975
## 67	ARHGAP27_2	1.780591e-03	3.388698e-02	-2.201105
## 68	ARHGEF10L	1.459662e-06	9.197650e-05	-2.623614
## 69	ARSF	4.766978e-05	1.713459e-03	-3.784077
## 70	ARTN	3.176918e-07	2.468441e-05	4.082105
## 71	ASB5	7.125052e-11	1.533964e-08	-6.676772
## 72	ASCL2	3.084618e-04	8.009157e-03	2.140080
## 73	ASPA	7.047749e-05	2.403678e-03	-3.541027
## 74	ASPG	9.079565e-04	1.963244e-02	-2.698803
## 75	ASPN_1	1.383102e-03	2.780591e-02	3.056957
## 76	ATG9B	1.247374e-03	2.534979e-02	-3.059404
## 77	ATOH8	2.180972e-03	4.013921e-02	-2.179397
## 78	ATP13A4	2.842360e-03	4.944941e-02	-2.644112
## 79	ATP1A2	1.253333e-27	8.634627e-24	-5.722853
## 80	AUNIP	7.373391e-04	1.633368e-02	2.291721
## 81	AURKA	1.131695e-04	3.533435e-03	2.334947
## 82	AURKB	5.455517e-04	1.276949e-02	2.031322
## 83	B3GNT8	1.930046e-03	3.639615e-02	-2.470325
## 84	B4GALNT1	1.430662e-17	1.642718e-14	4.803579
## 85	B4GALNT4	6.173947e-04	1.406577e-02	2.480870
## 86	BAMBI	2.267190e-04	6.264477e-03	3.335898
## 87	BARX1	5.179918e-11	1.176468e-08	7.907641
## 88	BARX2	1.119797e-04	3.506662e-03	-3.334623
## 89	BBOX1-AS1	1.693712e-04	4.958306e-03	3.418432
## 90	BCAN	2.670079e-08	2.982983e-06	4.563505
## 91	BCAS1	3.397841e-04	8.723799e-03	-2.567226
## 92	BICD1	1.525518e-03	3.005663e-02	1.915596
## 93	BIRC5	9.998321e-04	2.112938e-02	2.027227
## 94	BLNK	1.616126e-03	3.121691e-02	-2.391013
## 95	BMP8A	2.447328e-03	4.386936e-02	2.355312
## 96	BMPR1B	1.092653e-03	2.271926e-02	3.272940
## 97	BMS1P14	3.433392e-04	8.788536e-03	-2.667681
## 98	BMS1P23	1.610046e-04	4.746995e-03	-2.611368
## 99	BNIPL	2.892537e-04	7.625377e-03	-2.758131
## 100	BOC	1.705687e-04	4.979256e-03	-2.485833
## 101	BPIFB2	2.727619e-03	4.818327e-02	-7.359250
## 102	C15orf40	2.030561e-10	3.959211e-08	3.969106
## 103	C15orf62	1.435834e-05	6.314003e-04	-3.079208
## 104	C16orf74	7.270915e-05	2.463529e-03	2.810195
## 105	C1orf116	2.337206e-11	6.038172e-09	-3.698696
## 106	C1QTNF6	1.727401e-03	3.305734e-02	2.515549
## 107	C3orf70	2.929883e-04	7.679951e-03	-2.720467
## 108	C6orf132	2.440742e-03	4.378929e-02	-2.112112
## 109	C6orf15	7.098018e-04	1.580771e-02	5.588939
## 110	C6orf15_2	7.098018e-04	1.580771e-02	5.588939
## 111	C6orf15_3	7.098018e-04	1.580771e-02	5.588939

## 112	C6orf15_4	7.098018e-04	1.580771e-02	5.588939
## 113	C6orf15_5	6.402224e-04	1.447101e-02	8.584877
## 114	C7	6.192494e-21	1.599831e-17	-5.700613
## 115	C7orf57	2.406772e-03	4.329258e-02	-3.194505
## 116	CA13	4.409623e-05	1.601724e-03	-2.574776
## 117	CA9	5.437573e-07	3.902214e-05	7.002867
## 118	CAB39L	5.270562e-06	2.793128e-04	-2.557420
## 119	CABYR	4.702806e-07	3.446723e-05	3.815934
## 120	CACNA1B	2.196744e-14	1.227089e-11	7.311116
## 121	CACNB2	1.105349e-08	1.393010e-06	-3.233534
## 122	CACNG4	9.425408e-06	4.478261e-04	4.026712
## 123	CADM3	1.567751e-06	9.848716e-05	-3.750337
## 124	CALB1	4.769728e-04	1.144956e-02	4.569876
## 125	CAMK2N2	2.916652e-04	7.669384e-03	2.880172
## 126	CAPN14	1.572241e-03	3.059801e-02	-3.417891
## 127	CASC9	6.248353e-11	1.395881e-08	6.739083
## 128	CASQ2	4.079049e-09	5.620385e-07	-4.549791
## 129	CBS	2.357933e-05	9.481277e-04	3.373427
## 130	CCDC162P	8.330274e-12	2.295601e-09	4.473031
## 131	CCDC187	2.202148e-03	4.034929e-02	4.401733
## 132	CCL14	2.594485e-09	3.803037e-07	-3.920689
## 133	CCL14_1	2.594485e-09	3.803037e-07	-3.920689
## 134	CCL20	3.797190e-04	9.547485e-03	3.969695
## 135	CCN4	3.745370e-04	9.463241e-03	3.223451
## 136	CCN5	7.291776e-07	4.973810e-05	-3.655769
## 137	CCNA1	9.727653e-07	6.382576e-05	5.177291
## 138	CCNP	1.802282e-06	1.098807e-04	5.796424
## 139	CD1E	7.393371e-04	1.634178e-02	-3.201493
## 140	CD207	1.851222e-05	7.729504e-04	-3.468188
## 141	CD276	1.382656e-04	4.205689e-03	2.302272
## 142	CD300LG	8.900422e-09	1.142571e-06	-5.789410
## 143	CD83	2.299800e-03	4.184178e-02	1.943130
## 144	CDC45	4.965947e-06	2.650752e-04	2.018231
## 145	CDC6	2.503944e-06	1.466048e-04	2.646758
## 146	CDCA3	1.561002e-03	3.052299e-02	2.057956
## 147	CDCA5	5.011337e-04	1.183706e-02	2.009932
## 148	CDH11	5.625934e-04	1.309738e-02	2.886782
## 149	CDH19	1.690302e-04	4.955343e-03	-2.996838
## 150	CDKN3	7.529065e-06	3.731672e-04	3.185739
## 151	CDSN_3	9.790002e-04	2.077410e-02	5.565901
## 152	CDSN_4	9.790002e-04	2.077410e-02	5.565901
## 153	CDX2	4.993953e-04	1.180950e-02	6.244599
## 154	CEACAM1	1.336355e-05	6.004299e-04	-3.187936
## 155	CEACAM7	1.580903e-04	4.687819e-03	-3.955103
## 156	CECR2	5.511930e-04	1.287238e-02	2.898726
## 157	CENPA	1.580780e-05	6.816360e-04	2.440273
## 158	CENPF	1.048784e-04	3.329688e-03	2.116781
## 159	CENPH	1.474666e-03	2.924990e-02	1.894062
## 160	CENPM	1.972823e-03	3.689982e-02	2.001024
## 161	CENPV	2.746832e-03	4.839857e-02	2.332087
## 162	CEP55	1.507467e-03	2.978616e-02	1.776009
## 163	CES2	1.042277e-07	9.791722e-06	-2.966992
## 164	CFAP251	1.145070e-10	2.414930e-08	3.881225
## 165	CFAP47	1.214480e-05	5.563847e-04	5.487401

## 166	CFAP53	1.293823e-04	3.979275e-03	3.909695
## 167	CFD	2.173576e-07	1.782677e-05	-2.904503
## 168	CFD_1	2.173576e-07	1.782677e-05	-2.904503
## 169	CGNL1	2.220889e-05	8.965105e-04	-3.158161
## 170	CHGB	1.361649e-03	2.745615e-02	4.803079
## 171	CHODL	1.109903e-03	2.305474e-02	3.226112
## 172	CHP2	2.878407e-04	7.597818e-03	3.586071
## 173	CHRDL1	3.648760e-13	1.479285e-10	-5.102809
## 174	CHRNA1	9.113720e-05	2.968883e-03	5.610142
## 175	CHST2	8.357707e-05	2.754977e-03	2.965152
## 176	CILP	4.409699e-11	1.012663e-08	-4.203096
## 177	CILP2	9.968839e-04	2.111024e-02	2.848603
## 178	CIMAP2	1.793239e-05	7.533062e-04	-3.154746
## 179	CIP2A	1.139882e-04	3.553406e-03	2.312590
## 180	CITED2	4.155031e-04	1.024776e-02	-2.096602
## 181	CKS1B	1.558834e-04	4.649059e-03	2.558733
## 182	CKS2	2.142588e-04	5.984191e-03	3.131849
## 183	CLCA4	4.381302e-07	3.257293e-05	-4.382132
## 184	CLDN17	1.012485e-04	3.224717e-03	-3.949774
## 185	CLEC12A-AS1	1.044762e-13	4.907531e-11	5.095812
## 186	CLEC3B	8.441435e-04	1.842319e-02	-2.769500
## 187	CLIC3	2.924655e-04	7.679951e-03	-3.354401
## 188	CLSPN	3.832839e-06	2.112456e-04	2.324454
## 189	CNFN	9.185480e-08	8.748640e-06	-4.096514
## 190	CNGB1	1.065830e-03	2.220623e-02	4.438166
## 191	CNIH2	1.347200e-03	2.719134e-02	2.766163
## 192	CNN1	2.380517e-16	1.892328e-13	-4.508997
## 193	CNTN5	9.140337e-04	1.974007e-02	4.820636
## 194	CNTNAP2	4.379297e-04	1.074956e-02	3.855783
## 195	CNTNAP2_1	2.212718e-03	4.047120e-02	3.162057
## 196	COCH	1.334577e-08	1.622531e-06	4.014747
## 197	COL10A1	7.151233e-17	6.426160e-14	8.377233
## 198	COL11A1	6.034145e-15	3.708683e-12	8.011539
## 199	COL24A1	2.392427e-04	6.557914e-03	3.178407
## 200	COL27A1	1.563547e-03	3.054385e-02	2.455225
## 201	COL2A1	1.535421e-05	6.652849e-04	6.912131
## 202	COL7A1	8.507781e-04	1.854840e-02	2.132210
## 203	CPEB3	7.424140e-06	3.688944e-04	-2.359492
## 204	CPLX2	4.959148e-04	1.179467e-02	6.436563
## 205	CPNE6	8.028517e-05	2.663457e-03	-5.537337
## 206	CPNE6_1	8.028517e-05	2.663457e-03	-5.537337
## 207	CPXM1	1.422415e-03	2.851452e-02	2.855149
## 208	CRABP1	1.054582e-03	2.199405e-02	5.228691
## 209	CRCT1	4.585677e-07	3.373720e-05	-4.475599
## 210	CRISP2	2.012905e-05	8.303938e-04	-5.036277
## 211	CRISP3	3.130142e-11	7.889485e-09	-5.865594
## 212	CRMA	6.199882e-04	1.406577e-02	-3.599019
## 213	CRNN	6.126899e-05	2.117571e-03	-5.308179
## 214	CRTAC1	1.252044e-03	2.541969e-02	-3.497130
## 215	CRYBG1	1.714009e-06	1.067022e-04	-2.662419
## 216	CRYBG2	3.205208e-04	8.280655e-03	-2.968805
## 217	CSAG2_1	5.076371e-07	3.681348e-05	7.902203
## 218	CSAG3	5.937133e-06	3.090898e-04	7.234290
## 219	CSAG3_1	3.206431e-05	1.220452e-03	6.647558

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## 220      CSF2 1.198029e-03 2.463768e-02  5.552246
## 221      CSRP1 7.114144e-05 2.422325e-03 -2.060090
## 222      CST1 1.920540e-19 3.312485e-16 11.923317
## 223      CST2 1.786405e-04 5.149431e-03  5.555372
## 224      CSTB 3.308605e-07 2.551577e-05 -3.391982
## 225      CT45A5 2.201478e-03 4.034929e-02  5.827646
## 226      CTHRC1 6.071635e-07 4.218224e-05  4.517651
## 227      CTNNA3 2.325376e-04 6.408115e-03 -4.538274
## 228      CTSC 2.057773e-03 3.828087e-02  2.200636
## 229      CTTN-DT 1.712536e-04 4.985167e-03  3.415402
## 230      CTTN-DT_1 1.712536e-04 4.985167e-03  3.415402
## 231      CTTNBP2 1.784483e-07 1.562784e-05 -3.887677
## 232      CXCL1 2.429952e-05 9.751894e-04  4.126697
## 233      CXCL11 8.413262e-05 2.768874e-03  4.337692
## 234      CXCL12 1.174842e-04 3.645892e-03 -2.788505
## 235      CXCL14 4.873978e-04 1.163404e-02  3.026065
## 236      CXCL5 1.442096e-05 6.328077e-04  3.979228
## 237      CXCL6 9.331315e-05 3.013431e-03  3.963683
## 238      CXCR2 4.741313e-10 8.234745e-08 -4.041847
## 239      CYP11A1 7.321300e-05 2.470764e-03 -3.650514
## 240      CYP11A1_1 7.321300e-05 2.470764e-03 -3.650514
## 241      CYP1A1 4.566759e-06 2.464380e-04 -5.160696
## 242      CYP26A1 3.511465e-09 4.937072e-07  9.428985
## 243      CYP2C18 5.194153e-04 1.221306e-02 -2.844339
## 244      CYP2J2 2.231225e-03 4.073760e-02 -2.511379
## 245      CYP4B1 2.680204e-04 7.184753e-03 -3.010827
## 246      CYP4F12 7.127442e-05 2.422861e-03 -2.717821
## 247      CYP4F22 3.382853e-06 1.894764e-04 -4.389495
## 248      CYP4F29P 2.707595e-04 7.230049e-03 -4.192720
## 249      CYP4F35P 5.144243e-04 1.210948e-02 -4.083722
## 250      CYSRT1 1.583052e-05 6.816360e-04 -3.877197
## 251      DAAM2 1.033146e-05 4.875128e-04 -2.827132
## 252      DAPP1 4.104112e-04 1.015854e-02 -2.358937
## 253      DEFB4B_2 1.225680e-03 2.510640e-02  4.883737
## 254      DEGS2 9.447866e-07 6.258606e-05 -2.898118
## 255      DENND2C 2.949814e-04 7.707555e-03 -2.147567
## 256      DES 2.200340e-33 2.273832e-29 -5.823757
## 257      DHCR24 7.474934e-04 1.647036e-02 -1.712001
## 258      DHCR7-DT 4.277941e-04 1.051326e-02  3.296091
## 259      DHRS2 1.384746e-10 2.805876e-08  5.897005
## 260      DIP2C-AS1 6.502896e-05 2.240031e-03 -3.469270
## 261      DKK1 7.400767e-04 1.634178e-02  3.281840
## 262      DLGAP1-AS1 7.635522e-04 1.680628e-02  2.398370
## 263      DLGAP5 1.639670e-04 4.820582e-03  2.312016
## 264      DLL3 1.707995e-03 3.274660e-02  4.747036
## 265      DLX1 4.009265e-05 1.488488e-03  5.258218
## 266      DLX1_1 4.009265e-05 1.488488e-03  5.258218
## 267      DLX2 1.231073e-07 1.125832e-05  6.231012
## 268      DLX6 1.390444e-12 4.789617e-10  7.220500
## 269      DLX6-AS1 5.351712e-07 3.867454e-05  6.098045
## 270      DMP1 9.785835e-04 2.077410e-02  4.826783
## 271      DMRT1 1.970297e-03 3.688595e-02  5.252268
## 272      DNAH14 1.742569e-06 1.077101e-04  3.360521
## 273      DNAH17 6.193482e-04 1.406577e-02  3.865795

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## 274 DNAH5 3.576129e-04 9.080029e-03 3.512767
## 275 DNMT3B 3.206351e-11 7.984200e-09 2.935475
## 276 DOCK9 1.366046e-07 1.227541e-05 -2.201899
## 277 DOP1B 6.721441e-04 1.508347e-02 -2.007140
## 278 DPF1 5.549497e-07 3.918363e-05 3.464317
## 279 DPYSL5 2.323398e-03 4.219683e-02 6.130886
## 280 DRAXIN 2.148479e-03 3.964712e-02 2.887502
## 281 DSG3 1.566619e-03 3.057495e-02 -2.296234
## 282 DTL 1.183150e-03 2.438021e-02 2.162144
## 283 DUOX1 6.406509e-04 1.447101e-02 -2.576857
## 284 DUSP9 1.327409e-05 5.977099e-04 4.574561
## 285 DUXAP10 2.740257e-06 1.577846e-04 4.947445
## 286 DUXAP8 5.937322e-04 1.361960e-02 4.372594
## 287 DUXAP9 1.172300e-07 1.081656e-05 4.197566
## 288 DYNAP 2.813829e-03 4.911842e-02 -3.469109
## 289 E2F1 4.641661e-04 1.120723e-02 2.319064
## 290 E2F7 1.568869e-03 3.058998e-02 2.096709
## 291 ECM1 1.926419e-07 1.645257e-05 -3.988249
## 292 ECRG4 5.931261e-07 4.141463e-05 -4.068215
## 293 ECT2 3.769425e-05 1.416481e-03 1.963196
## 294 EGF 1.382770e-04 4.205689e-03 3.352511
## 295 EHD3 4.886588e-08 5.049800e-06 -3.195210
## 296 EHF 1.110833e-04 3.504450e-03 -2.245851
## 297 EIF5A2 1.960032e-03 3.676037e-02 1.952134
## 298 ELOVL4 1.994715e-07 1.685641e-05 -2.963806
## 299 EMID1 7.411824e-05 2.486196e-03 3.531609
## 300 EMP1 6.115482e-13 2.298087e-10 -3.946682
## 301 EMSY-DT 3.507450e-04 8.927584e-03 2.952592
## 302 EN1 1.474718e-09 2.326678e-07 6.308133
## 303 EN2 2.437165e-03 4.376311e-02 3.507301
## 304 ENDOU 3.835217e-12 1.148786e-09 -5.382425
## 305 ENTREP1 3.598046e-06 2.004431e-04 -2.811934
## 306 EPB41L4A 1.575207e-04 4.687819e-03 -1.956706
## 307 EPCAM 9.787774e-05 3.132637e-03 2.997321
## 308 EPGN 1.114822e-04 3.504450e-03 -3.633575
## 309 EPHA1 7.392239e-06 3.688944e-04 -2.597286
## 310 EPHA1_1 7.392239e-06 3.688944e-04 -2.597286
## 311 EPHB2 9.797420e-08 9.288673e-06 3.774833
## 312 EPHX2 4.243490e-05 1.557805e-03 -2.691414
## 313 EPHX3 1.215277e-03 2.491801e-02 -2.764085
## 314 EPS8L1 1.104730e-04 3.491218e-03 -3.436726
## 315 EPS8L2 6.798694e-05 2.330272e-03 -2.493373
## 316 EREG 1.166256e-03 2.412830e-02 -3.501221
## 317 ERFE 1.199550e-13 5.509399e-11 5.773914
## 318 ESM1 6.668553e-11 1.450796e-08 4.594029
## 319 ESPL1_1 2.751591e-05 1.077081e-03 -2.679451
## 320 ETFDH 5.411543e-05 1.908944e-03 -2.196594
## 321 EVA1A 1.778663e-05 7.487046e-04 3.765410
## 322 EVPL 2.888457e-05 1.115862e-03 -2.835649
## 323 F10 1.072602e-03 2.232480e-02 -2.558238
## 324 FABP4 6.029672e-05 2.090961e-03 6.104357
## 325 FADD 5.633620e-04 1.309738e-02 2.930394
## 326 FADS1 3.266785e-08 3.553574e-06 3.344929
## 327 FADS2 3.254258e-07 2.519064e-05 3.843030

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## 331	FAM135A	8.982869e-04	1.948142e-02	-1.968922
## 332	FAM25A	1.754910e-03	3.352169e-02	-3.286055
## 333	FAM3B	3.550996e-07	2.708191e-05	-3.749207
## 334	FAM3D	8.920157e-06	4.327742e-04	-3.042609
## 335	FAR2P1	2.785398e-04	7.390066e-03	7.708595
## 336	FBN2	6.324016e-09	8.432566e-07	4.851673
## 337	FCER1A	2.135612e-03	3.944488e-02	-3.319893
## 338	FCER1A_1	2.135612e-03	3.944488e-02	-3.319893
## 339	FCH02	4.011458e-05	1.488488e-03	-2.262602
## 340	FEZF1	8.334211e-05	2.751621e-03	6.130205
## 341	FEZF1-AS1	6.082014e-07	4.218224e-05	6.243414
## 342	FGF12	7.333436e-06	3.688944e-04	3.383388
## 343	FGF19	1.402323e-13	6.300698e-11	8.974206
## 344	FGF5	1.151590e-03	2.387269e-02	4.317060
## 345	FHAD1	9.705554e-04	2.067983e-02	2.983394
## 346	FHL1	5.736533e-07	4.019074e-05	-3.036548
## 347	FIBCD1	1.158394e-06	7.435309e-05	5.385084
## 348	FIRRE	4.827620e-06	2.591617e-04	5.189665
## 349	FLG	1.044296e-06	6.765987e-05	-5.023499
## 350	FLNC	3.944896e-04	9.870837e-03	-2.946375
## 351	FLOT1_1	9.342875e-04	2.007261e-02	2.897490
## 352	FNDC4	1.916855e-06	1.161805e-04	-3.153353
## 353	FOXD2-AS1	8.972985e-06	4.342276e-04	3.244359
## 354	FOXD3-AS1	1.741751e-03	3.330112e-02	4.171909
## 355	FOXI2	2.422558e-07	1.963507e-05	-5.026217
## 356	FOXI3	1.001781e-06	6.531486e-05	6.510808
## 357	FOXL1	6.457815e-06	3.328432e-04	4.268553
## 358	FOXL2	1.177326e-05	5.421798e-04	3.883344
## 359	FOXM1	2.036442e-04	5.726420e-03	2.070474
## 360	FOXO6_1	2.720452e-03	4.809778e-02	2.883230
## 361	FREM2	5.320575e-06	2.812420e-04	6.748405
## 362	FRMD4B	9.178679e-06	4.391318e-04	-2.075524
## 363	FUT3_1	6.662322e-07	4.574647e-05	-2.948489
## 364	FUT6	6.100989e-15	3.708683e-12	-5.359433
## 365	FXYD1	1.960107e-07	1.667140e-05	-3.317661
## 366	GAB1	6.214155e-04	1.408269e-02	-1.781714
## 367	GABRA4	7.356076e-05	2.476146e-03	-4.764576
## 368	GABRP	7.030203e-05	2.401657e-03	-3.348376
## 369	GAL	1.915175e-03	3.618174e-02	3.822356
## 370	GALNT17	2.575261e-06	1.490910e-04	-3.219810
## 371	GAP43	4.814730e-04	1.153080e-02	4.145143
## 372	GAST	1.383718e-04	4.205689e-03	5.874735
## 373	GATA4	1.682512e-05	7.150995e-04	6.294877
## 374	GBP6	1.242332e-07	1.131124e-05	-3.403819
## 375	GBX2	3.225447e-08	3.527171e-06	6.659396
## 376	GCKR	1.436394e-03	2.868348e-02	-3.574756
## 377	GCNT4	8.034691e-07	5.426830e-05	-2.359450
## 378	GCOM1	5.881366e-06	3.069598e-04	-4.147293
## 379	GDA	9.138959e-06	4.382460e-04	4.235641
## 380	GDF15	5.564657e-05	1.942742e-03	4.510794
## 381	GDF7	1.129179e-05	5.244467e-04	-2.969267

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## 382      GDPD2 5.819830e-04 1.343958e-02  3.872908
## 383      GDPD3 6.913378e-08 6.869505e-06 -3.643773
## 384      GFOD2 1.593199e-03 3.094759e-02 -2.041622
## 385      GFRA1 2.979166e-04 7.774420e-03 -3.071485
## 386      GGT6  2.544546e-06 1.477266e-04 -2.403564
## 387      GINS1 1.327456e-03 2.684526e-02  1.869682
## 388      GINS2 5.628778e-08 5.759187e-06  2.482497
## 389      GJB7  2.436688e-11 6.217464e-09  6.599639
## 390      GLI1  3.524557e-05 1.329298e-03  4.407792
## 391      GLI2  1.753106e-04 5.079354e-03  3.164524
## 392      GLTP  2.859525e-06 1.632615e-04 -2.422314
## 393      GNG7  9.354642e-04 2.007425e-02 -2.552658
## 394      GPC2  9.342494e-04 2.007261e-02  3.008774
## 395      GPD1L 2.698436e-07 2.150220e-05 -2.264351
## 396      GPIHBP1 9.854119e-06 4.660525e-04 -3.145514
## 397      GPR149 1.511222e-03 2.983185e-02  6.332652
## 398      GPR158 8.925844e-05 2.923603e-03  6.011792
## 399      GPR19  8.425959e-04 1.842319e-02  2.810312
## 400      GPR50 8.588241e-04 1.870409e-02  4.517511
## 401      GPT   4.861326e-06 2.602950e-04 -3.292211
## 402      GPT2  7.713634e-04 1.696015e-02 -2.076855
## 403      GPX3  2.222164e-04 6.156526e-03 -2.664937
## 404      GREM2 1.916465e-05 7.953716e-04 -3.623787
## 405      GRHL1 6.196216e-04 1.406577e-02 -2.308948
## 406      GRHL3 2.740694e-06 1.577846e-04 -2.953343
## 407      GRIN2D 8.973056e-05 2.934416e-03  3.737972
## 408      GRM4  1.228695e-04 3.807297e-03  5.264138
## 409      GRP   1.902890e-04 5.405488e-03  5.865513
## 410      GSC   5.397963e-04 1.264910e-02  5.295591
## 411      GTF2H2 1.424354e-03 2.852573e-02 -6.106284
## 412      GTF3C2-AS2 4.317188e-06 2.341933e-04 3.212989
## 413      GUCY1B2 2.812704e-03 4.911842e-02  4.496990
## 414      GYS2  2.948686e-09 4.211868e-07 -5.079477
## 415      H2AC19 7.614423e-12 2.147125e-09 21.362969
## 416      H2BC9  2.194906e-03 4.028803e-02  2.483033
## 417      H3C8  2.528918e-03 4.501953e-02  4.307085
## 418      HAGHL 4.481042e-04 1.093438e-02  2.241769
## 419      HAGLR 9.701335e-04 2.067983e-02  2.430394
## 420      HAGLROS 1.520283e-05 6.601094e-04  4.681027
## 421      HAP1  4.298065e-13 1.700489e-10  4.540555
## 422      HCG22_1 2.014483e-07 1.685641e-05 -7.966206
## 423      HCG22_2 2.014483e-07 1.685641e-05 -7.966206
## 424      HCG22_3 2.014483e-07 1.685641e-05 -7.966206
## 425      HCG22_6 2.122237e-07 1.754496e-05 -6.765059
## 426      HCG22_7 3.415320e-04 8.757795e-03 -10.678966
## 427      HES6  4.097620e-06 2.240466e-04  3.103272
## 428      HEY1  1.292486e-08 1.580657e-06  4.315672
## 429      HHIP  4.698340e-05 1.694682e-03  6.137370
## 430      HHIPL2 1.187642e-03 2.444839e-02  3.986500
## 431      HJURP 2.884248e-06 1.642194e-04  2.159983
## 432      HLA-DQA_5 2.359253e-03 4.269794e-02 -4.687193
## 433      HLA-DQB2_3 1.878587e-03 3.555553e-02 -3.314880
## 434      HLA-DQB2_5 1.878587e-03 3.555553e-02 -3.314880
## 435      HMCN2 1.286046e-04 3.961253e-03 -2.618386

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## 436	HMGA2	1.763324e-09	2.699583e-07	4.956659
## 437	HMGB3	4.012457e-06	2.199721e-04	2.301583
## 438	HMGCS2	3.650258e-13	1.479285e-10	-6.050288
## 439	HMMR	1.048631e-05	4.925704e-04	2.862831
## 440	HMX1	1.463395e-03	2.908217e-02	5.532836
## 441	HMX1_1	1.463395e-03	2.908217e-02	5.532836
## 442	HOMER3	8.648294e-05	2.841700e-03	1.989510
## 443	HOTAIR	3.980416e-04	9.938781e-03	5.537591
## 444	HOXA10	6.706960e-16	5.134053e-13	5.884920
## 445	HOXA10-AS	2.192250e-07	1.790886e-05	5.746006
## 446	HOXA11	1.152148e-08	1.442054e-06	6.793580
## 447	HOXA13	9.130063e-10	1.546723e-07	7.113792
## 448	HOXA9	4.982811e-04	1.180950e-02	3.707979
## 449	HOXB6	2.280451e-05	9.187594e-04	2.900030
## 450	HOXB7	1.646840e-14	9.454691e-12	5.055295
## 451	HOXB8	4.515242e-04	1.100304e-02	5.027547
## 452	HOXB9	4.566279e-07	3.373720e-05	7.661604
## 453	HOXC10	2.694633e-10	5.062971e-08	6.298655
## 454	HOXC11	1.450383e-05	6.337532e-04	6.339022
## 455	HOXC13	3.900678e-11	9.266576e-09	6.745942
## 456	HOXC13-AS	1.028874e-06	6.687033e-05	6.632678
## 457	HOXC9	1.216789e-05	5.563847e-04	3.680899
## 458	HOXD10	7.684336e-18	9.342344e-15	4.878219
## 459	HOXD11	2.204455e-38	4.556167e-34	6.974538
## 460	HOXD13	1.658926e-17	1.804562e-14	9.440985
## 461	HOXD8	4.633552e-04	1.120073e-02	1.772342
## 462	HPGD	4.751970e-04	1.143141e-02	-3.190964
## 463	HPSE2	1.067038e-26	5.513385e-23	-5.448579
## 464	HS3ST5	2.176537e-03	4.009328e-02	6.083482
## 465	HSPA6	6.034511e-04	1.378136e-02	3.475552
## 466	HSPB6	1.900253e-07	1.629644e-05	-3.771991
## 467	HSPB7	2.933348e-04	7.679951e-03	-2.813454
## 468	HTR2C	2.349033e-06	1.387138e-04	6.256706
## 469	HTR3B	3.182381e-06	1.787322e-04	-3.664048
## 470	IBSP	3.041223e-09	4.305206e-07	6.860903
## 471	IFI6	3.046941e-05	1.166188e-03	3.726198
## 472	IFIT1	1.384376e-03	2.780591e-02	2.932107
## 473	IGF2BP1	2.882556e-07	2.273918e-05	9.126923
## 474	IGF2BP2	4.012507e-08	4.252846e-06	3.861938
## 475	IGFBP3	1.114109e-09	1.798938e-07	4.050837
## 476	IGFL1	1.504463e-04	4.511571e-03	-3.450313
## 477	IL11	5.421667e-05	1.908944e-03	4.047767
## 478	IL12A	1.049385e-03	2.192991e-02	-3.182216
## 479	IL12A-AS1	2.916248e-06	1.655852e-04	-5.929746
## 480	IL13RA2	1.239171e-03	2.528251e-02	3.625986
## 481	IL18	5.898494e-04	1.357573e-02	-2.352964
## 482	IL1RN	4.763574e-05	1.713459e-03	-3.202644
## 483	IL24	6.191967e-04	1.406577e-02	3.712402
## 484	IL31RA	2.217981e-05	8.965105e-04	5.199156
## 485	IL34	2.575633e-07	2.079421e-05	-3.110092
## 486	IL36A	1.598893e-06	9.983662e-05	-4.527463
## 487	IL36B	3.668278e-04	9.279800e-03	-3.231847
## 488	IL36G	1.426389e-03	2.853882e-02	3.211320
## 489	INHBA	1.990388e-10	3.917843e-08	5.126958

## 490	IP6K3	1.663568e-09	2.604744e-07	-4.562975
## 491	IRS4	2.674034e-04	7.184753e-03	5.806283
## 492	ISG15	3.762948e-07	2.848814e-05	3.888150
## 493	IVL	2.000339e-04	5.647953e-03	-3.487145
## 494	JAKMIP3	1.462826e-07	1.303176e-05	3.659513
## 495	JPH2	1.374743e-10	2.805876e-08	-3.604444
## 496	KALRN	4.622513e-05	1.670247e-03	-2.681127
## 497	KANK1	3.723869e-09	5.200333e-07	-2.869435
## 498	KAT2B	3.957550e-07	2.961585e-05	-2.729732
## 499	KAZN	2.061132e-03	3.830888e-02	-1.864385
## 500	KCNA5	1.954340e-06	1.181062e-04	-4.599263
## 501	KCNB1	4.029968e-14	2.082284e-11	-4.926136
## 502	KCNC3	2.369560e-04	6.503861e-03	2.279744
## 503	KCNG3	1.283611e-05	5.817911e-04	4.666543
## 504	KCNK3	2.536901e-09	3.799469e-07	-3.113701
## 505	KCNMA1	2.782414e-05	1.085036e-03	-2.569439
## 506	KCNMB1	4.319061e-05	1.577144e-03	-2.902699
## 507	KCNMB2-AS1	4.177586e-11	9.811630e-09	6.696176
## 508	KCNMB3	1.473185e-05	6.410059e-04	3.088068
## 509	KCNS3	1.266605e-05	5.753449e-04	2.396974
## 510	KIAA1549	8.698604e-04	1.890460e-02	2.798982
## 511	KIF14	1.399515e-05	6.180592e-04	2.708076
## 512	KIF1A	1.005837e-09	1.663091e-07	5.641306
## 513	KIF26B	6.913291e-05	2.365628e-03	3.476663
## 514	KIF4A	9.545556e-04	2.042314e-02	1.869158
## 515	KILH	2.598771e-04	7.002789e-03	5.317647
## 516	KLHDC7B	1.981999e-04	5.611500e-03	3.785920
## 517	KLHDC7B-DT	2.777636e-05	1.085036e-03	4.068822
## 518	KLHDC8A	2.769617e-04	7.367109e-03	-2.763544
## 519	KLK11	2.025070e-06	1.220238e-04	-2.585626
## 520	KLK12	7.710714e-05	2.574556e-03	-3.361951
## 521	KLK13	1.255040e-05	5.713473e-04	-3.670627
## 522	KPRP	9.081003e-04	1.963244e-02	-4.343011
## 523	KREMEN2	6.302511e-04	1.426728e-02	2.312493
## 524	KRT13	2.987137e-05	1.147549e-03	-4.108900
## 525	KRT24	3.596713e-04	9.109911e-03	-4.743405
## 526	KRT3	6.554621e-09	8.684032e-07	-4.407911
## 527	KRT32	4.063507e-08	4.284927e-06	-3.992439
## 528	KRT4	3.634926e-11	8.735658e-09	-6.426344
## 529	KRT42P	2.690451e-04	7.202881e-03	4.645720
## 530	KRT75	2.954910e-09	4.211868e-07	5.826676
## 531	KRT78	6.306293e-11	1.395881e-08	-5.568714
## 532	KRT8	3.044149e-06	1.714345e-04	3.195239
## 533	KRTAP4-1_1	8.992134e-06	4.342276e-04	5.894576
## 534	LAMA1	5.812414e-06	3.041290e-04	3.906487
## 535	LAMB4	7.641550e-13	2.820278e-10	-5.023367
## 536	LAMC2	1.596060e-05	6.858083e-04	2.989928
## 537	LAMP3	1.370000e-03	2.759763e-02	2.366640
## 538	LAYN	1.774318e-03	3.386114e-02	2.962694
## 539	LCAL1	4.076958e-17	4.213129e-14	10.149662
## 540	LCN1	4.537270e-04	1.101954e-02	-5.664728
## 541	LCN10	7.436361e-04	1.640285e-02	-2.486435
## 542	LDB3	4.301304e-11	9.988692e-09	-3.588058
## 543	LGI1	6.931562e-06	3.537322e-04	-3.775251

## 544	LGI3	2.192393e-04	6.090372e-03	-2.862533
## 545	LHX2	1.680459e-10	3.339589e-08	7.663619
## 546	LHX5	1.141626e-07	1.058078e-05	6.444394
## 547	LINC00467	7.618781e-05	2.547977e-03	3.085382
## 548	LINC00491	1.883198e-05	7.831376e-04	4.095510
## 549	LINC00519	1.506185e-04	4.511571e-03	3.945064
## 550	LINC00622	2.739131e-03	4.830407e-02	3.460784
## 551	LINC00649	9.745589e-09	1.243345e-06	2.392794
## 552	LINC00942	4.089453e-12	1.190434e-09	8.723056
## 553	LINC01088	7.194003e-04	1.598771e-02	-3.744425
## 554	LINC01234	1.134370e-07	1.056088e-05	7.488093
## 555	LINC01269	1.118680e-04	3.506662e-03	-3.043004
## 556	LINC01297-DUXAP10-NBEAP6	1.823286e-04	5.233843e-03	2.419749
## 557	LINC01305	7.092271e-04	1.580771e-02	5.128829
## 558	LINC01614	3.392408e-06	1.894981e-04	5.702221
## 559	LINC01615	2.140653e-05	8.742275e-04	4.851814
## 560	LINC01873	1.452399e-03	2.891925e-02	2.275090
## 561	LINC02041_1	2.845073e-03	4.945497e-02	5.340062
## 562	LINC02043	2.385780e-12	7.704577e-10	4.801320
## 563	LINC02086	1.379844e-03	2.776886e-02	4.849915
## 564	LINC02137	1.921574e-03	3.626950e-02	4.677634
## 565	LINC02154	4.991375e-04	1.180950e-02	4.672197
## 566	LINC02159	1.604897e-03	3.105810e-02	4.926052
## 567	LINC02253	7.105362e-04	1.580771e-02	2.732653
## 568	LINC02321	2.257522e-03	4.118134e-02	4.116024
## 569	LINC02457	2.372334e-03	4.285718e-02	4.842186
## 570	LINC02487	3.779649e-04	9.514955e-03	-4.049008
## 571	LINC02538	1.669665e-03	3.212862e-02	-3.823813
## 572	LINC02560	3.967656e-05	1.482885e-03	-2.926668
## 573	LINC02561	4.465861e-04	1.091022e-02	4.875400
## 574	LINC02577	1.560988e-03	3.052299e-02	3.735854
## 575	LINC02609	1.942509e-03	3.653118e-02	3.145915
## 576	LINC02623	1.608860e-05	6.886073e-04	-6.379374
## 577	LINC02636	1.959903e-04	5.556555e-03	2.774930
## 578	LINC02827	8.345892e-07	5.618661e-05	5.902155
## 579	LINC02884	5.961354e-05	2.074230e-03	-3.404353
## 580	LINC02893	2.822798e-06	1.620600e-04	3.617737
## 581	LINC02994	8.261631e-05	2.736401e-03	5.265165
## 582	LINC02999	1.115700e-04	3.504450e-03	5.688034
## 583	LINC03011	1.340531e-03	2.708317e-02	2.982132
## 584	LINC03016	3.818535e-07	2.880346e-05	-5.737787
## 585	LINC03040	1.107416e-05	5.166607e-04	5.496827
## 586	LINC03050	1.904004e-04	5.405488e-03	3.307317
## 587	LINC03057	6.852237e-06	3.505496e-04	6.089477
## 588	LIPH	4.102884e-04	1.015854e-02	-2.458580
## 589	LM07	5.927733e-04	1.361271e-02	-2.460277
## 590	LMOD1	7.976556e-16	5.684809e-13	-4.246920
## 591	LMX1B	2.669118e-05	1.050768e-03	4.402419
## 592	LOC100287072	2.679207e-05	1.052735e-03	2.803376
## 593	LOC101927293	4.631992e-04	1.120073e-02	5.282676
## 594	LOC101927469	3.465828e-05	1.309538e-03	4.476802
## 595	LOC101928051	2.045519e-03	3.808719e-02	4.106970
## 596	LOC101928391	3.327469e-05	1.261874e-03	2.788825
## 597	LOC101928844	6.773019e-09	8.859794e-07	-5.006118

## 598	LOC101928844_1	6.773019e-09	8.859794e-07	-5.006118
## 599	LOC101929007	8.288299e-06	4.040155e-04	-4.586844
## 600	LOC101929594	1.104988e-05	5.166607e-04	2.914968
## 601	LOC101929748	2.832753e-06	1.621810e-04	6.275617
## 602	LOC102723546	2.615483e-12	8.316431e-10	7.828417
## 603	LOC102723686	2.339888e-06	1.385696e-04	-6.046804
## 604	LOC102723825	1.105177e-05	5.166607e-04	5.098425
## 605	LOC102723985	7.612839e-06	3.764166e-04	2.629702
## 606	LOC102724542	8.030317e-08	7.828801e-06	6.489403
## 607	LOC102724858	1.745144e-05	7.391113e-04	4.356653
## 608	LOC102725051	2.567293e-04	6.926999e-03	-4.378691
## 609	LOC102725238	8.850861e-04	1.921529e-02	2.434259
## 610	LOC105369147	2.070816e-04	5.807275e-03	2.534305
## 611	LOC105369266	6.994411e-07	4.786771e-05	-6.601385
## 612	LOC105369367	9.433439e-04	2.020418e-02	6.088664
## 613	LOC105369526	1.735969e-12	5.786938e-10	-6.948291
## 614	LOC105369601	8.038422e-06	3.965110e-04	3.122597
## 615	LOC105369887	3.053686e-04	7.938814e-03	5.090807
## 616	LOC105369975	2.513669e-05	1.001012e-03	-3.271354
## 617	LOC105370256	4.966846e-05	1.769910e-03	3.787447
## 618	LOC105370413	9.481329e-07	6.260706e-05	4.631686
## 619	LOC105370475	2.120212e-16	1.752822e-13	5.961380
## 620	LOC105370964	5.564694e-06	2.919063e-04	-3.761549
## 621	LOC105371175	1.722744e-09	2.657141e-07	-7.195873
## 622	LOC105371206	8.451085e-14	4.062024e-11	-6.451143
## 623	LOC105371207	3.481625e-04	8.872776e-03	-3.313280
## 624	LOC105371216	1.575425e-08	1.882132e-06	-6.089055
## 625	LOC105371217	3.481625e-04	8.872776e-03	-3.313280
## 626	LOC105371401	2.463162e-04	6.716179e-03	3.783830
## 627	LOC105371956	1.494170e-03	2.956789e-02	3.951683
## 628	LOC105372130	2.782324e-09	4.049653e-07	5.935220
## 629	LOC105373421	6.435163e-04	1.451986e-02	-3.466912
## 630	LOC105373582	7.420368e-06	3.688944e-04	3.976205
## 631	LOC105373949	3.886770e-04	9.749000e-03	5.572406
## 632	LOC105374122	2.121414e-06	1.270881e-04	6.692791
## 633	LOC105374264	1.458349e-06	9.197650e-05	2.902248
## 634	LOC105374816	2.051373e-04	5.760568e-03	5.896985
## 635	LOC105374848	1.210481e-03	2.484432e-02	-3.689391
## 636	LOC105375065	4.192623e-05	1.541871e-03	2.938649
## 637	LOC105375520	6.586259e-04	1.482841e-02	3.644871
## 638	LOC105375690	9.900106e-05	3.162525e-03	3.714075
## 639	LOC105375772	1.955389e-03	3.670661e-02	4.448575
## 640	LOC105375785	2.751768e-03	4.844423e-02	2.636012
## 641	LOC105376270	2.466468e-03	4.413589e-02	2.976998
## 642	LOC105376272	1.161395e-09	1.846439e-07	3.733960
## 643	LOC105377067	3.022947e-05	1.159151e-03	5.524255
## 644	LOC105377261	2.450276e-04	6.689870e-03	5.078891
## 645	LOC105377581	6.432302e-06	3.323570e-04	5.131912
## 646	LOC105377635	3.210609e-12	9.758363e-10	-6.152578
## 647	LOC105377911	1.807599e-04	5.196029e-03	5.230833
## 648	LOC105378030	2.141879e-05	8.742275e-04	5.681747
## 649	LOC105378675	1.052819e-03	2.197946e-02	5.232366
## 650	LOC105379524	5.320769e-05	1.886272e-03	5.985833
## 651	LOC105379539	2.172688e-03	4.005810e-02	-3.538696

## 652	LOC105379854	2.010462e-03	3.750202e-02	-3.116958
## 653	LOC107984006	1.631250e-03	3.145026e-02	-2.674685
## 654	LOC107984128	1.171757e-03	2.421788e-02	3.646776
## 655	LOC107984865_1	2.802332e-03	4.909029e-02	5.767151
## 656	LOC107985705	2.237191e-10	4.281320e-08	6.083644
## 657	LOC107985915	1.776312e-05	7.487046e-04	-3.512855
## 658	LOC107985962	2.172559e-04	6.051543e-03	2.637088
## 659	LOC107986056	1.002006e-03	2.115369e-02	5.275134
## 660	LOC107986141	1.412440e-03	2.834204e-02	-3.384648
## 661	LOC107986764	9.468542e-10	1.578192e-07	-5.830406
## 662	LOC107987138	2.853668e-03	4.956270e-02	5.377954
## 663	LOC107987365	2.158991e-10	4.170282e-08	-7.153943
## 664	LOC107987401	4.405458e-05	1.601724e-03	-3.358731
## 665	LOC107987423	7.998650e-20	1.653161e-16	25.190987
## 666	LOC110091776	2.139237e-04	5.982917e-03	-4.121904
## 667	LOC112267868	1.730610e-07	1.522053e-05	3.971436
## 668	LOC112268114	2.787450e-03	4.890579e-02	-2.664563
## 669	LOC112268153	7.425008e-06	3.688944e-04	-3.779089
## 670	LOC122513141	1.393968e-05	6.180592e-04	-2.371848
## 671	LOC124900467	2.203934e-04	6.114216e-03	4.173674
## 672	LOC124900509	2.935534e-04	7.679951e-03	-4.391009
## 673	LOC124900586	4.457980e-04	1.090385e-02	4.167246
## 674	LOC124900799	2.641627e-05	1.041930e-03	2.761224
## 675	LOC124901002	1.344690e-05	6.028643e-04	-5.641890
## 676	LOC124901321	2.333135e-04	6.412398e-03	4.174420
## 677	LOC124901380	8.329697e-05	2.751621e-03	5.363644
## 678	LOC124901465	4.756634e-04	1.143141e-02	-4.195062
## 679	LOC124901609	1.302581e-05	5.890974e-04	2.413012
## 680	LOC124901681	1.015463e-04	3.228861e-03	2.834843
## 681	LOC124902211	4.780609e-04	1.146237e-02	5.134138
## 682	LOC124902365	8.835280e-08	8.493375e-06	5.948565
## 683	LOC124902366	2.333028e-03	4.229740e-02	4.575091
## 684	LOC124902696	3.480932e-04	8.872776e-03	4.853257
## 685	LOC124902719	1.470133e-05	6.410059e-04	4.914521
## 686	LOC124902874	3.249927e-14	1.722295e-11	5.341204
## 687	LOC124902884	2.617012e-03	4.642782e-02	4.076092
## 688	LOC124902961	6.797012e-08	6.786505e-06	3.247779
## 689	LOC124903025	1.550352e-03	3.038915e-02	5.101163
## 690	LOC124903391	1.987504e-03	3.714080e-02	3.414647
## 691	LOC124903505	1.338915e-04	4.099659e-03	-3.115815
## 692	LOC124903541	1.776107e-03	3.386400e-02	3.764075
## 693	LOC124903758	8.295246e-04	1.816167e-02	-3.491529
## 694	LOC124903859	2.846530e-04	7.523283e-03	3.511914
## 695	LOC124904198	4.872765e-04	1.163404e-02	5.455711
## 696	LOC124904253	1.609238e-05	6.886073e-04	6.332531
## 697	LOC124904420	2.881610e-05	1.115302e-03	-4.961085
## 698	LOC124904423	4.765572e-06	2.564970e-04	3.349230
## 699	LOC124904447	5.573827e-07	3.918363e-05	5.409353
## 700	LOC124904544	3.287405e-11	8.088581e-09	4.759073
## 701	LOC124904731	1.996362e-03	3.727263e-02	5.125800
## 702	LOC124904915	6.556176e-08	6.642306e-06	-6.438119
## 703	LOC124905027	2.186641e-03	4.017200e-02	4.398057
## 704	LOC124905038	8.450426e-08	8.161374e-06	6.592308
## 705	LOC124905421	9.529299e-05	3.067781e-03	4.155461

## 706	LOC124905488	5.084848e-04	1.198331e-02	4.076661
## 707	LOC124906047	2.818391e-05	1.094934e-03	-4.133062
## 708	LOC124906245	1.128560e-06	7.266381e-05	3.189047
## 709	LOC124906252	7.373334e-06	3.688944e-04	-5.490473
## 710	LOC124906274	3.290964e-08	3.561133e-06	6.422909
## 711	LOC124906315	1.820535e-07	1.580959e-05	8.092887
## 712	LOC124907726	1.757922e-05	7.430009e-04	3.322772
## 713	LOC124907737	1.232837e-03	2.520305e-02	3.671566
## 714	LOC124907854	4.360649e-13	1.700489e-10	-6.208075
## 715	LOC124907910	2.222774e-08	2.524192e-06	-4.972562
## 716	LOC124907946	1.686445e-08	1.980423e-06	6.410888
## 717	LOC408186	5.629873e-04	1.309738e-02	2.922080
## 718	LOC440910	1.635978e-03	3.151202e-02	5.746305
## 719	LOC730101	1.938495e-03	3.650710e-02	3.148668
## 720	LOXL2	7.966121e-07	5.398157e-05	3.062593
## 721	LPIN1	8.711825e-06	4.236612e-04	-2.288133
## 722	LRP10	9.526271e-06	4.515802e-04	-1.939655
## 723	LRP12	3.320735e-04	8.557725e-03	2.409885
## 724	LRRC15	9.684445e-07	6.374462e-05	5.414281
## 725	LYPD1	1.626356e-03	3.138518e-02	3.990182
## 726	LYPD2	5.548029e-07	3.918363e-05	-4.132497
## 727	LYPD3	8.440537e-04	1.842319e-02	-2.620930
## 728	LYVE1	1.015611e-03	2.133196e-02	-2.797236
## 729	MAB21L2	1.788005e-06	1.093328e-04	-5.000498
## 730	MAB21L4	2.730976e-03	4.820137e-02	-3.259686
## 731	MACC1	1.596699e-03	3.098645e-02	-2.164663
## 732	MAFA	4.440151e-04	1.087311e-02	4.100720
## 733	MAGEA1	3.890719e-05	1.456764e-03	9.624323
## 734	MAGEA11	3.588390e-04	9.099980e-03	5.460437
## 735	MAGEA3	7.284244e-06	3.680947e-04	9.854543
## 736	MAGEA3_1	1.039281e-05	4.892908e-04	9.651252
## 737	MAGEA4	2.485773e-05	9.918137e-04	11.350465
## 738	MAGEA6_1	2.739448e-05	1.074363e-03	11.168207
## 739	MAL	8.215525e-06	4.023660e-04	-4.917325
## 740	MALL	2.565954e-05	1.017910e-03	-3.043013
## 741	MALLP2	6.019467e-04	1.377095e-02	-3.129838
## 742	MAPK12	4.809211e-10	8.283065e-08	3.302408
## 743	MAPT	4.396212e-10	7.878316e-08	-4.144386
## 744	MAPT_1	5.262637e-06	2.793128e-04	-4.255444
## 745	MAPT_2	1.461375e-07	1.303176e-05	-3.516632
## 746	MARCKSL1	1.275182e-07	1.155941e-05	2.569228
## 747	MC1R	6.127184e-04	1.397755e-02	2.324406
## 748	MCCC2_1	1.176000e-03	2.428129e-02	2.068511
## 749	MCIDAS	2.621500e-04	7.054838e-03	4.685459
## 750	MCM2	3.173350e-04	8.208611e-03	2.213711
## 751	MCOLN3	4.002930e-05	1.488488e-03	2.586861
## 752	MDK	5.418603e-07	3.902150e-05	2.902095
## 753	MECOM-AS1	2.285513e-07	1.859723e-05	-5.812803
## 754	MEGF10	2.302263e-03	4.184976e-02	3.789461
## 755	MELTF	1.327226e-04	4.069896e-03	2.761210
## 756	MELTF-AS1	3.535412e-07	2.706292e-05	3.025152
## 757	MEST	5.414268e-06	2.847381e-04	3.083017
## 758	METTL21A	1.482546e-03	2.937801e-02	2.175690
## 759	MFAP2	1.282661e-07	1.157644e-05	3.881065

## 760	MFAP2_1	3.337140e-07	2.564015e-05	3.744666
## 761	MGLL	2.031651e-04	5.720731e-03	-2.674648
## 762	MINDY1	8.245205e-06	4.028650e-04	-2.369815
## 763	MIR9-3HG	3.761687e-04	9.492862e-03	2.841015
## 764	MISP	1.181043e-15	7.628058e-13	4.820197
## 765	MLLT11	1.893854e-04	5.391485e-03	2.860576
## 766	MME	4.519750e-06	2.445398e-04	4.577817
## 767	MMP1	1.952686e-13	8.586834e-11	10.024847
## 768	MMP10	7.422020e-05	2.486196e-03	6.445193
## 769	MMP11	2.589043e-25	8.918389e-22	6.968003
## 770	MMP11_1	2.589043e-25	8.918389e-22	6.968003
## 771	MMP12	8.828905e-16	6.082527e-13	5.709569
## 772	MMP13	7.687597e-12	2.147125e-09	9.204225
## 773	MMP3	7.755367e-24	2.289827e-20	10.391534
## 774	MND1	2.165218e-08	2.486151e-06	3.052369
## 775	MPZL3	4.097288e-05	1.513426e-03	-2.403439
## 776	MS4A4E	5.788984e-04	1.339829e-02	2.561356
## 777	MSANTD3	2.450073e-05	9.794607e-04	2.156940
## 778	MSC	2.089611e-05	8.586098e-04	3.198001
## 779	MSI1	2.181062e-08	2.490507e-06	3.671964
## 780	MSX2	2.494772e-04	6.777814e-03	2.768051
## 781	MUC21	1.672653e-03	3.212862e-02	-4.933032
## 782	MUC21_1	2.695187e-12	8.440018e-10	-8.795640
## 783	MUC21_2	8.019181e-04	1.759453e-02	-5.661034
## 784	MUC21_3	2.371758e-12	7.704577e-10	-8.608956
## 785	MUC22	1.629013e-08	1.932439e-06	-9.896729
## 786	MUC22_1	1.081617e-08	1.371463e-06	-7.786738
## 787	MUC22_2	6.698438e-04	1.504819e-02	-7.605726
## 788	MUC22_7	8.174218e-06	4.019718e-04	-5.754626
## 789	MUC5B	3.347267e-05	1.267057e-03	-6.817035
## 790	MUCL1	1.720775e-04	4.995081e-03	5.139291
## 791	MUCL3	2.811209e-14	1.529002e-11	-8.114777
## 792	MUCL3_1	2.808169e-03	4.910256e-02	-6.814873
## 793	MUCL3_2	1.115197e-04	3.504450e-03	-5.878115
## 794	MUSTN1	1.370208e-04	4.183082e-03	-2.927784
## 795	MYBL2	4.303610e-08	4.515077e-06	2.325377
## 796	MYH11	4.993701e-07	3.634148e-05	-3.755102
## 797	MYH11_1	4.993701e-07	3.634148e-05	-3.755102
## 798	MYL9	2.966151e-05	1.141609e-03	-2.933443
## 799	MYOC	1.923254e-19	3.312485e-16	-9.146094
## 800	MYOCD	7.212244e-06	3.653497e-04	-3.837419
## 801	MYOZ1	1.012600e-04	3.224717e-03	-4.694806
## 802	MYRIP	4.682507e-04	1.129266e-02	-2.687636
## 803	MYZAP	2.844590e-07	2.252566e-05	-3.899927
## 804	N4BP3	1.594870e-04	4.722460e-03	-2.265274
## 805	NAT14	1.391425e-04	4.222904e-03	2.204311
## 806	NAT8L	2.228290e-03	4.071998e-02	2.926759
## 807	NCCRP1	9.035936e-04	1.957597e-02	-3.152976
## 808	NCK1-DT	4.154601e-05	1.530611e-03	3.020302
## 809	NDC80	7.263707e-04	1.612527e-02	2.354666
## 810	NDRG2	2.593479e-03	4.612911e-02	-1.874932
## 811	NEB	1.880655e-05	7.831376e-04	3.423178
## 812	NECAB2	1.494990e-03	2.956789e-02	4.087766
## 813	NEK10	2.229698e-04	6.169130e-03	-2.618473

## 814	NEK2	2.126430e-04	5.955157e-03	2.404138
## 815	NELL2	8.094926e-08	7.854739e-06	3.520369
## 816	NEURL3	8.603913e-04	1.871849e-02	3.316015
## 817	NEXN	1.866758e-04	5.329028e-03	-2.450713
## 818	NGB	1.430257e-04	4.302846e-03	5.573539
## 819	NIBAN2	6.040170e-10	1.031721e-07	-2.746913
## 820	NICOL1	1.971434e-05	8.149121e-04	2.851845
## 821	NIPAL1	6.348598e-11	1.395881e-08	-3.120640
## 822	NKAIN2	1.159201e-03	2.400639e-02	4.266265
## 823	NKX2-1	3.365263e-04	8.653582e-03	5.715618
## 824	NLRX1	1.342092e-06	8.534882e-05	-2.671439
## 825	NMB	1.842989e-07	1.587121e-05	4.095699
## 826	NMB_1	1.842989e-07	1.587121e-05	4.095699
## 827	NMRAL2P	5.560941e-07	3.918363e-05	4.624477
## 828	NOS2	1.408627e-04	4.256241e-03	4.376732
## 829	NOTUM	3.435812e-04	8.788536e-03	3.544654
## 830	NPAS2	9.128900e-05	2.968883e-03	-2.364852
## 831	NPBWR1	8.613938e-07	5.780288e-05	5.649350
## 832	NPY1R	4.092097e-04	1.015312e-02	-3.587613
## 833	NR0B1	4.314303e-05	1.577144e-03	6.295440
## 834	NRCAM	4.175129e-04	1.028505e-02	3.117400
## 835	NREP	1.537652e-03	3.026637e-02	2.361352
## 836	NRIP3	2.144539e-05	8.742275e-04	2.404298
## 837	NT5C2	5.066368e-04	1.195339e-02	-1.784537
## 838	NTRK3	1.684989e-05	7.150995e-04	-3.142386
## 839	NTS	5.161906e-05	1.833097e-03	6.147847
## 840	NUF2	4.586875e-07	3.373720e-05	2.885278
## 841	NXPH4	1.720730e-04	4.995081e-03	2.976568
## 842	OCLN_1	2.260104e-03	4.119209e-02	-2.585421
## 843	ODC1	3.981668e-04	9.938781e-03	3.023055
## 844	OGDHL	1.547121e-05	6.689520e-04	4.916908
## 845	OGN	2.495605e-04	6.777814e-03	-3.615193
## 846	OGN_1	2.495605e-04	6.777814e-03	-3.615193
## 847	ONECUT2	1.824649e-05	7.633976e-04	4.524312
## 848	OPRK1	2.585216e-05	1.021630e-03	7.181649
## 849	ORC6	1.391573e-05	6.180592e-04	2.499158
## 850	OTOP3	1.098274e-06	7.093479e-05	-3.629059
## 851	OTX1	2.071860e-03	3.842563e-02	2.573373
## 852	P2RX1	2.377193e-06	1.399767e-04	-3.418050
## 853	P2RX2	2.085307e-09	3.145921e-07	-6.521712
## 854	P2RY14	5.550078e-07	3.918363e-05	-3.416390
## 855	P3H4	2.048287e-05	8.433065e-04	2.658014
## 856	P4HA3	2.124189e-03	3.930416e-02	2.957375
## 857	PABIR3	4.882167e-05	1.743269e-03	2.799606
## 858	PADI1	9.150274e-05	2.968883e-03	-3.890382
## 859	PADI1_1	9.150274e-05	2.968883e-03	-3.890382
## 860	PAQR5	2.557773e-04	6.919379e-03	-2.510576
## 861	PAX3	2.635461e-03	4.671501e-02	6.215198
## 862	PAX6	1.636234e-08	1.932439e-06	6.834757
## 863	PAX7	4.539370e-10	8.018778e-08	8.914945
## 864	PAX9	3.969223e-07	2.961585e-05	-2.947128
## 865	PBK	9.363061e-04	2.007425e-02	2.290988
## 866	PBX2_2	1.437124e-12	4.869260e-10	22.065777
## 867	PCAT7	2.614539e-10	4.957551e-08	5.440406

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## 868      PCDH1 4.295060e-05 1.573941e-03 -2.501683
## 869      PCDHB9_1 9.832457e-04 2.084279e-02 2.413797
## 870      PCLAF 2.183426e-05 8.883277e-04 2.692258
## 871      PCP4 1.060519e-12 3.779105e-10 -5.700745
## 872      PCP4_1 1.060519e-12 3.779105e-10 -5.700745
## 873      PCSK1N 3.808541e-06 2.104677e-04 3.750836
## 874      PCSK2 1.539532e-03 3.026637e-02 -3.033709
## 875      PDE11A 2.829155e-03 4.930268e-02 4.633646
## 876      PDGFA 4.874722e-04 1.163404e-02 1.875907
## 877      PDK4 3.987510e-12 1.177341e-09 -4.453273
## 878      PDLIM2 7.854303e-07 5.339893e-05 -2.656734
## 879      PDZRN3 8.008219e-04 1.758915e-02 -2.401064
## 880      PDZRN4 4.744434e-09 6.409017e-07 -4.964039
## 881      PFN2 7.966882e-08 7.803769e-06 3.263089
## 882      PGM2L1 6.686159e-06 3.429021e-04 2.712992
## 883      PGM5 4.475768e-17 4.405008e-14 -4.915019
## 884      PGM5-AS1 1.357958e-11 3.598242e-09 -6.574068
## 885      PGM5P4 3.151616e-05 1.201801e-03 -5.638823
## 886      PHACTR2 4.100632e-05 1.513426e-03 -2.293035
## 887      PHACTR4 2.700326e-04 7.219965e-03 -1.856987
## 888      PHLDB3 2.186043e-03 4.017200e-02 -2.147347
## 889      PHYHIP 2.911076e-09 4.207421e-07 -3.863414
## 890      PI16 1.409410e-05 6.211019e-04 -4.560170
## 891      PIMREG 3.098055e-08 3.424097e-06 2.547777
## 892      PITX1 2.381226e-03 4.294518e-02 -2.245791
## 893      PKD1L1 3.767618e-04 9.496235e-03 2.553979
## 894      PKHD1L1 5.106729e-05 1.816624e-03 -3.380799
## 895      PKP3 1.991059e-04 5.629441e-03 -2.008591
## 896      PLA2G2A 4.277103e-04 1.051326e-02 -4.948860
## 897      PLA2G2C 2.983984e-06 1.685117e-04 -4.638091
## 898      PLA2G4B 1.763001e-04 5.096182e-03 -2.261183
## 899      PLAAT2 1.835967e-04 5.255645e-03 4.522061
## 900      PLAU 8.924327e-08 8.539259e-06 3.487790
## 901      PLCXD3 2.340611e-03 4.239768e-02 -3.340030
## 902      PLD6 5.885721e-04 1.356951e-02 2.224183
## 903      PLEKHA7 1.745834e-06 1.077101e-04 -2.312360
## 904      PLEKHG4B 9.170681e-05 2.970841e-03 2.862926
## 905      PLEKHM1 3.976752e-09 5.516209e-07 -2.722018
## 906      PLEKHM1_2 7.409799e-08 7.327547e-06 -2.506878
## 907      PLIN4 5.420294e-05 1.908944e-03 -2.621748
## 908      PLN 1.271498e-04 3.922287e-03 -2.997009
## 909      PLP1 7.572741e-16 5.589764e-13 -4.829035
## 910      PLPP4 1.949588e-05 8.074965e-04 4.244584
## 911      PMM1 1.445984e-03 2.884709e-02 -1.981647
## 912      POMC 1.243270e-03 2.530811e-02 2.856794
## 913      POPDC3 2.105389e-07 1.747558e-05 5.156923
## 914      POSTN 2.072991e-03 3.842563e-02 2.994432
## 915      POTEF_1 4.507484e-05 1.631535e-03 7.272708
## 916      POU6F2-AS2 5.564026e-05 1.942742e-03 5.163328
## 917      PPL 2.151941e-08 2.484711e-06 -3.450053
## 918      PPP1R12B 3.784147e-06 2.096803e-04 -2.239788
## 919      PPP1R1A 2.827731e-05 1.096502e-03 -3.787235
## 920      PPP1R1C 2.984095e-06 1.685117e-04 4.671039
## 921      PRAME 3.203421e-19 4.729165e-16 9.717650

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## 922	PRAME_1	3.203421e-19	4.729165e-16	9.717650
## 923	PRDM13	2.433312e-04	6.652339e-03	5.158176
## 924	PRELID3A	3.111817e-05	1.188818e-03	2.684926
## 925	PRELP	1.672549e-03	3.212862e-02	-2.698887
## 926	PROC	1.663064e-05	7.087052e-04	5.211048
## 927	PRR15	1.308674e-05	5.905608e-04	4.052911
## 928	PRR15-DT	2.206062e-03	4.038519e-02	3.373034
## 929	PRR20G	2.673459e-03	4.734795e-02	6.387081
## 930	PRR4	3.540414e-08	3.771819e-06	-7.109019
## 931	PRR4_1	3.540414e-08	3.771819e-06	-7.109019
## 932	PRR4_2	3.540414e-08	3.771819e-06	-7.109019
## 933	PRR5L	1.551217e-03	3.038915e-02	2.251606
## 934	PRSS21	2.440401e-06	1.432904e-04	4.829719
## 935	PRSS27	6.003224e-04	1.375550e-02	-3.582000
## 936	PRSS3	1.241568e-03	2.530644e-02	-2.889999
## 937	PRUNE2	2.569293e-09	3.803037e-07	-3.681802
## 938	PSCA	5.923937e-04	1.361271e-02	-3.259137
## 939	PTGIS	2.999263e-04	7.816995e-03	-3.399417
## 940	PTGS2	6.091846e-05	2.108983e-03	4.037431
## 941	PTH2R	5.982131e-05	2.077961e-03	6.219175
## 942	PTHLH	6.864420e-09	8.922882e-07	3.980194
## 943	PTK6	1.754722e-04	5.079354e-03	-2.678348
## 944	PTPRN	2.374273e-03	4.285718e-02	4.559153
## 945	PWWP3B	1.800291e-04	5.182231e-03	6.016714
## 946	PXDN	1.540556e-03	3.026637e-02	2.418928
## 947	PYGM	1.744267e-06	1.077101e-04	-3.147673
## 948	PYY2	1.178449e-07	1.082498e-05	6.875747
## 949	RAB15	1.014687e-03	2.133196e-02	2.093009
## 950	RAB25	1.599242e-03	3.100669e-02	-1.794638
## 951	RAB32	4.940038e-04	1.176275e-02	2.280534
## 952	RAB3B	6.643433e-04	1.494086e-02	3.337368
## 953	RAB40A	1.427837e-16	1.229606e-13	-5.250004
## 954	RAB9B	2.807589e-03	4.910256e-02	-2.611853
## 955	RABGGTA	6.763207e-05	2.321960e-03	-2.058172
## 956	RABGGTA_1	6.763207e-05	2.321960e-03	-2.058172
## 957	RAD51AP1	1.063589e-07	9.946724e-06	3.231669
## 958	RAET1E	1.244099e-03	2.530811e-02	-3.154733
## 959	RAG1	4.545504e-04	1.102658e-02	3.249189
## 960	RANBP9	3.137407e-04	8.125804e-03	-2.065572
## 961	RARG	8.138982e-04	1.783844e-02	-1.750511
## 962	RASL11B	1.510644e-04	4.511850e-03	3.428543
## 963	RBFOX3	8.175652e-19	1.126496e-15	-5.856158
## 964	RBM47	1.036379e-03	2.168004e-02	-2.114475
## 965	RBP1	9.618717e-05	3.091752e-03	3.148141
## 966	RBP4	5.810675e-04	1.343345e-02	4.120184
## 967	RBPMS2	4.334408e-05	1.579957e-03	-2.317533
## 968	RDH12	1.096810e-09	1.784950e-07	-4.161113
## 969	RELCH	3.830260e-04	9.618934e-03	-1.913041
## 970	RFC4	1.884428e-09	2.863777e-07	2.888049
## 971	RHCG	3.213789e-05	1.221004e-03	-3.397654
## 972	RHEBL1	9.289541e-05	3.004636e-03	3.271270
## 973	RIMS2	6.223337e-07	4.287464e-05	3.488222
## 974	RMND5B	4.439403e-04	1.087311e-02	-1.838266
## 975	RNASEH2A	3.049188e-04	7.937104e-03	1.991206

## 976	RNF222	1.000872e-06	6.531486e-05	-3.856026
## 977	RNF224	9.636083e-04	2.057423e-02	-3.310112
## 978	RNF225	1.362425e-04	4.165473e-03	-3.196677
## 979	RNFT2	2.564937e-05	1.017910e-03	3.181044
## 980	ROR1	1.231616e-03	2.520301e-02	-2.688994
## 981	ROS1	4.186410e-09	5.730114e-07	6.882220
## 982	RPL39L	5.246402e-08	5.394659e-06	3.476519
## 983	RRAD	2.474235e-03	4.423658e-02	-2.920293
## 984	SALL4	2.140600e-13	9.217068e-11	4.122564
## 985	SAMD5	1.810631e-06	1.100651e-04	-3.227946
## 986	SASH1	3.540705e-10	6.533864e-08	-3.172226
## 987	SCARA5	9.760922e-05	3.132589e-03	-3.944008
## 988	SCAT2	1.141086e-03	2.367869e-02	3.885584
## 989	SCAT8	1.939462e-03	3.650710e-02	3.737606
## 990	SCEL	6.072728e-06	3.153546e-04	-3.811042
## 991	SCG2	1.131766e-04	3.533435e-03	4.312047
## 992	SCGB1D2	1.779927e-03	3.388698e-02	-4.681143
## 993	SCGB2A2	1.775094e-06	1.089309e-04	-8.033597
## 994	SCGB3A1	1.412706e-04	4.256241e-03	-5.443222
## 995	SCIRT	2.835134e-03	4.936524e-02	3.159560
## 996	SCN5A	1.711758e-03	3.278833e-02	3.380014
## 997	SCN7A	5.855129e-17	5.500627e-14	-5.361083
## 998	SCN8A	9.791385e-05	3.132637e-03	2.934254
## 999	SCN9A	1.398237e-05	6.180592e-04	4.360995
## 1000	SCNN1B	2.915825e-08	3.240014e-06	-3.301441
## 1001	SCUBE3	4.978555e-04	1.180950e-02	3.427316
## 1002	SCX	2.904901e-04	7.648215e-03	2.411566
## 1003	SEC14L4	5.436615e-05	1.910952e-03	3.646183
## 1004	SERPINB1	1.608432e-03	3.109735e-02	-2.502206
## 1005	SERPINB11	7.353050e-04	1.630610e-02	-2.673095
## 1006	SERPINB12	1.580054e-04	4.687819e-03	-3.426843
## 1007	SERPINB2	6.979814e-04	1.562934e-02	-3.454279
## 1008	SERPINE1	7.981042e-05	2.656235e-03	3.139149
## 1009	SERPINH1	4.421737e-10	7.878316e-08	2.972488
## 1010	SEZ6L2	5.889225e-04	1.356951e-02	2.929850
## 1011	SFTA2_2	1.607418e-04	4.746016e-03	-5.290253
## 1012	SGCA	1.577508e-04	4.687819e-03	-2.942742
## 1013	SH3BGRL2	8.716296e-07	5.811239e-05	-3.267472
## 1014	SHH	1.027320e-03	2.153538e-02	5.190398
## 1015	SHISA2	1.230592e-04	3.807466e-03	4.512349
## 1016	SHOX2	8.540712e-09	1.103247e-06	7.259799
## 1017	SIM2	5.726257e-04	1.328286e-02	-2.665900
## 1018	SIX1	4.069584e-20	9.345575e-17	6.588428
## 1019	SIX2	9.305482e-10	1.563624e-07	3.881640
## 1020	SIX3	2.556274e-04	6.919379e-03	5.670471
## 1021	SIX4	1.177853e-05	5.421798e-04	3.375702
## 1022	SLC13A4	2.697109e-07	2.150220e-05	-3.522330
## 1023	SLC16A10	1.444491e-04	4.339352e-03	2.606154
## 1024	SLC16A6	4.229268e-06	2.300277e-04	-2.873453
## 1025	SLC24A2	7.862712e-05	2.621073e-03	4.592379
## 1026	SLC27A4	2.566104e-04	6.926999e-03	-2.011666
## 1027	SLC27A6	4.039618e-05	1.496251e-03	-3.382189
## 1028	SLC29A4	2.610296e-03	4.634846e-02	2.548636
## 1029	SLC2A4	4.732973e-08	4.915632e-06	-3.884614

## 1030	SLC2A4_1	4.732973e-08	4.915632e-06	-3.884614
## 1031	SLC30A3	7.112559e-06	3.611852e-04	4.767068
## 1032	SLC34A3	1.671779e-04	4.908001e-03	-3.556438
## 1033	SLC35C1	1.509611e-04	4.511850e-03	-2.074358
## 1034	SLC46A2	2.098118e-08	2.436174e-06	-4.700783
## 1035	SLC51A	4.978023e-04	1.180950e-02	4.201332
## 1036	SLC52A1	2.030464e-04	5.720731e-03	3.394591
## 1037	SLC5A12	7.059891e-06	3.593937e-04	4.947956
## 1038	SLC6A1	1.237002e-03	2.526320e-02	-3.261629
## 1039	SLC6A2	1.148041e-04	3.573450e-03	4.766898
## 1040	SLC7A5P2_1	2.517320e-03	4.489040e-02	3.406795
## 1041	SLC8A1-AS1	3.178601e-08	3.494432e-06	-6.547623
## 1042	SLC01A2	2.316120e-13	9.769302e-11	7.969568
## 1043	SLC04A1-AS2	1.585798e-06	9.931900e-05	-3.369559
## 1044	SLITRK3	3.650585e-06	2.028234e-04	-4.058491
## 1045	SLMAP	1.291355e-03	2.619209e-02	-1.593220
## 1046	SLURP1	1.030992e-07	9.729929e-06	-4.450451
## 1047	SMAGP	2.163190e-06	1.288439e-04	-2.251907
## 1048	SMC1B	1.770828e-04	5.111659e-03	5.405907
## 1049	SMOC1	3.366306e-04	8.653582e-03	3.624976
## 1050	SNCAIP	2.277406e-03	4.147086e-02	2.417071
## 1051	SNCB	9.110349e-05	2.968883e-03	5.623633
## 1052	SNHG33	2.767984e-03	4.860552e-02	2.623647
## 1053	SNORD3A	1.665177e-07	1.470764e-05	7.666877
## 1054	SNX10	2.704941e-07	2.150220e-05	2.789280
## 1055	SORBS1	1.375244e-10	2.805876e-08	-3.447275
## 1056	SORCS1	9.595141e-04	2.050800e-02	-3.180708
## 1057	SORT1	1.545014e-03	3.032511e-02	-1.890219
## 1058	SOST	6.534529e-08	6.642306e-06	7.517938
## 1059	SOX10	9.061043e-06	4.355201e-04	-3.422805
## 1060	SOX11	2.826034e-03	4.928985e-02	3.967565
## 1061	SOX4	1.791994e-03	3.407262e-02	2.103704
## 1062	SPC24	1.029815e-03	2.156455e-02	2.281515
## 1063	SPC25_1	9.308849e-04	2.004118e-02	2.859116
## 1064	SPINDOC	9.993194e-04	2.112938e-02	1.723117
## 1065	SPINK5	1.776162e-06	1.089309e-04	-4.404171
## 1066	SPNS2	1.803677e-07	1.572928e-05	-3.694177
## 1067	SPOCD1	8.188026e-06	4.019718e-04	3.833984
## 1068	SPP1	5.752457e-14	2.830757e-11	6.931695
## 1069	SPRR2G	1.005618e-03	2.120828e-02	3.929969
## 1070	SPRR3	1.249474e-05	5.700689e-04	-4.119999
## 1071	SPSB4	4.143043e-04	1.023040e-02	5.065519
## 1072	SPTLC3	2.384871e-03	4.297342e-02	-2.145970
## 1073	SRL	9.786618e-04	2.077410e-02	-2.819274
## 1074	ST6GALNAC1	7.400246e-04	1.634178e-02	-2.446651
## 1075	ST8SIA2	4.536053e-04	1.101954e-02	4.729849
## 1076	STIM1	2.743660e-04	7.307471e-03	-1.659554
## 1077	STK40	1.947326e-03	3.658849e-02	-2.033342
## 1078	STMN1	5.364338e-06	2.828320e-04	2.344041
## 1079	STRA6	2.154674e-04	6.009826e-03	3.321162
## 1080	STRA6_1	2.782547e-04	7.390066e-03	3.266690
## 1081	STRN	1.827147e-04	5.237654e-03	-1.833324
## 1082	STUM	3.060956e-07	2.396358e-05	-3.342904
## 1083	STX1A	2.432045e-04	6.652339e-03	2.560746

## 1084	SULT2B1	1.194528e-05	5.486333e-04	-3.267047
## 1085	SYCP2	9.168223e-04	1.977963e-02	3.160839
## 1086	SYNDIG1	5.279756e-04	1.240023e-02	3.745334
## 1087	SYNM	9.144651e-11	1.948471e-08	-3.533343
## 1088	SYNPO2	2.933802e-18	3.789739e-15	-4.868168
## 1089	SYNPO2L	3.043494e-07	2.391746e-05	-6.231343
## 1090	SYT1	4.263939e-12	1.223987e-09	4.128886
## 1091	SYT14	1.165198e-08	1.442054e-06	7.234694
## 1092	TACR1	4.883651e-05	1.743269e-03	-2.962693
## 1093	TACR2	6.451959e-05	2.226195e-03	-3.349569
## 1094	TAGLN	1.604899e-03	3.105810e-02	-2.189254
## 1095	TBX15	1.153048e-05	5.343315e-04	3.732520
## 1096	TBX6	3.674554e-07	2.792121e-05	-2.294096
## 1097	TCF15	2.703510e-03	4.783917e-02	3.481067
## 1098	TD02	2.523041e-06	1.468907e-04	5.301138
## 1099	TD02_1	2.523041e-06	1.468907e-04	5.301138
## 1100	TEC	2.763829e-03	4.857383e-02	-2.229229
## 1101	TENT5B	1.856360e-03	3.523162e-02	-2.562881
## 1102	TESMIN	4.562923e-04	1.105586e-02	3.156128
## 1103	TFAP2B	4.044693e-04	1.007177e-02	-4.246749
## 1104	TGFBI	2.100421e-05	8.613392e-04	3.502259
## 1105	TGFBR3	8.928982e-07	5.933897e-05	-2.745227
## 1106	TGM1	2.202551e-05	8.925948e-04	-3.507608
## 1107	TGM1_1	2.202551e-05	8.925948e-04	-3.507608
## 1108	TGM3	5.897911e-09	7.915456e-07	-4.972337
## 1109	TIAM1	1.079937e-04	3.418092e-03	-2.257659
## 1110	TJP1_1	1.359637e-05	6.069327e-04	-2.338450
## 1111	TJP1_2	1.359637e-05	6.069327e-04	-2.338450
## 1112	TJP3	5.761943e-04	1.335066e-02	-2.373186
## 1113	TK1	2.452839e-03	4.393005e-02	1.961834
## 1114	TM4SF19	1.320071e-03	2.672207e-02	3.008430
## 1115	TMEFF1	1.027380e-03	2.153538e-02	4.546940
## 1116	TMEM132C	1.988571e-11	5.202505e-09	-5.480825
## 1117	TMEM154	6.763835e-08	6.786163e-06	-2.908528
## 1118	TMEM35A	2.691721e-07	2.150220e-05	-3.736828
## 1119	TMEM44-AS1	1.203836e-03	2.473248e-02	2.755132
## 1120	TMEM79	4.033289e-04	1.005549e-02	-2.554608
## 1121	TMOD3	2.365252e-03	4.276905e-02	-1.752391
## 1122	TMPRSS11B	1.802152e-03	3.423426e-02	-4.210995
## 1123	TNFAIP6	8.886655e-05	2.915387e-03	3.271895
## 1124	TNFRSF18	1.196969e-06	7.659122e-05	3.158992
## 1125	TNXB_7	1.325575e-11	3.558049e-09	-4.581801
## 1126	TOLLIP	2.015824e-03	3.756813e-02	-1.658110
## 1127	TOLLIP_1	1.870016e-03	3.545825e-02	-1.663323
## 1128	TOM1	2.812058e-04	7.451232e-03	-1.908411
## 1129	TOML1L2	2.438309e-05	9.766468e-04	-2.036659
## 1130	TOP2A	1.434798e-03	2.867931e-02	2.071932
## 1131	TPM2	4.901793e-04	1.168515e-02	-2.361833
## 1132	TPRG1	1.445363e-05	6.328976e-04	-2.529544
## 1133	TREH	2.875734e-03	4.987014e-02	-2.396361
## 1134	TREM1	1.695229e-03	3.253202e-02	2.958320
## 1135	TREM2	8.676284e-07	5.803283e-05	4.014659
## 1136	TRIM29	2.740877e-04	7.307471e-03	-2.157665
## 1137	TRIM46	2.330362e-03	4.228614e-02	2.647856

## 1138	TRIP13	3.112460e-04	8.071308e-03	1.997837
## 1139	TRPM2-AS	2.074563e-06	1.246427e-04	6.184315
## 1140	TTC39A	9.294553e-06	4.426263e-04	-2.237704
## 1141	TTC9	1.882027e-04	5.365206e-03	-2.565271
## 1142	TTYH3	2.285098e-04	6.305528e-03	2.388186
## 1143	TUBB3	7.645872e-08	7.524994e-06	3.769970
## 1144	UBE2C	1.443628e-08	1.734703e-06	2.222365
## 1145	UBE2T	9.242600e-06	4.411687e-04	3.110640
## 1146	UBL3	9.017130e-06	4.344197e-04	-2.126612
## 1147	UCHL1	1.003493e-11	2.728973e-09	5.371256
## 1148	UGT1A8	1.225521e-12	4.293061e-10	-7.320981
## 1149	UNC13B	2.480866e-03	4.431680e-02	-1.685767
## 1150	UNC5B-AS1	2.803935e-05	1.091369e-03	3.976938
## 1151	UNC93A	1.700693e-04	4.971700e-03	-3.847726
## 1152	UPK1A	4.435137e-09	6.030619e-07	-5.137915
## 1153	UPK3B	2.509729e-04	6.807230e-03	-3.618064
## 1154	UPK3B_1	2.135985e-06	1.275911e-04	-3.658145
## 1155	USP2	9.263793e-04	1.996497e-02	-2.678990
## 1156	USP32P3	2.064825e-03	3.834303e-02	2.513311
## 1157	USP6NL	1.371863e-06	8.697441e-05	-2.298400
## 1158	UTS2R	9.514596e-05	3.067781e-03	3.040311
## 1159	VAX1	7.412510e-05	2.486196e-03	5.398244
## 1160	VGF	6.023291e-04	1.377095e-02	2.908115
## 1161	VIT	3.082910e-07	2.404437e-05	-4.231284
## 1162	VPS37D	1.816291e-05	7.614423e-04	2.890571
## 1163	VPS4B	1.605215e-04	4.746016e-03	-2.031815
## 1164	VSIG10L	1.561286e-10	3.132880e-08	-4.112878
## 1165	VSIG2	1.063380e-04	3.370850e-03	-3.334456
## 1166	WVA5B2	1.326630e-06	8.462587e-05	6.115367
## 1167	WDR54	1.173555e-05	5.421798e-04	2.446509
## 1168	WDR72	3.973141e-10	7.266981e-08	4.036126
## 1169	WNK4	3.207310e-12	9.758363e-10	-5.772009
## 1170	WNT11	4.607905e-10	8.070864e-08	5.019165
## 1171	WNT2	2.528722e-03	4.501953e-02	4.186985
## 1172	XCL1	4.156820e-06	2.266838e-04	3.932025
## 1173	XIRP1	2.044729e-07	1.704051e-05	7.117296
## 1174	XIST	9.408175e-16	6.272521e-13	-24.898870
## 1175	XPNPEP2	2.837938e-04	7.510180e-03	-3.535088
## 1176	ZBTB16	5.470011e-07	3.911909e-05	-2.903755
## 1177	ZDHHC15	4.850272e-05	1.737356e-03	-2.477272
## 1178	ZDHHC21	1.449244e-03	2.888425e-02	-1.877833
## 1179	ZIC1	2.511976e-03	4.483377e-02	5.559280
## 1180	ZIC2	2.272580e-06	1.349704e-04	4.318575
## 1181	ZIC5	1.713194e-09	2.657141e-07	7.867597
## 1182	ZNF114	5.551843e-05	1.942742e-03	3.289053
## 1183	ZNF185	3.989142e-06	2.192755e-04	-3.092249
## 1184	ZNF185_1	1.828335e-08	2.134917e-06	-3.513000
## 1185	ZNF365	7.262496e-05	2.463529e-03	-3.589701
## 1186	ZNF431	1.966448e-03	3.684729e-02	-2.045058
## 1187	ZNF469	4.976252e-06	2.650752e-04	3.830037
## 1188	ZNF536	3.270465e-04	8.438698e-03	-3.547536
## 1189	ZNF584-DT	4.081939e-04	1.014400e-02	2.528976
## 1190	ZNF677	1.159294e-04	3.603050e-03	-2.738006
## 1191	ZNF750	2.602564e-03	4.625090e-02	-2.333524

```
## 1192 ZNF823 2.401370e-03 4.323302e-02 -1.767326
```

```
# Get the entrez ids for the genes in degs df
library(AnnotationDbi)
library(org.Hs.eg.db)
```

```
annot <- select(
  org.Hs.eg.db,
  keys = degs$gene_id,
  columns = "ENTREZID",
  keytype = "SYMBOL"
)
```

```
annot
```

	SYMBOL	ENTREZID
## 1	A2ML1	144568
## 2	ABCA13	154664
## 3	ABCC2	1244
## 4	ABCG4	64137
## 5	ABLIM1	3983
## 6	ABLIM3	22885
## 7	ACER1	125981
## 8	ACKR1	2532
## 9	ACKR3	57007
## 10	ACOT11	26027
## 11	ACOX1	51
## 12	ACOX2	8309
## 13	ACOX3	8310
## 14	ACP3	55
## 15	ACTG2	72
## 16	ACVR1C	130399
## 17	ADAM12	8038
## 18	ADAM23	8745
## 19	ADAM32_1	<NA>
## 20	ADAM33	80332
## 21	ADAMTS12	81792
## 22	ADAMTS14	140766
## 23	ADAMTS7	11173
## 24	ADAMTS8	11095
## 25	ADAMTS9-AS1	101929335
## 26	ADAMTS9-AS2	100507098
## 27	ADCY5	111
## 28	ADD2	119
## 29	ADGRF2P	<NA>
## 30	ADGRG6	57211
## 31	ADH1B	125
## 32	AFDN	4301
## 33	AFF3	3899
## 34	AGFG2	3268
## 35	AGTR1	185
## 36	AIF1L	83543
## 37	AKR1B1	231

## 38	AKR1C1	1645
## 39	AKR1C2	1646
## 40	AKR1C3	8644
## 41	ALDH9A1	223
## 42	ALG1L1P	200810
## 43	ALOX12	239
## 44	ALOX12P2	245
## 45	ALOX15	246
## 46	ALX1	8092
## 47	AMOTL2	51421
## 48	ANGPTL1	9068
## 49	ANKRD19P	138649
## 50	ANKRD20A11P	391267
## 51	ANKRD20A5P	440482
## 52	ANKRD20A9P	284232
## 53	ANKRD20A9P_1	<NA>
## 54	ANKRD22	118932
## 55	ANKRD34B	340120
## 56	ANO1	55107
## 57	ANXA10	11199
## 58	ANXA9	8416
## 59	AOC3	8639
## 60	AOX1	316
## 61	APLN	8862
## 62	APOC1	341
## 63	AQP5	362
## 64	AQP7P1	375719
## 65	ARHGAP10	79658
## 66	ARHGAP27	201176
## 67	ARHGAP27_2	<NA>
## 68	ARHGEF10L	55160
## 69	ARSF	416
## 70	ARTN	9048
## 71	ASB5	140458
## 72	ASCL2	430
## 73	ASPA	443
## 74	ASPG	374569
## 75	ASPN_1	<NA>
## 76	ATG9B	285973
## 77	ATOH8	84913
## 78	ATP13A4	84239
## 79	ATP1A2	477
## 80	AUNIP	79000
## 81	AURKA	6790
## 82	AURKB	9212
## 83	B3GNT8	374907
## 84	B4GALNT1	2583
## 85	B4GALNT4	338707
## 86	BAMBI	25805
## 87	BARX1	56033
## 88	BARX2	8538
## 89	BBOX1-AS1	103695435
## 90	BCAN	63827
## 91	BCAS1	8537

## 92	BICD1	636
## 93	BIRC5	332
## 94	BLNK	29760
## 95	BMP8A	353500
## 96	BMPR1B	658
## 97	BMS1P14	101929959
## 98	BMS1P23	728034
## 99	BNIPL	149428
## 100	BOC	91653
## 101	BPIFB2	80341
## 102	C15orf40	123207
## 103	C15orf62	643338
## 104	C16orf74	404550
## 105	C1orf116	79098
## 106	C1QTNF6	114904
## 107	C3orf70	285382
## 108	C6orf132	647024
## 109	C6orf15	29113
## 110	C6orf15_2	<NA>
## 111	C6orf15_3	<NA>
## 112	C6orf15_4	<NA>
## 113	C6orf15_5	<NA>
## 114	C7	730
## 115	C7orf57	136288
## 116	CA13	377677
## 117	CA9	768
## 118	CAB39L	81617
## 119	CABYR	26256
## 120	CACNA1B	774
## 121	CACNB2	783
## 122	CACNG4	27092
## 123	CADM3	57863
## 124	CALB1	793
## 125	CAMK2N2	94032
## 126	CAPN14	440854
## 127	CASC9	101805492
## 128	CASQ2	845
## 129	CBS	875
## 130	CCDC162P	221262
## 131	CCDC187	399693
## 132	CCL14	6358
## 133	CCL14_1	<NA>
## 134	CCL20	6364
## 135	CCN4	8840
## 136	CCN5	8839
## 137	CCNA1	8900
## 138	CCNP	79935
## 139	CD1E	913
## 140	CD207	50489
## 141	CD276	80381
## 142	CD300LG	146894
## 143	CD83	9308
## 144	CDC45	8318
## 145	CDC6	990

## 146	CDCA3	83461
## 147	CDCA5	113130
## 148	CDH11	1009
## 149	CDH19	28513
## 150	CDKN3	1033
## 151	CDSN_3	<NA>
## 152	CDSN_4	<NA>
## 153	CDX2	1045
## 154	CEACAM1	634
## 155	CEACAM7	1087
## 156	CECR2	27443
## 157	CENPA	1058
## 158	CENPF	1063
## 159	CENPH	64946
## 160	CENPM	79019
## 161	CENPV	201161
## 162	CEP55	55165
## 163	CES2	8824
## 164	CFAP251	144406
## 165	CFAP47	286464
## 166	CFAP53	220136
## 167	CFD	1675
## 168	CFD_1	<NA>
## 169	CGNL1	84952
## 170	CHGB	1114
## 171	CHODL	140578
## 172	CHP2	63928
## 173	CHRDL1	91851
## 174	CHRNA1	1134
## 175	CHST2	9435
## 176	CILP	8483
## 177	CILP2	148113
## 178	CIMAP2	163747
## 179	CIP2A	57650
## 180	CITED2	10370
## 181	CKS1B	1163
## 182	CKS2	1164
## 183	CLCA4	22802
## 184	CLDN17	26285
## 185	CLEC12A-AS1	400002
## 186	CLEC3B	7123
## 187	CLIC3	9022
## 188	CLSPN	63967
## 189	CNFN	84518
## 190	CNGB1	1258
## 191	CNIH2	254263
## 192	CNN1	1264
## 193	CNTN5	53942
## 194	CNTNAP2	26047
## 195	CNTNAP2_1	<NA>
## 196	COCH	1690
## 197	COL10A1	1300
## 198	COL11A1	1301
## 199	COL24A1	255631

## 200	COL27A1	85301
## 201	COL2A1	1280
## 202	COL7A1	1294
## 203	CPEB3	22849
## 204	CPLX2	10814
## 205	CPNE6	9362
## 206	CPNE6_1	<NA>
## 207	CPXM1	56265
## 208	CRABP1	1381
## 209	CRCT1	54544
## 210	CRISP2	7180
## 211	CRISP3	10321
## 212	CRMA	253868
## 213	CRNN	49860
## 214	CRTAC1	55118
## 215	CRYBG1	202
## 216	CRYBG2	55057
## 217	CSAG2_1	<NA>
## 218	CSAG3	389903
## 219	CSAG3_1	<NA>
## 220	CSF2	1437
## 221	CSRP1	1465
## 222	CST1	1469
## 223	CST2	1470
## 224	CSTB	1476
## 225	CT45A5	441521
## 226	CTHRC1	115908
## 227	CTNNA3	29119
## 228	CTSC	1075
## 229	CTTN-DT	121233926
## 230	CTTN-DT_1	<NA>
## 231	CTTNBP2	83992
## 232	CXCL1	2919
## 233	CXCL11	6373
## 234	CXCL12	6387
## 235	CXCL14	9547
## 236	CXCL5	6374
## 237	CXCL6	6372
## 238	CXCR2	3579
## 239	CYP11A1	1583
## 240	CYP11A1_1	<NA>
## 241	CYP1A1	1543
## 242	CYP26A1	1592
## 243	CYP2C18	1562
## 244	CYP2J2	1573
## 245	CYP4B1	1580
## 246	CYP4F12	66002
## 247	CYP4F22	126410
## 248	CYP4F29P	54055
## 249	CYP4F35P	284233
## 250	CYSRT1	375791
## 251	DAAM2	23500
## 252	DAPP1	27071
## 253	DEFB4B_2	<NA>

## 254	DEGS2	123099
## 255	DENNND2C	163259
## 256	DES	1674
## 257	DHCR24	1718
## 258	DHCR7-DT	129810502
## 259	DHRS2	10202
## 260	DIP2C-AS1	414235
## 261	DKK1	22943
## 262	DLGAP1-AS1	649446
## 263	DLGAP5	9787
## 264	DLL3	10683
## 265	DLX1	1745
## 266	DLX1_1	<NA>
## 267	DLX2	1746
## 268	DLX6	1750
## 269	DLX6-AS1	285987
## 270	DMP1	1758
## 271	DMRT1	1761
## 272	DNAH14	127602
## 273	DNAH17	8632
## 274	DNAH5	1767
## 275	DNMT3B	1789
## 276	DOCK9	23348
## 277	DOP1B	9980
## 278	DPF1	8193
## 279	DPYSL5	56896
## 280	DRAXIN	374946
## 281	DSG3	1830
## 282	DTL	51514
## 283	DUOX1	53905
## 284	DUSP9	1852
## 285	DUXAP10	503639
## 286	DUXAP8	503637
## 287	DUXAP9	503638
## 288	DYNAP	284254
## 289	E2F1	1869
## 290	E2F7	144455
## 291	ECM1	1893
## 292	ECRG4	84417
## 293	ECT2	1894
## 294	EGF	1950
## 295	EHD3	30845
## 296	EHF	26298
## 297	EIF5A2	56648
## 298	ELOVL4	6785
## 299	EMID1	129080
## 300	EMP1	2012
## 301	EMSY-DT	124902718
## 302	EN1	2019
## 303	EN2	2020
## 304	ENDOU	8909
## 305	ENTREP1	9413
## 306	EPB41L4A	64097
## 307	EPCAM	4072

## 308	EPGN	255324
## 309	EPHA1	2041
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## 311	EPHB2	2048
## 312	EPHX2	2053
## 313	EPHX3	79852
## 314	EPS8L1	54869
## 315	EPS8L2	64787
## 316	EREG	2069
## 317	ERFE	151176
## 318	ESM1	11082
## 319	ESPL1_1	<NA>
## 320	ETFDH	2110
## 321	EVA1A	84141
## 322	EVPL	2125
## 323	F10	2159
## 324	FABP4	2167
## 325	FADD	8772
## 326	FADS1	3992
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## 328	FADS3	3995
## 329	FALEC	100874054
## 330	FAM107A	11170
## 331	FAM135A	57579
## 332	FAM25A	643161
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## 334	FAM3D	131177
## 335	FAR2P1	440905
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## 337	FCER1A	2205
## 338	FCER1A_1	<NA>
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## 340	FEZF1	389549
## 341	FEZF1-AS1	154860
## 342	FGF12	2257
## 343	FGF19	9965
## 344	FGF5	2250
## 345	FHAD1	114827
## 346	FHL1	2273
## 347	FIBCD1	84929
## 348	FIRRE	286467
## 349	FLG	2312
## 350	FLNC	2318
## 351	FLOT1_1	<NA>
## 352	FNDC4	64838
## 353	FOXD2-AS1	84793
## 354	FOXD3-AS1	100996301
## 355	FOXI2	399823
## 356	FOXI3	344167
## 357	FOXL1	2300
## 358	FOXL2	668
## 359	FOXM1	2305
## 360	FOXO6_1	<NA>
## 361	FREM2	341640

## 362	FRMD4B	23150
## 363	FUT3_1	<NA>
## 364	FUT6	2528
## 365	FXYD1	5348
## 366	GAB1	2549
## 367	GABRA4	2557
## 368	GABRP	2568
## 369	GAL	51083
## 370	GALNT17	64409
## 371	GAP43	2596
## 372	GAST	2520
## 373	GATA4	2626
## 374	GBP6	163351
## 375	GBX2	2637
## 376	GCKR	2646
## 377	GCNT4	51301
## 378	GCOM1	145781
## 379	GDA	9615
## 380	GDF15	9518
## 381	GDF7	151449
## 382	GDPD2	54857
## 383	GDPD3	79153
## 384	GFOD2	81577
## 385	GFRA1	2674
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## 1018	SIX1	6495
## 1019	SIX2	10736
## 1020	SIX3	6496
## 1021	SIX4	51804
## 1022	SLC13A4	26266
## 1023	SLC16A10	117247
## 1024	SLC16A6	9120
## 1025	SLC24A2	25769
## 1026	SLC27A4	10999
## 1027	SLC27A6	28965
## 1028	SLC29A4	222962
## 1029	SLC2A4	6517
## 1030	SLC2A4_1	<NA>
## 1031	SLC30A3	7781
## 1032	SLC34A3	142680
## 1033	SLC35C1	55343
## 1034	SLC46A2	57864
## 1035	SLC51A	200931
## 1036	SLC52A1	55065
## 1037	SLC5A12	159963
## 1038	SLC6A1	6529
## 1039	SLC6A2	6530
## 1040	SLC7A5P2_1	<NA>
## 1041	SLC8A1-AS1	100128590
## 1042	SLCO1A2	6579
## 1043	SLCO4A1-AS2	101928465
## 1044	SLITRK3	22865
## 1045	SLMAP	7871
## 1046	SLURP1	57152
## 1047	SMAGP	57228
## 1048	SMC1B	27127
## 1049	SMOC1	64093
## 1050	SNCAIP	9627
## 1051	SNCB	6620
## 1052	SNHG33	100505687
## 1053	SNORD3A	780851
## 1054	SNX10	29887
## 1055	SORBS1	10580
## 1056	SORCS1	114815
## 1057	SORT1	6272
## 1058	SOST	50964
## 1059	SOX10	6663
## 1060	SOX11	6664
## 1061	SOX4	6659
## 1062	SPC24	147841
## 1063	SPC25_1	<NA>

## 1064	SPINDOC	144097
## 1065	SPINK5	11005
## 1066	SPNS2	124976
## 1067	SPOCD1	90853
## 1068	SPP1	6696
## 1069	SPRR2G	6706
## 1070	SPRR3	6707
## 1071	SPSB4	92369
## 1072	SPTLC3	55304
## 1073	SRL	6345
## 1074	ST6GALNAC1	55808
## 1075	ST8SIA2	8128
## 1076	STIM1	6786
## 1077	STK40	83931
## 1078	STMN1	3925
## 1079	STRA6	64220
## 1080	STRA6_1	<NA>
## 1081	STRN	6801
## 1082	STUM	375057
## 1083	STX1A	6804
## 1084	SULT2B1	6820
## 1085	SYCP2	10388
## 1086	SYNDIG1	79953
## 1087	SYNM	23336
## 1088	SYNP02	171024
## 1089	SYNP02L	79933
## 1090	SYT1	6857
## 1091	SYT14	255928
## 1092	TACR1	6869
## 1093	TACR2	6865
## 1094	TAGLN	6876
## 1095	TBX15	6913
## 1096	TBX6	6911
## 1097	TCF15	6939
## 1098	TD02	6999
## 1099	TD02_1	<NA>
## 1100	TEC	7006
## 1101	TEC	100124696
## 1102	TENT5B	115572
## 1103	TESMIN	9633
## 1104	TFAP2B	7021
## 1105	TGFBI	7045
## 1106	TGFBR3	7049
## 1107	TGM1	7051
## 1108	TGM1_1	<NA>
## 1109	TGM3	7053
## 1110	TIAM1	7074
## 1111	TJP1_1	<NA>
## 1112	TJP1_2	<NA>
## 1113	TJP3	27134
## 1114	TK1	7083
## 1115	TM4SF19	116211
## 1116	TMEFF1	8577
## 1117	TMEM132C	92293

## 1118	TMEM154	201799
## 1119	TMEM35A	59353
## 1120	TMEM44-AS1	100507297
## 1121	TMEM79	84283
## 1122	TMOD3	29766
## 1123	TMPRSS11B	132724
## 1124	TNFAIP6	7130
## 1125	TNFRSF18	8784
## 1126	TNXB_7	<NA>
## 1127	TOLLIP	54472
## 1128	TOLLIP_1	<NA>
## 1129	TOM1	10043
## 1130	TOM1L2	146691
## 1131	TOP2A	7153
## 1132	TPM2	7169
## 1133	TPRG1	285386
## 1134	TREH	11181
## 1135	TREM1	54210
## 1136	TREM2	54209
## 1137	TRIM29	23650
## 1138	TRIM46	80128
## 1139	TRIP13	9319
## 1140	TRPM2-AS	101928607
## 1141	TTC39A	22996
## 1142	TTC9	23508
## 1143	TTYH3	80727
## 1144	TUBB3	10381
## 1145	UBE2C	11065
## 1146	UBE2T	29089
## 1147	UBL3	5412
## 1148	UCHL1	7345
## 1149	UGT1A8	54576
## 1150	UNC13B	10497
## 1151	UNC5B-AS1	728978
## 1152	UNC93A	54346
## 1153	UPK1A	11045
## 1154	UPK3B	105375355
## 1155	UPK3B_1	<NA>
## 1156	USP2	9099
## 1157	USP32P3	347716
## 1158	USP6NL	9712
## 1159	UTS2R	2837
## 1160	VAX1	11023
## 1161	VGF	7425
## 1162	VIT	5212
## 1163	VPS37D	155382
## 1164	VPS4B	9525
## 1165	VSIG10L	147645
## 1166	VSIG2	23584
## 1167	WWA5B2	90113
## 1168	WDR54	84058
## 1169	WDR72	256764
## 1170	WNK4	65266
## 1171	WNT11	7481

```

## 1172          WNT2      7472
## 1173          XCL1      6375
## 1174          XIRP1    165904
## 1175          XIST      7503
## 1176          XPNPEP2   7512
## 1177          ZBTB16    7704
## 1178          ZDHHC15   158866
## 1179          ZDHHC21   340481
## 1180          ZIC1      7545
## 1181          ZIC2      7546
## 1182          ZIC5      85416
## 1183          ZNF114    163071
## 1184          ZNF185    7739
## 1185          ZNF185_1  <NA>
## 1186          ZNF365    22891
## 1187          ZNF431    170959
## 1188          ZNF469    84627
## 1189          ZNF536    9745
## 1190          ZNF584-DT  123706526
## 1191          ZNF677    342926
## 1192          ZNF750    79755
## 1193          ZNF823    55552

```

```

# Check whether there are more than one entrez id mapped to a gene
annot$n <- sapply(strsplit(as.character(annot$ENTREZID), ","), length)
idx <- which(annot$n > 2)
annot[idx, ] # Output: 0 rows

```

```

## [1] SYMBOL ENTREZID n
## <0 rows> (or 0-length row.names)

```

```
# Merge annot$ENTREZID with degs
```

```

degs <- degs %>%
  left_join(
    annot %>% dplyr::select(SYMBOL, ENTREZID),
    by = c("gene_id" = "SYMBOL")
  )

```

```
degs
```

	gene_id	P.Value	adj.P.Val	logFC	ENTREZID
## 1	A2ML1	2.573314e-08	2.890503e-06	-4.293321	144568
## 2	ABCA13	5.316538e-04	1.247244e-02	3.345934	154664
## 3	ABCC2	3.676964e-05	1.384253e-03	4.392891	1244
## 4	ABCG4	5.561698e-04	1.297395e-02	-2.724376	64137
## 5	ABLIM1	2.930047e-05	1.129817e-03	-2.430274	3983
## 6	ABLIM3	7.328133e-05	2.470764e-03	-3.245304	22885
## 7	ACER1	1.425965e-08	1.723499e-06	-5.248480	125981
## 8	ACKR1	4.453428e-05	1.614797e-03	-2.736071	2532
## 9	ACKR3	2.758081e-03	4.851406e-02	2.305740	57007
## 10	ACOT11	2.096467e-03	3.882596e-02	-2.159534	26027
## 11	ACOX1	1.412286e-04	4.256241e-03	-1.731111	51

## 12	ACOX2	1.617922e-05	6.908930e-04	-2.729130	8309
## 13	ACOX3	6.520376e-06	3.352317e-04	-2.288334	8310
## 14	ACP3	2.678986e-04	7.184753e-03	-2.778791	55
## 15	ACTG2	4.499720e-14	2.268298e-11	-4.835173	72
## 16	ACVR1C	1.389269e-05	6.180592e-04	3.812397	130399
## 17	ADAM12	5.332085e-05	1.887047e-03	3.635348	8038
## 18	ADAM23	6.673881e-08	6.728575e-06	3.126805	8745
## 19	ADAM32_1	2.425635e-03	4.359393e-02	3.392516	<NA>
## 20	ADAM33	1.302545e-04	4.000149e-03	-2.710397	80332
## 21	ADAMTS12	1.536425e-07	1.362868e-05	4.218968	81792
## 22	ADAMTS14	1.863847e-04	5.328075e-03	2.976114	140766
## 23	ADAMTS7	4.132060e-04	1.021548e-02	2.388743	11173
## 24	ADAMTS8	3.909642e-07	2.938344e-05	-4.018085	11095
## 25	ADAMTS9-AS1	1.300500e-14	7.679641e-12	-6.334097	101929335
## 26	ADAMTS9-AS2	5.473167e-04	1.279631e-02	-3.608809	100507098
## 27	ADCY5	6.178884e-07	4.271076e-05	-2.934183	111
## 28	ADD2	3.880832e-05	1.455699e-03	3.187843	119
## 29	ADGRF2P	1.471048e-03	2.920616e-02	-2.664184	<NA>
## 30	ADGRG6	2.802716e-03	4.909029e-02	-1.836851	57211
## 31	ADH1B	5.770564e-13	2.208630e-10	-5.267329	125
## 32	AFDN	2.331455e-04	6.412398e-03	-2.066169	4301
## 33	AFF3	1.069216e-09	1.753853e-07	-3.039541	3899
## 34	AGFG2	2.544316e-08	2.873548e-06	-2.589640	3268
## 35	AGTR1	2.876195e-03	4.987014e-02	-2.970563	185
## 36	AIF1L	5.478900e-05	1.922545e-03	-2.870940	83543
## 37	AKR1B1	1.570847e-03	3.059801e-02	2.135055	231
## 38	AKR1C1	4.519830e-04	1.100304e-02	3.726399	1645
## 39	AKR1C2	5.637977e-05	1.965020e-03	3.385866	1646
## 40	AKR1C3	6.955118e-04	1.559093e-02	3.769443	8644
## 41	ALDH9A1	4.083514e-04	1.014400e-02	-1.673958	223
## 42	ALG1L1P	4.285828e-10	7.770131e-08	6.114394	200810
## 43	ALOX12	1.158416e-08	1.442054e-06	-4.441886	239
## 44	ALOX12P2	1.217839e-08	1.498232e-06	3.406411	245
## 45	ALOX15	1.142238e-09	1.830061e-07	6.651260	246
## 46	ALX1	6.187678e-06	3.205186e-04	5.859546	8092
## 47	AMOTL2	5.682106e-04	1.319526e-02	-2.116939	51421
## 48	ANGPTL1	3.320303e-10	6.182345e-08	-3.884158	9068
## 49	ANKRD19P	6.541929e-04	1.474467e-02	2.808824	138649
## 50	ANKRD20A11P	3.557681e-04	9.044299e-03	-4.221272	391267
## 51	ANKRD20A5P	1.247145e-04	3.852913e-03	-3.993332	440482
## 52	ANKRD20A9P	1.403292e-04	4.246449e-03	-4.154720	284232
## 53	ANKRD20A9P_1	1.403292e-04	4.246449e-03	-4.154720	<NA>
## 54	ANKRD22	1.012590e-03	2.131182e-02	-2.053582	118932
## 55	ANKRD34B	1.632805e-04	4.807238e-03	5.244232	340120
## 56	ANO1	4.714316e-04	1.135612e-02	3.208133	55107
## 57	ANXA10	1.577296e-03	3.066750e-02	5.729620	11199
## 58	ANXA9	2.183325e-04	6.073346e-03	-3.105041	8416
## 59	AOC3	1.518325e-03	2.994346e-02	-2.395572	8639
## 60	AOX1	3.493113e-11	8.493607e-09	-4.180983	316
## 61	APLN	1.894643e-03	3.582660e-02	2.670513	8862
## 62	APOC1	1.128272e-05	5.244467e-04	3.831572	341
## 63	AQP5	1.182394e-03	2.438021e-02	-3.573483	362
## 64	AQP7P1	4.849868e-05	1.737356e-03	-4.143373	375719
## 65	ARHGAP10	1.182649e-10	2.468988e-08	-2.823413	79658

## 66	ARHGAP27	3.900443e-04	9.771438e-03	-2.180975	201176
## 67	ARHGAP27_2	1.780591e-03	3.388698e-02	-2.201105	<NA>
## 68	ARHGEF10L	1.459662e-06	9.197650e-05	-2.623614	55160
## 69	ARSF	4.766978e-05	1.713459e-03	-3.784077	416
## 70	ARTN	3.176918e-07	2.468441e-05	4.082105	9048
## 71	ASB5	7.125052e-11	1.533964e-08	-6.676772	140458
## 72	ASCL2	3.084618e-04	8.009157e-03	2.140080	430
## 73	ASPA	7.047749e-05	2.403678e-03	-3.541027	443
## 74	ASPG	9.079565e-04	1.963244e-02	-2.698803	374569
## 75	ASPN_1	1.383102e-03	2.780591e-02	3.056957	<NA>
## 76	ATG9B	1.247374e-03	2.534979e-02	-3.059404	285973
## 77	ATOH8	2.180972e-03	4.013921e-02	-2.179397	84913
## 78	ATP13A4	2.842360e-03	4.944941e-02	-2.644112	84239
## 79	ATP1A2	1.253333e-27	8.634627e-24	-5.722853	477
## 80	AUNIP	7.373391e-04	1.633368e-02	2.291721	79000
## 81	AURKA	1.131695e-04	3.533435e-03	2.334947	6790
## 82	AURKB	5.455517e-04	1.276949e-02	2.031322	9212
## 83	B3GNT8	1.930046e-03	3.639615e-02	-2.470325	374907
## 84	B4GALNT1	1.430662e-17	1.642718e-14	4.803579	2583
## 85	B4GALNT4	6.173947e-04	1.406577e-02	2.480870	338707
## 86	BAMBI	2.267190e-04	6.264477e-03	3.335898	25805
## 87	BARX1	5.179918e-11	1.176468e-08	7.907641	56033
## 88	BARX2	1.119797e-04	3.506662e-03	-3.334623	8538
## 89	BBOX1-AS1	1.693712e-04	4.958306e-03	3.418432	103695435
## 90	BCAN	2.670079e-08	2.982983e-06	4.563505	63827
## 91	BCAS1	3.397841e-04	8.723799e-03	-2.567226	8537
## 92	BICD1	1.525518e-03	3.005663e-02	1.915596	636
## 93	BIRC5	9.998321e-04	2.112938e-02	2.027227	332
## 94	BLNK	1.616126e-03	3.121691e-02	-2.391013	29760
## 95	BMP8A	2.447328e-03	4.386936e-02	2.355312	353500
## 96	BMPR1B	1.092653e-03	2.271926e-02	3.272940	658
## 97	BMS1P14	3.433392e-04	8.788536e-03	-2.667681	101929959
## 98	BMS1P23	1.610046e-04	4.746995e-03	-2.611368	728034
## 99	BNIPL	2.892537e-04	7.625377e-03	-2.758131	149428
## 100	BOC	1.705687e-04	4.979256e-03	-2.485833	91653
## 101	BPIFB2	2.727619e-03	4.818327e-02	-7.359250	80341
## 102	C15orf40	2.030561e-10	3.959211e-08	3.969106	123207
## 103	C15orf62	1.435834e-05	6.314003e-04	-3.079208	643338
## 104	C16orf74	7.270915e-05	2.463529e-03	2.810195	404550
## 105	C1orf116	2.337206e-11	6.038172e-09	-3.698696	79098
## 106	C1QTNF6	1.727401e-03	3.305734e-02	2.515549	114904
## 107	C3orf70	2.929883e-04	7.679951e-03	-2.720467	285382
## 108	C6orf132	2.440742e-03	4.378929e-02	-2.112112	647024
## 109	C6orf15	7.098018e-04	1.580771e-02	5.588939	29113
## 110	C6orf15_2	7.098018e-04	1.580771e-02	5.588939	<NA>
## 111	C6orf15_3	7.098018e-04	1.580771e-02	5.588939	<NA>
## 112	C6orf15_4	7.098018e-04	1.580771e-02	5.588939	<NA>
## 113	C6orf15_5	6.402224e-04	1.447101e-02	8.584877	<NA>
## 114	C7	6.192494e-21	1.599831e-17	-5.700613	730
## 115	C7orf57	2.406772e-03	4.329258e-02	-3.194505	136288
## 116	CA13	4.409623e-05	1.601724e-03	-2.574776	377677
## 117	CA9	5.437573e-07	3.902214e-05	7.002867	768
## 118	CAB39L	5.270562e-06	2.793128e-04	-2.557420	81617
## 119	CABYR	4.702806e-07	3.446723e-05	3.815934	26256

## 120	CACNA1B	2.196744e-14	1.227089e-11	7.311116	774
## 121	CACNB2	1.105349e-08	1.393010e-06	-3.233534	783
## 122	CACNG4	9.425408e-06	4.478261e-04	4.026712	27092
## 123	CADM3	1.567751e-06	9.848716e-05	-3.750337	57863
## 124	CALB1	4.769728e-04	1.144956e-02	4.569876	793
## 125	CAMK2N2	2.916652e-04	7.669384e-03	2.880172	94032
## 126	CAPN14	1.572241e-03	3.059801e-02	-3.417891	440854
## 127	CASC9	6.248353e-11	1.395881e-08	6.739083	101805492
## 128	CASQ2	4.079049e-09	5.620385e-07	-4.549791	845
## 129	CBS	2.357933e-05	9.481277e-04	3.373427	875
## 130	CCDC162P	8.330274e-12	2.295601e-09	4.473031	221262
## 131	CCDC187	2.202148e-03	4.034929e-02	4.401733	399693
## 132	CCL14	2.594485e-09	3.803037e-07	-3.920689	6358
## 133	CCL14_1	2.594485e-09	3.803037e-07	-3.920689	<NA>
## 134	CCL20	3.797190e-04	9.547485e-03	3.969695	6364
## 135	CCN4	3.745370e-04	9.463241e-03	3.223451	8840
## 136	CCN5	7.291776e-07	4.973810e-05	-3.655769	8839
## 137	CCNA1	9.727653e-07	6.382576e-05	5.177291	8900
## 138	CCNP	1.802282e-06	1.098807e-04	5.796424	79935
## 139	CD1E	7.393371e-04	1.634178e-02	-3.201493	913
## 140	CD207	1.851222e-05	7.729504e-04	-3.468188	50489
## 141	CD276	1.382656e-04	4.205689e-03	2.302272	80381
## 142	CD300LG	8.900422e-09	1.142571e-06	-5.789410	146894
## 143	CD83	2.299800e-03	4.184178e-02	1.943130	9308
## 144	CDC45	4.965947e-06	2.650752e-04	2.018231	8318
## 145	CDC6	2.503944e-06	1.466048e-04	2.646758	990
## 146	CDCA3	1.561002e-03	3.052299e-02	2.057956	83461
## 147	CDCA5	5.011337e-04	1.183706e-02	2.009932	113130
## 148	CDH11	5.625934e-04	1.309738e-02	2.886782	1009
## 149	CDH19	1.690302e-04	4.955343e-03	-2.996838	28513
## 150	CDKN3	7.529065e-06	3.731672e-04	3.185739	1033
## 151	CDSN_3	9.790002e-04	2.077410e-02	5.565901	<NA>
## 152	CDSN_4	9.790002e-04	2.077410e-02	5.565901	<NA>
## 153	CDX2	4.993953e-04	1.180950e-02	6.244599	1045
## 154	CEACAM1	1.336355e-05	6.004299e-04	-3.187936	634
## 155	CEACAM7	1.580903e-04	4.687819e-03	-3.955103	1087
## 156	CECR2	5.511930e-04	1.287238e-02	2.898726	27443
## 157	CENPA	1.580780e-05	6.816360e-04	2.440273	1058
## 158	CENPF	1.048784e-04	3.329688e-03	2.116781	1063
## 159	CENPH	1.474666e-03	2.924990e-02	1.894062	64946
## 160	CENPM	1.972823e-03	3.689982e-02	2.001024	79019
## 161	CENPV	2.746832e-03	4.839857e-02	2.332087	201161
## 162	CEP55	1.507467e-03	2.978616e-02	1.776009	55165
## 163	CES2	1.042277e-07	9.791722e-06	-2.966992	8824
## 164	CFAP251	1.145070e-10	2.414930e-08	3.881225	144406
## 165	CFAP47	1.214480e-05	5.563847e-04	5.487401	286464
## 166	CFAP53	1.293823e-04	3.979275e-03	3.909695	220136
## 167	CFD	2.173576e-07	1.782677e-05	-2.904503	1675
## 168	CFD_1	2.173576e-07	1.782677e-05	-2.904503	<NA>
## 169	CGNL1	2.220889e-05	8.965105e-04	-3.158161	84952
## 170	CHGB	1.361649e-03	2.745615e-02	4.803079	1114
## 171	CHODL	1.109903e-03	2.305474e-02	3.226112	140578
## 172	CHP2	2.878407e-04	7.597818e-03	3.586071	63928
## 173	CHRDL1	3.648760e-13	1.479285e-10	-5.102809	91851

## 174	CHRNA1	9.113720e-05	2.968883e-03	5.610142	1134
## 175	CHST2	8.357707e-05	2.754977e-03	2.965152	9435
## 176	CILP	4.409699e-11	1.012663e-08	-4.203096	8483
## 177	CILP2	9.968839e-04	2.111024e-02	2.848603	148113
## 178	CIMAP2	1.793239e-05	7.533062e-04	-3.154746	163747
## 179	CIP2A	1.139882e-04	3.553406e-03	2.312590	57650
## 180	CITED2	4.155031e-04	1.024776e-02	-2.096602	10370
## 181	CKS1B	1.558834e-04	4.649059e-03	2.558733	1163
## 182	CKS2	2.142588e-04	5.984191e-03	3.131849	1164
## 183	CLCA4	4.381302e-07	3.257293e-05	-4.382132	22802
## 184	CLDN17	1.012485e-04	3.224717e-03	-3.949774	26285
## 185	CLEC12A-AS1	1.044762e-13	4.907531e-11	5.095812	400002
## 186	CLEC3B	8.441435e-04	1.842319e-02	-2.769500	7123
## 187	CLIC3	2.924655e-04	7.679951e-03	-3.354401	9022
## 188	CLSPN	3.832839e-06	2.112456e-04	2.324454	63967
## 189	CNFN	9.185480e-08	8.748640e-06	-4.096514	84518
## 190	CNGB1	1.065830e-03	2.220623e-02	4.438166	1258
## 191	CNIH2	1.347200e-03	2.719134e-02	2.766163	254263
## 192	CNN1	2.380517e-16	1.892328e-13	-4.508997	1264
## 193	CNTN5	9.140337e-04	1.974007e-02	4.820636	53942
## 194	CNTNAP2	4.379297e-04	1.074956e-02	3.855783	26047
## 195	CNTNAP2_1	2.212718e-03	4.047120e-02	3.162057	<NA>
## 196	COCH	1.334577e-08	1.622531e-06	4.014747	1690
## 197	COL10A1	7.151233e-17	6.426160e-14	8.377233	1300
## 198	COL11A1	6.034145e-15	3.708683e-12	8.011539	1301
## 199	COL24A1	2.392427e-04	6.557914e-03	3.178407	255631
## 200	COL27A1	1.563547e-03	3.054385e-02	2.455225	85301
## 201	COL2A1	1.535421e-05	6.652849e-04	6.912131	1280
## 202	COL7A1	8.507781e-04	1.854840e-02	2.132210	1294
## 203	CPEB3	7.424140e-06	3.688944e-04	-2.359492	22849
## 204	CPLX2	4.959148e-04	1.179467e-02	6.436563	10814
## 205	CPNE6	8.028517e-05	2.663457e-03	-5.537337	9362
## 206	CPNE6_1	8.028517e-05	2.663457e-03	-5.537337	<NA>
## 207	CPXM1	1.422415e-03	2.851452e-02	2.855149	56265
## 208	CRABP1	1.054582e-03	2.199405e-02	5.228691	1381
## 209	CRCT1	4.585677e-07	3.373720e-05	-4.475599	54544
## 210	CRISP2	2.012905e-05	8.303938e-04	-5.036277	7180
## 211	CRISP3	3.130142e-11	7.889485e-09	-5.865594	10321
## 212	CRMA	6.199882e-04	1.406577e-02	-3.599019	253868
## 213	CRNN	6.126899e-05	2.117571e-03	-5.308179	49860
## 214	CRTAC1	1.252044e-03	2.541969e-02	-3.497130	55118
## 215	CRYBG1	1.714009e-06	1.067022e-04	-2.662419	202
## 216	CRYBG2	3.205208e-04	8.280655e-03	-2.968805	55057
## 217	CSAG2_1	5.076371e-07	3.681348e-05	7.902203	<NA>
## 218	CSAG3	5.937133e-06	3.090898e-04	7.234290	389903
## 219	CSAG3_1	3.206431e-05	1.220452e-03	6.647558	<NA>
## 220	CSF2	1.198029e-03	2.463768e-02	5.552246	1437
## 221	CSRP1	7.114144e-05	2.422325e-03	-2.060090	1465
## 222	CST1	1.920540e-19	3.312485e-16	11.923317	1469
## 223	CST2	1.786405e-04	5.149431e-03	5.555372	1470
## 224	CSTB	3.308605e-07	2.551577e-05	-3.391982	1476
## 225	CT45A5	2.201478e-03	4.034929e-02	5.827646	441521
## 226	CTHRC1	6.071635e-07	4.218224e-05	4.517651	115908
## 227	CTNNA3	2.325376e-04	6.408115e-03	-4.538274	29119

## 228	CTSC	2.057773e-03	3.828087e-02	2.200636	1075
## 229	CTTN-DT	1.712536e-04	4.985167e-03	3.415402	121233926
## 230	CTTN-DT_1	1.712536e-04	4.985167e-03	3.415402	<NA>
## 231	CTTNBP2	1.784483e-07	1.562784e-05	-3.887677	83992
## 232	CXCL1	2.429952e-05	9.751894e-04	4.126697	2919
## 233	CXCL11	8.413262e-05	2.768874e-03	4.337692	6373
## 234	CXCL12	1.174842e-04	3.645892e-03	-2.788505	6387
## 235	CXCL14	4.873978e-04	1.163404e-02	3.026065	9547
## 236	CXCL5	1.442096e-05	6.328077e-04	3.979228	6374
## 237	CXCL6	9.331315e-05	3.013431e-03	3.963683	6372
## 238	CXCR2	4.741313e-10	8.234745e-08	-4.041847	3579
## 239	CYP11A1	7.321300e-05	2.470764e-03	-3.650514	1583
## 240	CYP11A1_1	7.321300e-05	2.470764e-03	-3.650514	<NA>
## 241	CYP1A1	4.566759e-06	2.464380e-04	-5.160696	1543
## 242	CYP26A1	3.511465e-09	4.937072e-07	9.428985	1592
## 243	CYP2C18	5.194153e-04	1.221306e-02	-2.844339	1562
## 244	CYP2J2	2.231225e-03	4.073760e-02	-2.511379	1573
## 245	CYP4B1	2.680204e-04	7.184753e-03	-3.010827	1580
## 246	CYP4F12	7.127442e-05	2.422861e-03	-2.717821	66002
## 247	CYP4F22	3.382853e-06	1.894764e-04	-4.389495	126410
## 248	CYP4F29P	2.707595e-04	7.230049e-03	-4.192720	54055
## 249	CYP4F35P	5.144243e-04	1.210948e-02	-4.083722	284233
## 250	CYSRT1	1.583052e-05	6.816360e-04	-3.877197	375791
## 251	DAAM2	1.033146e-05	4.875128e-04	-2.827132	23500
## 252	DAPP1	4.104112e-04	1.015854e-02	-2.358937	27071
## 253	DEFB4B_2	1.225680e-03	2.510640e-02	4.883737	<NA>
## 254	DEGS2	9.447866e-07	6.258606e-05	-2.898118	123099
## 255	DENND2C	2.949814e-04	7.707555e-03	-2.147567	163259
## 256	DES	2.200340e-33	2.273832e-29	-5.823757	1674
## 257	DHCR24	7.474934e-04	1.647036e-02	-1.712001	1718
## 258	DHCR7-DT	4.277941e-04	1.051326e-02	3.296091	129810502
## 259	DHRS2	1.384746e-10	2.805876e-08	5.897005	10202
## 260	DIP2C-AS1	6.502896e-05	2.240031e-03	-3.469270	414235
## 261	DKK1	7.400767e-04	1.634178e-02	3.281840	22943
## 262	DLGAP1-AS1	7.635522e-04	1.680628e-02	2.398370	649446
## 263	DLGAP5	1.639670e-04	4.820582e-03	2.312016	9787
## 264	DLL3	1.707995e-03	3.274660e-02	4.747036	10683
## 265	DLX1	4.009265e-05	1.488488e-03	5.258218	1745
## 266	DLX1_1	4.009265e-05	1.488488e-03	5.258218	<NA>
## 267	DLX2	1.231073e-07	1.125832e-05	6.231012	1746
## 268	DLX6	1.390444e-12	4.789617e-10	7.220500	1750
## 269	DLX6-AS1	5.351712e-07	3.867454e-05	6.098045	285987
## 270	DMP1	9.785835e-04	2.077410e-02	4.826783	1758
## 271	DMRT1	1.970297e-03	3.688595e-02	5.252268	1761
## 272	DNAH14	1.742569e-06	1.077101e-04	3.360521	127602
## 273	DNAH17	6.193482e-04	1.406577e-02	3.865795	8632
## 274	DNAH5	3.576129e-04	9.080029e-03	3.512767	1767
## 275	DNMT3B	3.206351e-11	7.984200e-09	2.935475	1789
## 276	DOCK9	1.366046e-07	1.227541e-05	-2.201899	23348
## 277	DOP1B	6.721441e-04	1.508347e-02	-2.007140	9980
## 278	DPF1	5.549497e-07	3.918363e-05	3.464317	8193
## 279	DPYSL5	2.323398e-03	4.219683e-02	6.130886	56896
## 280	DRAXIN	2.148479e-03	3.964712e-02	2.887502	374946
## 281	DSG3	1.566619e-03	3.057495e-02	-2.296234	1830

## 282	DTL	1.183150e-03	2.438021e-02	2.162144	51514
## 283	DUOX1	6.406509e-04	1.447101e-02	-2.576857	53905
## 284	DUSP9	1.327409e-05	5.977099e-04	4.574561	1852
## 285	DUXAP10	2.740257e-06	1.577846e-04	4.947445	503639
## 286	DUXAP8	5.937322e-04	1.361960e-02	4.372594	503637
## 287	DUXAP9	1.172300e-07	1.081656e-05	4.197566	503638
## 288	DYNAP	2.813829e-03	4.911842e-02	-3.469109	284254
## 289	E2F1	4.641661e-04	1.120723e-02	2.319064	1869
## 290	E2F7	1.568869e-03	3.058998e-02	2.096709	144455
## 291	ECM1	1.926419e-07	1.645257e-05	-3.988249	1893
## 292	ECRG4	5.931261e-07	4.141463e-05	-4.068215	84417
## 293	ECT2	3.769425e-05	1.416481e-03	1.963196	1894
## 294	EGF	1.382770e-04	4.205689e-03	3.352511	1950
## 295	EHD3	4.886588e-08	5.049800e-06	-3.195210	30845
## 296	EHF	1.110833e-04	3.504450e-03	-2.245851	26298
## 297	EIF5A2	1.960032e-03	3.676037e-02	1.952134	56648
## 298	ELOVL4	1.994715e-07	1.685641e-05	-2.963806	6785
## 299	EMID1	7.411824e-05	2.486196e-03	3.531609	129080
## 300	EMP1	6.115482e-13	2.298087e-10	-3.946682	2012
## 301	EMSY-DT	3.507450e-04	8.927584e-03	2.952592	124902718
## 302	EN1	1.474718e-09	2.326678e-07	6.308133	2019
## 303	EN2	2.437165e-03	4.376311e-02	3.507301	2020
## 304	ENDOU	3.835217e-12	1.148786e-09	-5.382425	8909
## 305	ENTREP1	3.598046e-06	2.004431e-04	-2.811934	9413
## 306	EPB41L4A	1.575207e-04	4.687819e-03	-1.956706	64097
## 307	EPCAM	9.787774e-05	3.132637e-03	2.997321	4072
## 308	EPGN	1.114822e-04	3.504450e-03	-3.633575	255324
## 309	EPHA1	7.392239e-06	3.688944e-04	-2.597286	2041
## 310	EPHA1_1	7.392239e-06	3.688944e-04	-2.597286	<NA>
## 311	EPHB2	9.797420e-08	9.288673e-06	3.774833	2048
## 312	EPHX2	4.243490e-05	1.557805e-03	-2.691414	2053
## 313	EPHX3	1.215277e-03	2.491801e-02	-2.764085	79852
## 314	EPS8L1	1.104730e-04	3.491218e-03	-3.436726	54869
## 315	EPS8L2	6.798694e-05	2.330272e-03	-2.493373	64787
## 316	EREG	1.166256e-03	2.412830e-02	-3.501221	2069
## 317	ERFE	1.199550e-13	5.509399e-11	5.773914	151176
## 318	ESM1	6.668553e-11	1.450796e-08	4.594029	11082
## 319	ESPL1_1	2.751591e-05	1.077081e-03	-2.679451	<NA>
## 320	ETFDH	5.411543e-05	1.908944e-03	-2.196594	2110
## 321	EVA1A	1.778663e-05	7.487046e-04	3.765410	84141
## 322	EVPL	2.888457e-05	1.115862e-03	-2.835649	2125
## 323	F10	1.072602e-03	2.232480e-02	-2.558238	2159
## 324	FABP4	6.029672e-05	2.090961e-03	6.104357	2167
## 325	FADD	5.633620e-04	1.309738e-02	2.930394	8772
## 326	FADS1	3.266785e-08	3.553574e-06	3.344929	3992
## 327	FADS2	3.254258e-07	2.519064e-05	3.843030	9415
## 328	FADS3	1.008979e-03	2.125747e-02	2.100228	3995
## 329	FALEC	1.311990e-03	2.658451e-02	-3.179931	100874054
## 330	FAM107A	2.582856e-05	1.021630e-03	-3.042066	11170
## 331	FAM135A	8.982869e-04	1.948142e-02	-1.968922	57579
## 332	FAM25A	1.754910e-03	3.352169e-02	-3.286055	643161
## 333	FAM3B	3.550996e-07	2.708191e-05	-3.749207	54097
## 334	FAM3D	8.920157e-06	4.327742e-04	-3.042609	131177
## 335	FAR2P1	2.785398e-04	7.390066e-03	7.708595	440905

## 336	FBN2	6.324016e-09	8.432566e-07	4.851673	2201
## 337	FCER1A	2.135612e-03	3.944488e-02	-3.319893	2205
## 338	FCER1A_1	2.135612e-03	3.944488e-02	-3.319893	<NA>
## 339	FCH02	4.011458e-05	1.488488e-03	-2.262602	115548
## 340	FEZF1	8.334211e-05	2.751621e-03	6.130205	389549
## 341	FEZF1-AS1	6.082014e-07	4.218224e-05	6.243414	154860
## 342	FGF12	7.333436e-06	3.688944e-04	3.383388	2257
## 343	FGF19	1.402323e-13	6.300698e-11	8.974206	9965
## 344	FGF5	1.151590e-03	2.387269e-02	4.317060	2250
## 345	FHAD1	9.705554e-04	2.067983e-02	2.983394	114827
## 346	FHL1	5.736533e-07	4.019074e-05	-3.036548	2273
## 347	FIBCD1	1.158394e-06	7.435309e-05	5.385084	84929
## 348	FIRRE	4.827620e-06	2.591617e-04	5.189665	286467
## 349	FLG	1.044296e-06	6.765987e-05	-5.023499	2312
## 350	FLNC	3.944896e-04	9.870837e-03	-2.946375	2318
## 351	FLOT1_1	9.342875e-04	2.007261e-02	2.897490	<NA>
## 352	FNDC4	1.916855e-06	1.161805e-04	-3.153353	64838
## 353	FOXD2-AS1	8.972985e-06	4.342276e-04	3.244359	84793
## 354	FOXD3-AS1	1.741751e-03	3.330112e-02	4.171909	100996301
## 355	FOXI2	2.422558e-07	1.963507e-05	-5.026217	399823
## 356	FOXI3	1.001781e-06	6.531486e-05	6.510808	344167
## 357	FOXL1	6.457815e-06	3.328432e-04	4.268553	2300
## 358	FOXL2	1.177326e-05	5.421798e-04	3.883344	668
## 359	FOXM1	2.036442e-04	5.726420e-03	2.070474	2305
## 360	FOXO6_1	2.720452e-03	4.809778e-02	2.883230	<NA>
## 361	FREM2	5.320575e-06	2.812420e-04	6.748405	341640
## 362	FRMD4B	9.178679e-06	4.391318e-04	-2.075524	23150
## 363	FUT3_1	6.662322e-07	4.574647e-05	-2.948489	<NA>
## 364	FUT6	6.100989e-15	3.708683e-12	-5.359433	2528
## 365	FXYD1	1.960107e-07	1.667140e-05	-3.317661	5348
## 366	GAB1	6.214155e-04	1.408269e-02	-1.781714	2549
## 367	GABRA4	7.356076e-05	2.476146e-03	-4.764576	2557
## 368	GABRP	7.030203e-05	2.401657e-03	-3.348376	2568
## 369	GAL	1.915175e-03	3.618174e-02	3.822356	51083
## 370	GALNT17	2.575261e-06	1.490910e-04	-3.219810	64409
## 371	GAP43	4.814730e-04	1.153080e-02	4.145143	2596
## 372	GAST	1.383718e-04	4.205689e-03	5.874735	2520
## 373	GATA4	1.682512e-05	7.150995e-04	6.294877	2626
## 374	GBP6	1.242332e-07	1.131124e-05	-3.403819	163351
## 375	GBX2	3.225447e-08	3.527171e-06	6.659396	2637
## 376	GCKR	1.436394e-03	2.868348e-02	-3.574756	2646
## 377	GCNT4	8.034691e-07	5.426830e-05	-2.359450	51301
## 378	GCOM1	5.881366e-06	3.069598e-04	-4.147293	145781
## 379	GDA	9.138959e-06	4.382460e-04	4.235641	9615
## 380	GDF15	5.564657e-05	1.942742e-03	4.510794	9518
## 381	GDF7	1.129179e-05	5.244467e-04	-2.969267	151449
## 382	GDPD2	5.819830e-04	1.343958e-02	3.872908	54857
## 383	GDPD3	6.913378e-08	6.869505e-06	-3.643773	79153
## 384	GFOD2	1.593199e-03	3.094759e-02	-2.041622	81577
## 385	GFRA1	2.979166e-04	7.774420e-03	-3.071485	2674
## 386	GGT6	2.544546e-06	1.477266e-04	-2.403564	124975
## 387	GINS1	1.327456e-03	2.684526e-02	1.869682	9837
## 388	GINS2	5.628778e-08	5.759187e-06	2.482497	51659
## 389	GJB7	2.436688e-11	6.217464e-09	6.599639	375519

## 390	GLI1	3.524557e-05	1.329298e-03	4.407792	2735
## 391	GLI2	1.753106e-04	5.079354e-03	3.164524	2736
## 392	GLTP	2.859525e-06	1.632615e-04	-2.422314	51228
## 393	GNG7	9.354642e-04	2.007425e-02	-2.552658	2788
## 394	GPC2	9.342494e-04	2.007261e-02	3.008774	221914
## 395	GPD1L	2.698436e-07	2.150220e-05	-2.264351	23171
## 396	GPIHBP1	9.854119e-06	4.660525e-04	-3.145514	338328
## 397	GPR149	1.511222e-03	2.983185e-02	6.332652	344758
## 398	GPR158	8.925844e-05	2.923603e-03	6.011792	57512
## 399	GPR19	8.425959e-04	1.842319e-02	2.810312	2842
## 400	GPR50	8.588241e-04	1.870409e-02	4.517511	9248
## 401	GPT	4.861326e-06	2.602950e-04	-3.292211	2875
## 402	GPT2	7.713634e-04	1.696015e-02	-2.076855	84706
## 403	GPX3	2.222164e-04	6.156526e-03	-2.664937	2878
## 404	GREM2	1.916465e-05	7.953716e-04	-3.623787	64388
## 405	GRHL1	6.196216e-04	1.406577e-02	-2.308948	29841
## 406	GRHL3	2.740694e-06	1.577846e-04	-2.953343	57822
## 407	GRIN2D	8.973056e-05	2.934416e-03	3.737972	2906
## 408	GRM4	1.228695e-04	3.807297e-03	5.264138	2914
## 409	GRP	1.902890e-04	5.405488e-03	5.865513	2922
## 410	GSC	5.397963e-04	1.264910e-02	5.295591	145258
## 411	GTF2H2	1.424354e-03	2.852573e-02	-6.106284	2966
## 412	GTF3C2-AS2	4.317188e-06	2.341933e-04	3.212989	105374363
## 413	GUCY1B2	2.812704e-03	4.911842e-02	4.496990	2974
## 414	GYS2	2.948686e-09	4.211868e-07	-5.079477	2998
## 415	H2AC19	7.614423e-12	2.147125e-09	21.362969	723790
## 416	H2BC9	2.194906e-03	4.028803e-02	2.483033	8345
## 417	H3C8	2.528918e-03	4.501953e-02	4.307085	8355
## 418	HAGHL	4.481042e-04	1.093438e-02	2.241769	84264
## 419	HAGLR	9.701335e-04	2.067983e-02	2.430394	401022
## 420	HAGLROS	1.520283e-05	6.601094e-04	4.681027	102800310
## 421	HAP1	4.298065e-13	1.700489e-10	4.540555	9001
## 422	HCG22_1	2.014483e-07	1.685641e-05	-7.966206	<NA>
## 423	HCG22_2	2.014483e-07	1.685641e-05	-7.966206	<NA>
## 424	HCG22_3	2.014483e-07	1.685641e-05	-7.966206	<NA>
## 425	HCG22_6	2.122237e-07	1.754496e-05	-6.765059	<NA>
## 426	HCG22_7	3.415320e-04	8.757795e-03	-10.678966	<NA>
## 427	HES6	4.097620e-06	2.240466e-04	3.103272	55502
## 428	HEY1	1.292486e-08	1.580657e-06	4.315672	23462
## 429	HHIP	4.698340e-05	1.694682e-03	6.137370	64399
## 430	HHIPL2	1.187642e-03	2.444839e-02	3.986500	79802
## 431	HJURP	2.884248e-06	1.642194e-04	2.159983	55355
## 432	HLA-DOA_5	2.359253e-03	4.269794e-02	-4.687193	<NA>
## 433	HLA-DQB2_3	1.878587e-03	3.555553e-02	-3.314880	<NA>
## 434	HLA-DQB2_5	1.878587e-03	3.555553e-02	-3.314880	<NA>
## 435	HMCN2	1.286046e-04	3.961253e-03	-2.618386	256158
## 436	HMGA2	1.763324e-09	2.699583e-07	4.956659	8091
## 437	HMGB3	4.012457e-06	2.199721e-04	2.301583	3149
## 438	HMGCS2	3.650258e-13	1.479285e-10	-6.050288	3158
## 439	HMMR	1.048631e-05	4.925704e-04	2.862831	3161
## 440	HMX1	1.463395e-03	2.908217e-02	5.532836	3166
## 441	HMX1_1	1.463395e-03	2.908217e-02	5.532836	<NA>
## 442	HOMER3	8.648294e-05	2.841700e-03	1.989510	9454
## 443	HOTAIR	3.980416e-04	9.938781e-03	5.537591	100124700

## 444	HOXA10	6.706960e-16	5.134053e-13	5.884920	3206
## 445	HOXA10-AS	2.192250e-07	1.790886e-05	5.746006	100874323
## 446	HOXA11	1.152148e-08	1.442054e-06	6.793580	3207
## 447	HOXA13	9.130063e-10	1.546723e-07	7.113792	3209
## 448	HOXA9	4.982811e-04	1.180950e-02	3.707979	3205
## 449	HOXB6	2.280451e-05	9.187594e-04	2.900030	3216
## 450	HOXB7	1.646840e-14	9.454691e-12	5.055295	3217
## 451	HOXB8	4.515242e-04	1.100304e-02	5.027547	3218
## 452	HOXB9	4.566279e-07	3.373720e-05	7.661604	3219
## 453	HOXC10	2.694633e-10	5.062971e-08	6.298655	3226
## 454	HOXC11	1.450383e-05	6.337532e-04	6.339022	3227
## 455	HOXC13	3.900678e-11	9.266576e-09	6.745942	3229
## 456	HOXC13-AS	1.028874e-06	6.687033e-05	6.632678	100874366
## 457	HOXC9	1.216789e-05	5.563847e-04	3.680899	3225
## 458	HOXD10	7.684336e-18	9.342344e-15	4.878219	3236
## 459	HOXD11	2.204455e-38	4.556167e-34	6.974538	3237
## 460	HOXD13	1.658926e-17	1.804562e-14	9.440985	3239
## 461	HOXD8	4.633552e-04	1.120073e-02	1.772342	3234
## 462	HPGD	4.751970e-04	1.143141e-02	-3.190964	3248
## 463	HPSE2	1.067038e-26	5.513385e-23	-5.448579	60495
## 464	HS3ST5	2.176537e-03	4.009328e-02	6.083482	222537
## 465	HSPA6	6.034511e-04	1.378136e-02	3.475552	3310
## 466	HSPB6	1.900253e-07	1.629644e-05	-3.771991	126393
## 467	HSPB7	2.933348e-04	7.679951e-03	-2.813454	27129
## 468	HTR2C	2.349033e-06	1.387138e-04	6.256706	3358
## 469	HTR3B	3.182381e-06	1.787322e-04	-3.664048	9177
## 470	IBSP	3.041223e-09	4.305206e-07	6.860903	3381
## 471	IFI6	3.046941e-05	1.166188e-03	3.726198	2537
## 472	IFIT1	1.384376e-03	2.780591e-02	2.932107	3434
## 473	IGF2BP1	2.882556e-07	2.273918e-05	9.126923	10642
## 474	IGF2BP2	4.012507e-08	4.252846e-06	3.861938	10644
## 475	IGFBP3	1.114109e-09	1.798938e-07	4.050837	3486
## 476	IGFL1	1.504463e-04	4.511571e-03	-3.450313	374918
## 477	IL11	5.421667e-05	1.908944e-03	4.047767	3589
## 478	IL12A	1.049385e-03	2.192991e-02	-3.182216	3592
## 479	IL12A-AS1	2.916248e-06	1.655852e-04	-5.929746	101928376
## 480	IL13RA2	1.239171e-03	2.528251e-02	3.625986	3598
## 481	IL18	5.898494e-04	1.357573e-02	-2.352964	3606
## 482	IL1RN	4.763574e-05	1.713459e-03	-3.202644	3557
## 483	IL24	6.191967e-04	1.406577e-02	3.712402	11009
## 484	IL31RA	2.217981e-05	8.965105e-04	5.199156	133396
## 485	IL34	2.575633e-07	2.079421e-05	-3.110092	146433
## 486	IL36A	1.598893e-06	9.983662e-05	-4.527463	27179
## 487	IL36B	3.668278e-04	9.279800e-03	-3.231847	27177
## 488	IL36G	1.426389e-03	2.853882e-02	3.211320	56300
## 489	INHBA	1.990388e-10	3.917843e-08	5.126958	3624
## 490	IP6K3	1.663568e-09	2.604744e-07	-4.562975	117283
## 491	IRS4	2.674034e-04	7.184753e-03	5.806283	8471
## 492	ISG15	3.762948e-07	2.848814e-05	3.888150	9636
## 493	IVL	2.000339e-04	5.647953e-03	-3.487145	3713
## 494	JAKMIP3	1.462826e-07	1.303176e-05	3.659513	282973
## 495	JPH2	1.374743e-10	2.805876e-08	-3.604444	57158
## 496	KALRN	4.622513e-05	1.670247e-03	-2.681127	8997
## 497	KANK1	3.723869e-09	5.200333e-07	-2.869435	23189

## 498	KAT2B	3.957550e-07	2.961585e-05	-2.729732	8850
## 499	KAZN	2.061132e-03	3.830888e-02	-1.864385	23254
## 500	KCNA5	1.954340e-06	1.181062e-04	-4.599263	3741
## 501	KCNB1	4.029968e-14	2.082284e-11	-4.926136	3745
## 502	KCNC3	2.369560e-04	6.503861e-03	2.279744	3748
## 503	KCNG3	1.283611e-05	5.817911e-04	4.666543	170850
## 504	KCNK3	2.536901e-09	3.799469e-07	-3.113701	3777
## 505	KCNMA1	2.782414e-05	1.085036e-03	-2.569439	3778
## 506	KCNMB1	4.319061e-05	1.577144e-03	-2.902699	3779
## 507	KCNMB2-AS1	4.177586e-11	9.811630e-09	6.696176	104797538
## 508	KCNMB3	1.473185e-05	6.410059e-04	3.088068	27094
## 509	KCNS3	1.266605e-05	5.753449e-04	2.396974	3790
## 510	KIAA1549	8.698604e-04	1.890460e-02	2.798982	57670
## 511	KIF14	1.399515e-05	6.180592e-04	2.708076	9928
## 512	KIF1A	1.005837e-09	1.663091e-07	5.641306	547
## 513	KIF26B	6.913291e-05	2.365628e-03	3.476663	55083
## 514	KIF4A	9.545556e-04	2.042314e-02	1.869158	24137
## 515	KILH	2.598771e-04	7.002789e-03	5.317647	101927136
## 516	KLHDC7B	1.981999e-04	5.611500e-03	3.785920	113730
## 517	KLHDC7B-DT	2.777636e-05	1.085036e-03	4.068822	105373098
## 518	KLHDC8A	2.769617e-04	7.367109e-03	-2.763544	55220
## 519	KLK11	2.025070e-06	1.220238e-04	-2.585626	11012
## 520	KLK12	7.710714e-05	2.574556e-03	-3.361951	43849
## 521	KLK13	1.255040e-05	5.713473e-04	-3.670627	26085
## 522	KPRP	9.081003e-04	1.963244e-02	-4.343011	448834
## 523	KREMEN2	6.302511e-04	1.426728e-02	2.312493	79412
## 524	KRT13	2.987137e-05	1.147549e-03	-4.108900	3860
## 525	KRT24	3.596713e-04	9.109911e-03	-4.743405	192666
## 526	KRT3	6.554621e-09	8.684032e-07	-4.407911	3850
## 527	KRT32	4.063507e-08	4.284927e-06	-3.992439	3882
## 528	KRT4	3.634926e-11	8.735658e-09	-6.426344	3851
## 529	KRT42P	2.690451e-04	7.202881e-03	4.645720	284116
## 530	KRT75	2.954910e-09	4.211868e-07	5.826676	9119
## 531	KRT78	6.306293e-11	1.395881e-08	-5.568714	196374
## 532	KRT8	3.044149e-06	1.714345e-04	3.195239	3856
## 533	KRTAP4-1_1	8.992134e-06	4.342276e-04	5.894576	<NA>
## 534	LAMA1	5.812414e-06	3.041290e-04	3.906487	284217
## 535	LAMB4	7.641550e-13	2.820278e-10	-5.023367	22798
## 536	LAMC2	1.596060e-05	6.858083e-04	2.989928	3918
## 537	LAMP3	1.370000e-03	2.759763e-02	2.366640	27074
## 538	LAYN	1.774318e-03	3.386114e-02	2.962694	143903
## 539	LCAL1	4.076958e-17	4.213129e-14	10.149662	80078
## 540	LCN1	4.537270e-04	1.101954e-02	-5.664728	3933
## 541	LCN10	7.436361e-04	1.640285e-02	-2.486435	414332
## 542	LDB3	4.301304e-11	9.988692e-09	-3.588058	11155
## 543	LGI1	6.931562e-06	3.537322e-04	-3.775251	9211
## 544	LGI3	2.192393e-04	6.090372e-03	-2.862533	203190
## 545	LHX2	1.680459e-10	3.339589e-08	7.663619	9355
## 546	LHX5	1.141626e-07	1.058078e-05	6.444394	64211
## 547	LINC00467	7.618781e-05	2.547977e-03	3.085382	84791
## 548	LINC00491	1.883198e-05	7.831376e-04	4.095510	285708
## 549	LINC00519	1.506185e-04	4.511571e-03	3.945064	161342
## 550	LINC00622	2.739131e-03	4.830407e-02	3.460784	644242
## 551	LINC00649	9.745589e-09	1.243345e-06	2.392794	100506334

## 552	LINC00942	4.089453e-12	1.190434e-09	8.723056	100292680
## 553	LINC01088	7.194003e-04	1.598771e-02	-3.744425	100505875
## 554	LINC01234	1.134370e-07	1.056088e-05	7.488093	100506465
## 555	LINC01269	1.118680e-04	3.506662e-03	-3.043004	103695436
## 556	LINC01297-DUXAP10-NBEAP6	1.823286e-04	5.233843e-03	2.419749	115801414
## 557	LINC01305	7.092271e-04	1.580771e-02	5.128829	285084
## 558	LINC01614	3.392408e-06	1.894981e-04	5.702221	105373869
## 559	LINC01615	2.140653e-05	8.742275e-04	4.851814	101929484
## 560	LINC01873	1.452399e-03	2.891925e-02	2.275090	729348
## 561	LINC02041_1	2.845073e-03	4.945497e-02	5.340062	<NA>
## 562	LINC02043	2.385780e-12	7.704577e-10	4.801320	102724699
## 563	LINC02086	1.379844e-03	2.776886e-02	4.849915	105371809
## 564	LINC02137	1.921574e-03	3.626950e-02	4.677634	105371295
## 565	LINC02154	4.991375e-04	1.180950e-02	4.672197	109729169
## 566	LINC02159	1.604897e-03	3.105810e-02	4.926052	285629
## 567	LINC02253	7.105362e-04	1.580771e-02	2.732653	107984764
## 568	LINC02321	2.257522e-03	4.118134e-02	4.116024	105370623
## 569	LINC02457	2.372334e-03	4.285718e-02	4.842186	107984439
## 570	LINC02487	3.779649e-04	9.514955e-03	-4.049008	441178
## 571	LINC02538	1.669665e-03	3.212862e-02	-3.823813	401286
## 572	LINC02560	3.967656e-05	1.482885e-03	-2.926668	110806301
## 573	LINC02561	4.465861e-04	1.091022e-02	4.875400	110806285
## 574	LINC02577	1.560988e-03	3.052299e-02	3.735854	111216280
## 575	LINC02609	1.942509e-03	3.653118e-02	3.145915	105378853
## 576	LINC02623	1.608860e-05	6.886073e-04	-6.379374	101929445
## 577	LINC02636	1.959903e-04	5.556555e-03	2.774930	102723350
## 578	LINC02827	8.345892e-07	5.618661e-05	5.902155	105369606
## 579	LINC02884	5.961354e-05	2.074230e-03	-3.404353	105378909
## 580	LINC02893	2.822798e-06	1.620600e-04	3.617737	440173
## 581	LINC02994	8.261631e-05	2.736401e-03	5.265165	101928978
## 582	LINC02999	1.115700e-04	3.504450e-03	5.688034	102546229
## 583	LINC03011	1.340531e-03	2.708317e-02	2.982132	100289098
## 584	LINC03016	3.818535e-07	2.880346e-05	-5.737787	101927354
## 585	LINC03040	1.107416e-05	5.166607e-04	5.496827	221416
## 586	LINC03050	1.904004e-04	5.405488e-03	3.307317	126568848
## 587	LINC03057	6.852237e-06	3.505496e-04	6.089477	127379715
## 588	LIPH	4.102884e-04	1.015854e-02	-2.458580	200879
## 589	LM07	5.927733e-04	1.361271e-02	-2.460277	4008
## 590	LMOD1	7.976556e-16	5.684809e-13	-4.246920	25802
## 591	LMX1B	2.669118e-05	1.050768e-03	4.402419	4010
## 592	LOC100287072	2.679207e-05	1.052735e-03	2.803376	100287072
## 593	LOC101927293	4.631992e-04	1.120073e-02	5.282676	101927293
## 594	LOC101927469	3.465828e-05	1.309538e-03	4.476802	101927469
## 595	LOC101928051	2.045519e-03	3.808719e-02	4.106970	101928051
## 596	LOC101928391	3.327469e-05	1.261874e-03	2.788825	101928391
## 597	LOC101928844	6.773019e-09	8.859794e-07	-5.006118	101928844
## 598	LOC101928844_1	6.773019e-09	8.859794e-07	-5.006118	<NA>
## 599	LOC101929007	8.288299e-06	4.040155e-04	-4.586844	101929007
## 600	LOC101929594	1.104988e-05	5.166607e-04	2.914968	101929594
## 601	LOC101929748	2.832753e-06	1.621810e-04	6.275617	101929748
## 602	LOC102723546	2.615483e-12	8.316431e-10	7.828417	102723546
## 603	LOC102723686	2.339888e-06	1.385696e-04	-6.046804	102723686
## 604	LOC102723825	1.105177e-05	5.166607e-04	5.098425	102723825
## 605	LOC102723985	7.612839e-06	3.764166e-04	2.629702	102723985

## 606	LOC102724542	8.030317e-08	7.828801e-06	6.489403	102724542
## 607	LOC102724858	1.745144e-05	7.391113e-04	4.356653	102724858
## 608	LOC102725051	2.567293e-04	6.926999e-03	-4.378691	102725051
## 609	LOC102725238	8.850861e-04	1.921529e-02	2.434259	102725238
## 610	LOC105369147	2.070816e-04	5.807275e-03	2.534305	105369147
## 611	LOC105369266	6.994411e-07	4.786771e-05	-6.601385	105369266
## 612	LOC105369367	9.433439e-04	2.020418e-02	6.088664	105369367
## 613	LOC105369526	1.735969e-12	5.786938e-10	-6.948291	105369526
## 614	LOC105369601	8.038422e-06	3.965110e-04	3.122597	105369601
## 615	LOC105369887	3.053686e-04	7.938814e-03	5.090807	105369887
## 616	LOC105369975	2.513669e-05	1.001012e-03	-3.271354	105369975
## 617	LOC105370256	4.966846e-05	1.769910e-03	3.787447	105370256
## 618	LOC105370413	9.481329e-07	6.260706e-05	4.631686	105370413
## 619	LOC105370475	2.120212e-16	1.752822e-13	5.961380	105370475
## 620	LOC105370964	5.564694e-06	2.919063e-04	-3.761549	105370964
## 621	LOC105371175	1.722744e-09	2.657141e-07	-7.195873	105371175
## 622	LOC105371206	8.451085e-14	4.062024e-11	-6.451143	105371206
## 623	LOC105371207	3.481625e-04	8.872776e-03	-3.313280	105371207
## 624	LOC105371216	1.575425e-08	1.882132e-06	-6.089055	105371216
## 625	LOC105371217	3.481625e-04	8.872776e-03	-3.313280	105371217
## 626	LOC105371401	2.463162e-04	6.716179e-03	3.783830	105371401
## 627	LOC105371956	1.494170e-03	2.956789e-02	3.951683	105371956
## 628	LOC105372130	2.782324e-09	4.049653e-07	5.935220	105372130
## 629	LOC105373421	6.435163e-04	1.451986e-02	-3.466912	105373421
## 630	LOC105373582	7.420368e-06	3.688944e-04	3.976205	105373582
## 631	LOC105373949	3.886770e-04	9.749000e-03	5.572406	105373949
## 632	LOC105374122	2.121414e-06	1.270881e-04	6.692791	105374122
## 633	LOC105374264	1.458349e-06	9.197650e-05	2.902248	105374264
## 634	LOC105374816	2.051373e-04	5.760568e-03	5.896985	105374816
## 635	LOC105374848	1.210481e-03	2.484432e-02	-3.689391	105374848
## 636	LOC105375065	4.192623e-05	1.541871e-03	2.938649	105375065
## 637	LOC105375520	6.586259e-04	1.482841e-02	3.644871	105375520
## 638	LOC105375690	9.900106e-05	3.162525e-03	3.714075	105375690
## 639	LOC105375772	1.955389e-03	3.670661e-02	4.448575	105375772
## 640	LOC105375785	2.751768e-03	4.844423e-02	2.636012	105375785
## 641	LOC105376270	2.466468e-03	4.413589e-02	2.976998	105376270
## 642	LOC105376272	1.161395e-09	1.846439e-07	3.733960	105376272
## 643	LOC105377067	3.022947e-05	1.159151e-03	5.524255	105377067
## 644	LOC105377261	2.450276e-04	6.689870e-03	5.078891	105377261
## 645	LOC105377581	6.432302e-06	3.323570e-04	5.131912	105377581
## 646	LOC105377635	3.210609e-12	9.758363e-10	-6.152578	105377635
## 647	LOC105377911	1.807599e-04	5.196029e-03	5.230833	105377911
## 648	LOC105378030	2.141879e-05	8.742275e-04	5.681747	105378030
## 649	LOC105378675	1.052819e-03	2.197946e-02	5.232366	105378675
## 650	LOC105379524	5.320769e-05	1.886272e-03	5.985833	105379524
## 651	LOC105379539	2.172688e-03	4.005810e-02	-3.538696	105379539
## 652	LOC105379854	2.010462e-03	3.750202e-02	-3.116958	105379854
## 653	LOC107984006	1.631250e-03	3.145026e-02	-2.674685	107984006
## 654	LOC107984128	1.171757e-03	2.421788e-02	3.646776	107984128
## 655	LOC107984865_1	2.802332e-03	4.909029e-02	5.767151	<NA>
## 656	LOC107985705	2.237191e-10	4.281320e-08	6.083644	107985705
## 657	LOC107985915	1.776312e-05	7.487046e-04	-3.512855	107985915
## 658	LOC107985962	2.172559e-04	6.051543e-03	2.637088	107985962
## 659	LOC107986056	1.002006e-03	2.115369e-02	5.275134	107986056

## 660	LOC107986141	1.412440e-03	2.834204e-02	-3.384648	107986141
## 661	LOC107986764	9.468542e-10	1.578192e-07	-5.830406	107986764
## 662	LOC107987138	2.853668e-03	4.956270e-02	5.377954	107987138
## 663	LOC107987365	2.158991e-10	4.170282e-08	-7.153943	107987365
## 664	LOC107987401	4.405458e-05	1.601724e-03	-3.358731	107987401
## 665	LOC107987423	7.998650e-20	1.653161e-16	25.190987	107987423
## 666	LOC110091776	2.139237e-04	5.982917e-03	-4.121904	110091776
## 667	LOC112267868	1.730610e-07	1.522053e-05	3.971436	112267868
## 668	LOC112268114	2.787450e-03	4.890579e-02	-2.664563	112268114
## 669	LOC112268153	7.425008e-06	3.688944e-04	-3.779089	112268153
## 670	LOC122513141	1.393968e-05	6.180592e-04	-2.371848	122513141
## 671	LOC124900467	2.203934e-04	6.114216e-03	4.173674	124900467
## 672	LOC124900509	2.935534e-04	7.679951e-03	-4.391009	124900509
## 673	LOC124900586	4.457980e-04	1.090385e-02	4.167246	124900586
## 674	LOC124900799	2.641627e-05	1.041930e-03	2.761224	124900799
## 675	LOC124901002	1.344690e-05	6.028643e-04	-5.641890	124901002
## 676	LOC124901321	2.333135e-04	6.412398e-03	4.174420	124901321
## 677	LOC124901380	8.329697e-05	2.751621e-03	5.363644	124901380
## 678	LOC124901465	4.756634e-04	1.143141e-02	-4.195062	124901465
## 679	LOC124901609	1.302581e-05	5.890974e-04	2.413012	124901609
## 680	LOC124901681	1.015463e-04	3.228861e-03	2.834843	124901681
## 681	LOC124902211	4.780609e-04	1.146237e-02	5.134138	124902211
## 682	LOC124902365	8.835280e-08	8.493375e-06	5.948565	124902365
## 683	LOC124902366	2.333028e-03	4.229740e-02	4.575091	124902366
## 684	LOC124902696	3.480932e-04	8.872776e-03	4.853257	124902696
## 685	LOC124902719	1.470133e-05	6.410059e-04	4.914521	124902719
## 686	LOC124902874	3.249927e-14	1.722295e-11	5.341204	124902874
## 687	LOC124902884	2.617012e-03	4.642782e-02	4.076092	124902884
## 688	LOC124902961	6.797012e-08	6.786505e-06	3.247779	124902961
## 689	LOC124903025	1.550352e-03	3.038915e-02	5.101163	124903025
## 690	LOC124903391	1.987504e-03	3.714080e-02	3.414647	124903391
## 691	LOC124903505	1.338915e-04	4.099659e-03	-3.115815	124903505
## 692	LOC124903541	1.776107e-03	3.386400e-02	3.764075	124903541
## 693	LOC124903758	8.295246e-04	1.816167e-02	-3.491529	124903758
## 694	LOC124903859	2.846530e-04	7.523283e-03	3.511914	124903859
## 695	LOC124904198	4.872765e-04	1.163404e-02	5.455711	124904198
## 696	LOC124904253	1.609238e-05	6.886073e-04	6.332531	124904253
## 697	LOC124904420	2.881610e-05	1.115302e-03	-4.961085	124904420
## 698	LOC124904423	4.765572e-06	2.564970e-04	3.349230	124904423
## 699	LOC124904447	5.573827e-07	3.918363e-05	5.409353	124904447
## 700	LOC124904544	3.287405e-11	8.088581e-09	4.759073	124904544
## 701	LOC124904731	1.996362e-03	3.727263e-02	5.125800	124904731
## 702	LOC124904915	6.556176e-08	6.642306e-06	-6.438119	124904915
## 703	LOC124905027	2.186641e-03	4.017200e-02	4.398057	124905027
## 704	LOC124905038	8.450426e-08	8.161374e-06	6.592308	124905038
## 705	LOC124905421	9.529299e-05	3.067781e-03	4.155461	124905421
## 706	LOC124905488	5.084848e-04	1.198331e-02	4.076661	124905488
## 707	LOC124906047	2.818391e-05	1.094934e-03	-4.133062	124906047
## 708	LOC124906245	1.128560e-06	7.266381e-05	3.189047	124906245
## 709	LOC124906252	7.373334e-06	3.688944e-04	-5.490473	124906252
## 710	LOC124906274	3.290964e-08	3.561133e-06	6.422909	124906274
## 711	LOC124906315	1.820535e-07	1.580959e-05	8.092887	124906315
## 712	LOC124907726	1.757922e-05	7.430009e-04	3.322772	124907726
## 713	LOC124907737	1.232837e-03	2.520305e-02	3.671566	124907737

## 714	LOC124907854	4.360649e-13	1.700489e-10	-6.208075	124907854
## 715	LOC124907910	2.222774e-08	2.524192e-06	-4.972562	124907910
## 716	LOC124907946	1.686445e-08	1.980423e-06	6.410888	124907946
## 717	LOC408186	5.629873e-04	1.309738e-02	2.922080	408186
## 718	LOC440910	1.635978e-03	3.151202e-02	5.746305	440910
## 719	LOC730101	1.938495e-03	3.650710e-02	3.148668	730101
## 720	LOXL2	7.966121e-07	5.398157e-05	3.062593	4017
## 721	LPIN1	8.711825e-06	4.236612e-04	-2.288133	23175
## 722	LRP10	9.526271e-06	4.515802e-04	-1.939655	26020
## 723	LRP12	3.320735e-04	8.557725e-03	2.409885	29967
## 724	RRRC15	9.684445e-07	6.374462e-05	5.414281	131578
## 725	LYPD1	1.626356e-03	3.138518e-02	3.990182	116372
## 726	LYPD2	5.548029e-07	3.918363e-05	-4.132497	137797
## 727	LYPD3	8.440537e-04	1.842319e-02	-2.620930	27076
## 728	LYVE1	1.015611e-03	2.133196e-02	-2.797236	10894
## 729	MAB21L2	1.788005e-06	1.093328e-04	-5.000498	10586
## 730	MAB21L4	2.730976e-03	4.820137e-02	-3.259686	79919
## 731	MACC1	1.596699e-03	3.098645e-02	-2.164663	346389
## 732	MAFA	4.440151e-04	1.087311e-02	4.100720	389692
## 733	MAGEA1	3.890719e-05	1.456764e-03	9.624323	4100
## 734	MAGEA11	3.588390e-04	9.099980e-03	5.460437	4110
## 735	MAGEA3	7.284244e-06	3.680947e-04	9.854543	4102
## 736	MAGEA3_1	1.039281e-05	4.892908e-04	9.651252	<NA>
## 737	MAGEA4	2.485773e-05	9.918137e-04	11.350465	4103
## 738	MAGEA6_1	2.739448e-05	1.074363e-03	11.168207	<NA>
## 739	MAL	8.215525e-06	4.023660e-04	-4.917325	4118
## 740	MALL	2.565954e-05	1.017910e-03	-3.043013	7851
## 741	MALLP2	6.019467e-04	1.377095e-02	-3.129838	105374855
## 742	MAPK12	4.809211e-10	8.283065e-08	3.302408	6300
## 743	MAPT	4.396212e-10	7.878316e-08	-4.144386	4137
## 744	MAPT_1	5.262637e-06	2.793128e-04	-4.255444	<NA>
## 745	MAPT_2	1.461375e-07	1.303176e-05	-3.516632	<NA>
## 746	MARCKSL1	1.275182e-07	1.155941e-05	2.569228	65108
## 747	MC1R	6.127184e-04	1.397755e-02	2.324406	4157
## 748	MCCC2_1	1.176000e-03	2.428129e-02	2.068511	<NA>
## 749	MCIDAS	2.621500e-04	7.054838e-03	4.685459	345643
## 750	MCM2	3.173350e-04	8.208611e-03	2.213711	4171
## 751	MCOLN3	4.002930e-05	1.488488e-03	2.586861	55283
## 752	MDK	5.418603e-07	3.902150e-05	2.902095	4192
## 753	MECOM-AS1	2.285513e-07	1.859723e-05	-5.812803	105374205
## 754	MEGF10	2.302263e-03	4.184976e-02	3.789461	84466
## 755	MELTF	1.327226e-04	4.069896e-03	2.761210	4241
## 756	MELTF-AS1	3.535412e-07	2.706292e-05	3.025152	100507057
## 757	MEST	5.414268e-06	2.847381e-04	3.083017	4232
## 758	METTL21A	1.482546e-03	2.937801e-02	2.175690	151194
## 759	MFAP2	1.282661e-07	1.157644e-05	3.881065	4237
## 760	MFAP2_1	3.337140e-07	2.564015e-05	3.744666	<NA>
## 761	MGLL	2.031651e-04	5.720731e-03	-2.674648	11343
## 762	MINDY1	8.245205e-06	4.028650e-04	-2.369815	55793
## 763	MIR9-3HG	3.761687e-04	9.492862e-03	2.841015	254559
## 764	MISP	1.181043e-15	7.628058e-13	4.820197	126353
## 765	MLLT11	1.893854e-04	5.391485e-03	2.860576	10962
## 766	MME	4.519750e-06	2.445398e-04	4.577817	4311
## 767	MMP1	1.952686e-13	8.586834e-11	10.024847	4312

## 768	MMP10	7.422020e-05	2.486196e-03	6.445193	4319
## 769	MMP11	2.589043e-25	8.918389e-22	6.968003	4320
## 770	MMP11_1	2.589043e-25	8.918389e-22	6.968003	<NA>
## 771	MMP12	8.828905e-16	6.082527e-13	5.709569	4321
## 772	MMP13	7.687597e-12	2.147125e-09	9.204225	4322
## 773	MMP3	7.755367e-24	2.289827e-20	10.391534	4314
## 774	MND1	2.165218e-08	2.486151e-06	3.052369	84057
## 775	MPZL3	4.097288e-05	1.513426e-03	-2.403439	196264
## 776	MS4A4E	5.788984e-04	1.339829e-02	2.561356	643680
## 777	MSANTD3	2.450073e-05	9.794607e-04	2.156940	91283
## 778	MSC	2.089611e-05	8.586098e-04	3.198001	9242
## 779	MSI1	2.181062e-08	2.490507e-06	3.671964	4440
## 780	MSX2	2.494772e-04	6.777814e-03	2.768051	4488
## 781	MUC21	1.672653e-03	3.212862e-02	-4.933032	394263
## 782	MUC21_1	2.695187e-12	8.440018e-10	-8.795640	<NA>
## 783	MUC21_2	8.019181e-04	1.759453e-02	-5.661034	<NA>
## 784	MUC21_3	2.371758e-12	7.704577e-10	-8.608956	<NA>
## 785	MUC22	1.629013e-08	1.932439e-06	-9.896729	100507679
## 786	MUC22_1	1.081617e-08	1.371463e-06	-7.786738	<NA>
## 787	MUC22_2	6.698438e-04	1.504819e-02	-7.605726	<NA>
## 788	MUC22_7	8.174218e-06	4.019718e-04	-5.754626	<NA>
## 789	MUC5B	3.347267e-05	1.267057e-03	-6.817035	727897
## 790	MUCL1	1.720775e-04	4.995081e-03	5.139291	118430
## 791	MUCL3	2.811209e-14	1.529002e-11	-8.114777	135656
## 792	MUCL3_1	2.808169e-03	4.910256e-02	-6.814873	<NA>
## 793	MUCL3_2	1.115197e-04	3.504450e-03	-5.878115	<NA>
## 794	MUSTN1	1.370208e-04	4.183082e-03	-2.927784	389125
## 795	MYBL2	4.303610e-08	4.515077e-06	2.325377	4605
## 796	MYH11	4.993701e-07	3.634148e-05	-3.755102	4629
## 797	MYH11_1	4.993701e-07	3.634148e-05	-3.755102	<NA>
## 798	MYL9	2.966151e-05	1.141609e-03	-2.933443	10398
## 799	MYOC	1.923254e-19	3.312485e-16	-9.146094	4653
## 800	MYOCD	7.212244e-06	3.653497e-04	-3.837419	93649
## 801	MYOZ1	1.012600e-04	3.224717e-03	-4.694806	58529
## 802	MYRIP	4.682507e-04	1.129266e-02	-2.687636	25924
## 803	MYZAP	2.844590e-07	2.252566e-05	-3.899927	100820829
## 804	N4BP3	1.594870e-04	4.722460e-03	-2.265274	23138
## 805	NAT14	1.391425e-04	4.222904e-03	2.204311	57106
## 806	NAT8L	2.228290e-03	4.071998e-02	2.926759	339983
## 807	NCCRP1	9.035936e-04	1.957597e-02	-3.152976	342897
## 808	NCK1-DT	4.154601e-05	1.530611e-03	3.020302	101927597
## 809	NDC80	7.263707e-04	1.612527e-02	2.354666	10403
## 810	NDRG2	2.593479e-03	4.612911e-02	-1.874932	57447
## 811	NEB	1.880655e-05	7.831376e-04	3.423178	4703
## 812	NECAB2	1.494990e-03	2.956789e-02	4.087766	54550
## 813	NEK10	2.229698e-04	6.169130e-03	-2.618473	152110
## 814	NEK2	2.126430e-04	5.955157e-03	2.404138	4751
## 815	NELL2	8.094926e-08	7.854739e-06	3.520369	4753
## 816	NEURL3	8.603913e-04	1.871849e-02	3.316015	93082
## 817	NEXN	1.866758e-04	5.329028e-03	-2.450713	91624
## 818	NGB	1.430257e-04	4.302846e-03	5.573539	58157
## 819	NIBAN2	6.040170e-10	1.031721e-07	-2.746913	64855
## 820	NICOL1	1.971434e-05	8.149121e-04	2.851845	401115
## 821	NIPAL1	6.348598e-11	1.395881e-08	-3.120640	152519

## 822	NKAIN2	1.159201e-03	2.400639e-02	4.266265	154215
## 823	NKX2-1	3.365263e-04	8.653582e-03	5.715618	7080
## 824	NLRX1	1.342092e-06	8.534882e-05	-2.671439	79671
## 825	NMB	1.842989e-07	1.587121e-05	4.095699	4828
## 826	NMB_1	1.842989e-07	1.587121e-05	4.095699	<NA>
## 827	NMRAL2P	5.560941e-07	3.918363e-05	4.624477	344887
## 828	NOS2	1.408627e-04	4.256241e-03	4.376732	4843
## 829	NOTUM	3.435812e-04	8.788536e-03	3.544654	147111
## 830	NPAS2	9.128900e-05	2.968883e-03	-2.364852	4862
## 831	NPBWR1	8.613938e-07	5.780288e-05	5.649350	2831
## 832	NPY1R	4.092097e-04	1.015312e-02	-3.587613	4886
## 833	NROB1	4.314303e-05	1.577144e-03	6.295440	190
## 834	NRCAM	4.175129e-04	1.028505e-02	3.117400	4897
## 835	NREP	1.537652e-03	3.026637e-02	2.361352	9315
## 836	NRIP3	2.144539e-05	8.742275e-04	2.404298	56675
## 837	NT5C2	5.066368e-04	1.195339e-02	-1.784537	22978
## 838	NTRK3	1.684989e-05	7.150995e-04	-3.142386	4916
## 839	NTS	5.161906e-05	1.833097e-03	6.147847	4922
## 840	NUF2	4.586875e-07	3.373720e-05	2.885278	83540
## 841	NXPH4	1.720730e-04	4.995081e-03	2.976568	11247
## 842	OCLN_1	2.260104e-03	4.119209e-02	-2.585421	<NA>
## 843	ODC1	3.981668e-04	9.938781e-03	3.023055	4953
## 844	OGDHL	1.547121e-05	6.689520e-04	4.916908	55753
## 845	OGN	2.495605e-04	6.777814e-03	-3.615193	4969
## 846	OGN_1	2.495605e-04	6.777814e-03	-3.615193	<NA>
## 847	ONECUT2	1.824649e-05	7.633976e-04	4.524312	9480
## 848	OPRK1	2.585216e-05	1.021630e-03	7.181649	4986
## 849	ORC6	1.391573e-05	6.180592e-04	2.499158	23594
## 850	OTOP3	1.098274e-06	7.093479e-05	-3.629059	347741
## 851	OTX1	2.071860e-03	3.842563e-02	2.573373	5013
## 852	P2RX1	2.377193e-06	1.399767e-04	-3.418050	5023
## 853	P2RX2	2.085307e-09	3.145921e-07	-6.521712	22953
## 854	P2RY14	5.550078e-07	3.918363e-05	-3.416390	9934
## 855	P3H4	2.048287e-05	8.433065e-04	2.658014	10609
## 856	P4HA3	2.124189e-03	3.930416e-02	2.957375	283208
## 857	PABIR3	4.882167e-05	1.743269e-03	2.799606	159091
## 858	PADI1	9.150274e-05	2.968883e-03	-3.890382	29943
## 859	PADI1_1	9.150274e-05	2.968883e-03	-3.890382	<NA>
## 860	PAQR5	2.557773e-04	6.919379e-03	-2.510576	54852
## 861	PAX3	2.635461e-03	4.671501e-02	6.215198	5077
## 862	PAX6	1.636234e-08	1.932439e-06	6.834757	5080
## 863	PAX7	4.539370e-10	8.018778e-08	8.914945	5081
## 864	PAX9	3.969223e-07	2.961585e-05	-2.947128	5083
## 865	PBK	9.363061e-04	2.007425e-02	2.290988	55872
## 866	PBX2_2	1.437124e-12	4.869260e-10	22.065777	<NA>
## 867	PCAT7	2.614539e-10	4.957551e-08	5.440406	101928099
## 868	PCDH1	4.295060e-05	1.573941e-03	-2.501683	5097
## 869	PCDHB9_1	9.832457e-04	2.084279e-02	2.413797	<NA>
## 870	PCLAF	2.183426e-05	8.883277e-04	2.692258	9768
## 871	PCP4	1.060519e-12	3.779105e-10	-5.700745	5121
## 872	PCP4_1	1.060519e-12	3.779105e-10	-5.700745	<NA>
## 873	PCSK1N	3.808541e-06	2.104677e-04	3.750836	27344
## 874	PCSK2	1.539532e-03	3.026637e-02	-3.033709	5126
## 875	PDE11A	2.829155e-03	4.930268e-02	4.633646	50940

## 876	PDGFA	4.874722e-04	1.163404e-02	1.875907	5154
## 877	PDK4	3.987510e-12	1.177341e-09	-4.453273	5166
## 878	PDLIM2	7.854303e-07	5.339893e-05	-2.656734	64236
## 879	PDZRN3	8.008219e-04	1.758915e-02	-2.401064	23024
## 880	PDZRN4	4.744434e-09	6.409017e-07	-4.964039	29951
## 881	PFN2	7.966882e-08	7.803769e-06	3.263089	5217
## 882	PGM2L1	6.686159e-06	3.429021e-04	2.712992	283209
## 883	PGM5	4.475768e-17	4.405008e-14	-4.915019	5239
## 884	PGM5-AS1	1.357958e-11	3.598242e-09	-6.574068	572558
## 885	PGM5P4	3.151616e-05	1.201801e-03	-5.638823	729468
## 886	PHACTR2	4.100632e-05	1.513426e-03	-2.293035	9749
## 887	PHACTR4	2.700326e-04	7.219965e-03	-1.856987	65979
## 888	PHLDB3	2.186043e-03	4.017200e-02	-2.147347	653583
## 889	PHYHIP	2.911076e-09	4.207421e-07	-3.863414	9796
## 890	PI16	1.409410e-05	6.211019e-04	-4.560170	221476
## 891	PIMREG	3.098055e-08	3.424097e-06	2.547777	54478
## 892	PITX1	2.381226e-03	4.294518e-02	-2.245791	5307
## 893	PKD1L1	3.767618e-04	9.496235e-03	2.553979	168507
## 894	PKHD1L1	5.106729e-05	1.816624e-03	-3.380799	93035
## 895	PKP3	1.991059e-04	5.629441e-03	-2.008591	11187
## 896	PLA2G2A	4.277103e-04	1.051326e-02	-4.948860	5320
## 897	PLA2G2C	2.983984e-06	1.685117e-04	-4.638091	391013
## 898	PLA2G4B	1.763001e-04	5.096182e-03	-2.261183	100137049
## 899	PLAAC2	1.835967e-04	5.255645e-03	4.522061	54979
## 900	PLAU	8.924327e-08	8.539259e-06	3.487790	5328
## 901	PLCXD3	2.340611e-03	4.239768e-02	-3.340030	345557
## 902	PLD6	5.885721e-04	1.356951e-02	2.224183	201164
## 903	PLEKHA7	1.745834e-06	1.077101e-04	-2.312360	144100
## 904	PLEKHG4B	9.170681e-05	2.970841e-03	2.862926	153478
## 905	PLEKHM1	3.976752e-09	5.516209e-07	-2.722018	9842
## 906	PLEKHM1_2	7.409799e-08	7.327547e-06	-2.506878	<NA>
## 907	PLIN4	5.420294e-05	1.908944e-03	-2.621748	729359
## 908	PLN	1.271498e-04	3.922287e-03	-2.997009	5350
## 909	PLP1	7.572741e-16	5.589764e-13	-4.829035	5354
## 910	PLPP4	1.949588e-05	8.074965e-04	4.244584	196051
## 911	PMM1	1.445984e-03	2.884709e-02	-1.981647	5372
## 912	POMC	1.243270e-03	2.530811e-02	2.856794	5443
## 913	POPDC3	2.105389e-07	1.747558e-05	5.156923	64208
## 914	POSTN	2.072991e-03	3.842563e-02	2.994432	10631
## 915	POTEF_1	4.507484e-05	1.631535e-03	7.272708	<NA>
## 916	POU6F2-AS2	5.564026e-05	1.942742e-03	5.163328	100689074
## 917	PPL	2.151941e-08	2.484711e-06	-3.450053	5493
## 918	PPP1R12B	3.784147e-06	2.096803e-04	-2.239788	4660
## 919	PPP1R1A	2.827731e-05	1.096502e-03	-3.787235	5502
## 920	PPP1R1C	2.984095e-06	1.685117e-04	4.671039	151242
## 921	PRAME	3.203421e-19	4.729165e-16	9.717650	23532
## 922	PRAME_1	3.203421e-19	4.729165e-16	9.717650	<NA>
## 923	PRDM13	2.433312e-04	6.652339e-03	5.158176	59336
## 924	PRELID3A	3.111817e-05	1.188818e-03	2.684926	10650
## 925	PRELP	1.672549e-03	3.212862e-02	-2.698887	5549
## 926	PROC	1.663064e-05	7.087052e-04	5.211048	5624
## 927	PRR15	1.308674e-05	5.905608e-04	4.052911	222171
## 928	PRR15-DT	2.206062e-03	4.038519e-02	3.373034	107986700
## 929	PRR20G	2.673459e-03	4.734795e-02	6.387081	100419008

## 930	PRR4	3.540414e-08	3.771819e-06	-7.109019	11272
## 931	PRR4_1	3.540414e-08	3.771819e-06	-7.109019	<NA>
## 932	PRR4_2	3.540414e-08	3.771819e-06	-7.109019	<NA>
## 933	PRR5L	1.551217e-03	3.038915e-02	2.251606	79899
## 934	PRSS21	2.440401e-06	1.432904e-04	4.829719	10942
## 935	PRSS27	6.003224e-04	1.375550e-02	-3.582000	83886
## 936	PRSS3	1.241568e-03	2.530644e-02	-2.889999	5646
## 937	PRUNE2	2.569293e-09	3.803037e-07	-3.681802	158471
## 938	PSCA	5.923937e-04	1.361271e-02	-3.259137	8000
## 939	PTGIS	2.999263e-04	7.816995e-03	-3.399417	5740
## 940	PTGS2	6.091846e-05	2.108983e-03	4.037431	5743
## 941	PTH2R	5.982131e-05	2.077961e-03	6.219175	5746
## 942	PTHLH	6.864420e-09	8.922882e-07	3.980194	5744
## 943	PTK6	1.754722e-04	5.079354e-03	-2.678348	5753
## 944	PTPRN	2.374273e-03	4.285718e-02	4.559153	5798
## 945	PWWP3B	1.800291e-04	5.182231e-03	6.016714	139221
## 946	PXDN	1.540556e-03	3.026637e-02	2.418928	7837
## 947	PYGM	1.744267e-06	1.077101e-04	-3.147673	5837
## 948	PYY2	1.178449e-07	1.082498e-05	6.875747	23615
## 949	RAB15	1.014687e-03	2.133196e-02	2.093009	376267
## 950	RAB25	1.599242e-03	3.100669e-02	-1.794638	57111
## 951	RAB32	4.940038e-04	1.176275e-02	2.280534	10981
## 952	RAB3B	6.643433e-04	1.494086e-02	3.337368	5865
## 953	RAB40A	1.427837e-16	1.229606e-13	-5.250004	142684
## 954	RAB9B	2.807589e-03	4.910256e-02	-2.611853	51209
## 955	RABGGTA	6.763207e-05	2.321960e-03	-2.058172	5875
## 956	RABGGTA_1	6.763207e-05	2.321960e-03	-2.058172	<NA>
## 957	RAD51AP1	1.063589e-07	9.946724e-06	3.231669	10635
## 958	RAET1E	1.244099e-03	2.530811e-02	-3.154733	135250
## 959	RAG1	4.545504e-04	1.102658e-02	3.249189	5896
## 960	RANBP9	3.137407e-04	8.125804e-03	-2.065572	10048
## 961	RARG	8.138982e-04	1.783844e-02	-1.750511	5916
## 962	RASL11B	1.510644e-04	4.511850e-03	3.428543	65997
## 963	RBFOX3	8.175652e-19	1.126496e-15	-5.856158	146713
## 964	RBM47	1.036379e-03	2.168004e-02	-2.114475	54502
## 965	RBP1	9.618717e-05	3.091752e-03	3.148141	5947
## 966	RBP4	5.810675e-04	1.343345e-02	4.120184	5950
## 967	RBPMS2	4.334408e-05	1.579957e-03	-2.317533	348093
## 968	RDH12	1.096810e-09	1.784950e-07	-4.161113	145226
## 969	RELCH	3.830260e-04	9.618934e-03	-1.913041	57614
## 970	RFC4	1.884428e-09	2.863777e-07	2.888049	5984
## 971	RHCG	3.213789e-05	1.221004e-03	-3.397654	51458
## 972	RHEBL1	9.289541e-05	3.004636e-03	3.271270	121268
## 973	RIMS2	6.223337e-07	4.287464e-05	3.488222	9699
## 974	RMND5B	4.439403e-04	1.087311e-02	-1.838266	64777
## 975	RNASEH2A	3.049188e-04	7.937104e-03	1.991206	10535
## 976	RNF222	1.000872e-06	6.531486e-05	-3.856026	643904
## 977	RNF224	9.636083e-04	2.057423e-02	-3.310112	643596
## 978	RNF225	1.362425e-04	4.165473e-03	-3.196677	646862
## 979	RNFT2	2.564937e-05	1.017910e-03	3.181044	84900
## 980	ROR1	1.231616e-03	2.520301e-02	-2.688994	4919
## 981	ROS1	4.186410e-09	5.730114e-07	6.882220	6098
## 982	RPL39L	5.246402e-08	5.394659e-06	3.476519	116832
## 983	RRAD	2.474235e-03	4.423658e-02	-2.920293	6236

## 984	SALL4	2.140600e-13	9.217068e-11	4.122564	57167
## 985	SAMD5	1.810631e-06	1.100651e-04	-3.227946	389432
## 986	SASH1	3.540705e-10	6.533864e-08	-3.172226	23328
## 987	SCARA5	9.760922e-05	3.132589e-03	-3.944008	286133
## 988	SCAT2	1.141086e-03	2.367869e-02	3.885584	112935960
## 989	SCAT8	1.939462e-03	3.650710e-02	3.737606	112935969
## 990	SCEL	6.072728e-06	3.153546e-04	-3.811042	8796
## 991	SCG2	1.131766e-04	3.533435e-03	4.312047	7857
## 992	SCGB1D2	1.779927e-03	3.388698e-02	-4.681143	10647
## 993	SCGB2A2	1.775094e-06	1.089309e-04	-8.033597	4250
## 994	SCGB3A1	1.412706e-04	4.256241e-03	-5.443222	92304
## 995	SCIRT	2.835134e-03	4.936524e-02	3.159560	101929705
## 996	SCN5A	1.711758e-03	3.278833e-02	3.380014	6331
## 997	SCN7A	5.855129e-17	5.500627e-14	-5.361083	6332
## 998	SCN8A	9.791385e-05	3.132637e-03	2.934254	6334
## 999	SCN9A	1.398237e-05	6.180592e-04	4.360995	6335
## 1000	SCNN1B	2.915825e-08	3.240014e-06	-3.301441	6338
## 1001	SCUBE3	4.978555e-04	1.180950e-02	3.427316	222663
## 1002	SCX	2.904901e-04	7.648215e-03	2.411566	642658
## 1003	SEC14L4	5.436615e-05	1.910952e-03	3.646183	284904
## 1004	SERPINB1	1.608432e-03	3.109735e-02	-2.502206	1992
## 1005	SERPINB11	7.353050e-04	1.630610e-02	-2.673095	89778
## 1006	SERPINB12	1.580054e-04	4.687819e-03	-3.426843	89777
## 1007	SERPINB2	6.979814e-04	1.562934e-02	-3.454279	5055
## 1008	SERPINE1	7.981042e-05	2.656235e-03	3.139149	5054
## 1009	SERPINH1	4.421737e-10	7.878316e-08	2.972488	871
## 1010	SEZ6L2	5.889225e-04	1.356951e-02	2.929850	26470
## 1011	SFTA2_2	1.607418e-04	4.746016e-03	-5.290253	<NA>
## 1012	SGCA	1.577508e-04	4.687819e-03	-2.942742	6442
## 1013	SH3BGRL2	8.716296e-07	5.811239e-05	-3.267472	83699
## 1014	SHH	1.027320e-03	2.153538e-02	5.190398	6469
## 1015	SHISA2	1.230592e-04	3.807466e-03	4.512349	387914
## 1016	SHOX2	8.540712e-09	1.103247e-06	7.259799	6474
## 1017	SIM2	5.726257e-04	1.328286e-02	-2.665900	6493
## 1018	SIX1	4.069584e-20	9.345575e-17	6.588428	6495
## 1019	SIX2	9.305482e-10	1.563624e-07	3.881640	10736
## 1020	SIX3	2.556274e-04	6.919379e-03	5.670471	6496
## 1021	SIX4	1.177853e-05	5.421798e-04	3.375702	51804
## 1022	SLC13A4	2.697109e-07	2.150220e-05	-3.522330	26266
## 1023	SLC16A10	1.444491e-04	4.339352e-03	2.606154	117247
## 1024	SLC16A6	4.229268e-06	2.300277e-04	-2.873453	9120
## 1025	SLC24A2	7.862712e-05	2.621073e-03	4.592379	25769
## 1026	SLC27A4	2.566104e-04	6.926999e-03	-2.011666	10999
## 1027	SLC27A6	4.039618e-05	1.496251e-03	-3.382189	28965
## 1028	SLC29A4	2.610296e-03	4.634846e-02	2.548636	222962
## 1029	SLC2A4	4.732973e-08	4.915632e-06	-3.884614	6517
## 1030	SLC2A4_1	4.732973e-08	4.915632e-06	-3.884614	<NA>
## 1031	SLC30A3	7.112559e-06	3.611852e-04	4.767068	7781
## 1032	SLC34A3	1.671779e-04	4.908001e-03	-3.556438	142680
## 1033	SLC35C1	1.509611e-04	4.511850e-03	-2.074358	55343
## 1034	SLC46A2	2.098118e-08	2.436174e-06	-4.700783	57864
## 1035	SLC51A	4.978023e-04	1.180950e-02	4.201332	200931
## 1036	SLC52A1	2.030464e-04	5.720731e-03	3.394591	55065
## 1037	SLC5A12	7.059891e-06	3.593937e-04	4.947956	159963

## 1038	SLC6A1	1.237002e-03	2.526320e-02	-3.261629	6529
## 1039	SLC6A2	1.148041e-04	3.573450e-03	4.766898	6530
## 1040	SLC7A5P2_1	2.517320e-03	4.489040e-02	3.406795	<NA>
## 1041	SLC8A1-AS1	3.178601e-08	3.494432e-06	-6.547623	100128590
## 1042	SLCO1A2	2.316120e-13	9.769302e-11	7.969568	6579
## 1043	SLCO4A1-AS2	1.585798e-06	9.931900e-05	-3.369559	101928465
## 1044	SLITRK3	3.650585e-06	2.028234e-04	-4.058491	22865
## 1045	SLMAP	1.291355e-03	2.619209e-02	-1.593220	7871
## 1046	SLURP1	1.030992e-07	9.729929e-06	-4.450451	57152
## 1047	SMAGP	2.163190e-06	1.288439e-04	-2.251907	57228
## 1048	SMC1B	1.770828e-04	5.111659e-03	5.405907	27127
## 1049	SMOC1	3.366306e-04	8.653582e-03	3.624976	64093
## 1050	SNCAIP	2.277406e-03	4.147086e-02	2.417071	9627
## 1051	SNCB	9.110349e-05	2.968883e-03	5.623633	6620
## 1052	SNHG33	2.767984e-03	4.860552e-02	2.623647	100505687
## 1053	SNORD3A	1.665177e-07	1.470764e-05	7.666877	780851
## 1054	SNX10	2.704941e-07	2.150220e-05	2.789280	29887
## 1055	SORBS1	1.375244e-10	2.805876e-08	-3.447275	10580
## 1056	SORCS1	9.595141e-04	2.050800e-02	-3.180708	114815
## 1057	SORT1	1.545014e-03	3.032511e-02	-1.890219	6272
## 1058	SOST	6.534529e-08	6.642306e-06	7.517938	50964
## 1059	SOX10	9.061043e-06	4.355201e-04	-3.422805	6663
## 1060	SOX11	2.826034e-03	4.928985e-02	3.967565	6664
## 1061	SOX4	1.791994e-03	3.407262e-02	2.103704	6659
## 1062	SPC24	1.029815e-03	2.156455e-02	2.281515	147841
## 1063	SPC25_1	9.308849e-04	2.004118e-02	2.859116	<NA>
## 1064	SPINDOC	9.993194e-04	2.112938e-02	1.723117	144097
## 1065	SPINK5	1.776162e-06	1.089309e-04	-4.404171	11005
## 1066	SPNS2	1.803677e-07	1.572928e-05	-3.694177	124976
## 1067	SPOCD1	8.188026e-06	4.019718e-04	3.833984	90853
## 1068	SPP1	5.752457e-14	2.830757e-11	6.931695	6696
## 1069	SPRR2G	1.005618e-03	2.120828e-02	3.929969	6706
## 1070	SPRR3	1.249474e-05	5.700689e-04	-4.119999	6707
## 1071	SPSB4	4.143043e-04	1.023040e-02	5.065519	92369
## 1072	SPTLC3	2.384871e-03	4.297342e-02	-2.145970	55304
## 1073	SRL	9.786618e-04	2.077410e-02	-2.819274	6345
## 1074	ST6GALNAC1	7.400246e-04	1.634178e-02	-2.446651	55808
## 1075	ST8SIA2	4.536053e-04	1.101954e-02	4.729849	8128
## 1076	STIM1	2.743660e-04	7.307471e-03	-1.659554	6786
## 1077	STK40	1.947326e-03	3.658849e-02	-2.033342	83931
## 1078	STMN1	5.364338e-06	2.828320e-04	2.344041	3925
## 1079	STRA6	2.154674e-04	6.009826e-03	3.321162	64220
## 1080	STRA6_1	2.782547e-04	7.390066e-03	3.266690	<NA>
## 1081	STRN	1.827147e-04	5.237654e-03	-1.833324	6801
## 1082	STUM	3.060956e-07	2.396358e-05	-3.342904	375057
## 1083	STX1A	2.432045e-04	6.652339e-03	2.560746	6804
## 1084	SULT2B1	1.194528e-05	5.486333e-04	-3.267047	6820
## 1085	SYCP2	9.168223e-04	1.977963e-02	3.160839	10388
## 1086	SYNDIG1	5.279756e-04	1.240023e-02	3.745334	79953
## 1087	SYNM	9.144651e-11	1.948471e-08	-3.533343	23336
## 1088	SYNPO2	2.933802e-18	3.789739e-15	-4.868168	171024
## 1089	SYNPO2L	3.043494e-07	2.391746e-05	-6.231343	79933
## 1090	SYT1	4.263939e-12	1.223987e-09	4.128886	6857
## 1091	SYT14	1.165198e-08	1.442054e-06	7.234694	255928

## 1092	TACR1	4.883651e-05	1.743269e-03	-2.962693	6869
## 1093	TACR2	6.451959e-05	2.226195e-03	-3.349569	6865
## 1094	TAGLN	1.604899e-03	3.105810e-02	-2.189254	6876
## 1095	TBX15	1.153048e-05	5.343315e-04	3.732520	6913
## 1096	TBX6	3.674554e-07	2.792121e-05	-2.294096	6911
## 1097	TCF15	2.703510e-03	4.783917e-02	3.481067	6939
## 1098	TD02	2.523041e-06	1.468907e-04	5.301138	6999
## 1099	TD02_1	2.523041e-06	1.468907e-04	5.301138	<NA>
## 1100	TEC	2.763829e-03	4.857383e-02	-2.229229	7006
## 1101	TEC	2.763829e-03	4.857383e-02	-2.229229	100124696
## 1102	TENT5B	1.856360e-03	3.523162e-02	-2.562881	115572
## 1103	TESMIN	4.562923e-04	1.105586e-02	3.156128	9633
## 1104	TFAP2B	4.044693e-04	1.007177e-02	-4.246749	7021
## 1105	TGFBI	2.100421e-05	8.613392e-04	3.502259	7045
## 1106	TGFBR3	8.928982e-07	5.933897e-05	-2.745227	7049
## 1107	TGM1	2.202551e-05	8.925948e-04	-3.507608	7051
## 1108	TGM1_1	2.202551e-05	8.925948e-04	-3.507608	<NA>
## 1109	TGM3	5.897911e-09	7.915456e-07	-4.972337	7053
## 1110	TIAM1	1.079937e-04	3.418092e-03	-2.257659	7074
## 1111	TJP1_1	1.359637e-05	6.069327e-04	-2.338450	<NA>
## 1112	TJP1_2	1.359637e-05	6.069327e-04	-2.338450	<NA>
## 1113	TJP3	5.761943e-04	1.335066e-02	-2.373186	27134
## 1114	TK1	2.452839e-03	4.393005e-02	1.961834	7083
## 1115	TM4SF19	1.320071e-03	2.672207e-02	3.008430	116211
## 1116	TMEFF1	1.027380e-03	2.153538e-02	4.546940	8577
## 1117	TMEM132C	1.988571e-11	5.202505e-09	-5.480825	92293
## 1118	TMEM154	6.763835e-08	6.786163e-06	-2.908528	201799
## 1119	TMEM35A	2.691721e-07	2.150220e-05	-3.736828	59353
## 1120	TMEM44-AS1	1.203836e-03	2.473248e-02	2.755132	100507297
## 1121	TMEM79	4.033289e-04	1.005549e-02	-2.554608	84283
## 1122	TMOD3	2.365252e-03	4.276905e-02	-1.752391	29766
## 1123	TMPRSS11B	1.802152e-03	3.423426e-02	-4.210995	132724
## 1124	TNFAIP6	8.886655e-05	2.915387e-03	3.271895	7130
## 1125	TNFRSF18	1.196969e-06	7.659122e-05	3.158992	8784
## 1126	TNXB_7	1.325575e-11	3.558049e-09	-4.581801	<NA>
## 1127	TOLLIP	2.015824e-03	3.756813e-02	-1.658110	54472
## 1128	TOLLIP_1	1.870016e-03	3.545825e-02	-1.663323	<NA>
## 1129	TOM1	2.812058e-04	7.451232e-03	-1.908411	10043
## 1130	TOM1L2	2.438309e-05	9.766468e-04	-2.036659	146691
## 1131	TOP2A	1.434798e-03	2.867931e-02	2.071932	7153
## 1132	TPM2	4.901793e-04	1.168515e-02	-2.361833	7169
## 1133	TPRG1	1.445363e-05	6.328976e-04	-2.529544	285386
## 1134	TREH	2.875734e-03	4.987014e-02	-2.396361	11181
## 1135	TREM1	1.695229e-03	3.253202e-02	2.958320	54210
## 1136	TREM2	8.676284e-07	5.803283e-05	4.014659	54209
## 1137	TRIM29	2.740877e-04	7.307471e-03	-2.157665	23650
## 1138	TRIM46	2.330362e-03	4.228614e-02	2.647856	80128
## 1139	TRIP13	3.112460e-04	8.071308e-03	1.997837	9319
## 1140	TRPM2-AS	2.074563e-06	1.246427e-04	6.184315	101928607
## 1141	TTC39A	9.294553e-06	4.426263e-04	-2.237704	22996
## 1142	TTC9	1.882027e-04	5.365206e-03	-2.565271	23508
## 1143	TTYH3	2.285098e-04	6.305528e-03	2.388186	80727
## 1144	TUBB3	7.645872e-08	7.524994e-06	3.769970	10381
## 1145	UBE2C	1.443628e-08	1.734703e-06	2.222365	11065

## 1146	UBE2T	9.242600e-06	4.411687e-04	3.110640	29089
## 1147	UBL3	9.017130e-06	4.344197e-04	-2.126612	5412
## 1148	UCHL1	1.003493e-11	2.728973e-09	5.371256	7345
## 1149	UGT1A8	1.225521e-12	4.293061e-10	-7.320981	54576
## 1150	UNC13B	2.480866e-03	4.431680e-02	-1.685767	10497
## 1151	UNC5B-AS1	2.803935e-05	1.091369e-03	3.976938	728978
## 1152	UNC93A	1.700693e-04	4.971700e-03	-3.847726	54346
## 1153	UPK1A	4.435137e-09	6.030619e-07	-5.137915	11045
## 1154	UPK3B	2.509729e-04	6.807230e-03	-3.618064	105375355
## 1155	UPK3B_1	2.135985e-06	1.275911e-04	-3.658145	<NA>
## 1156	USP2	9.263793e-04	1.996497e-02	-2.678990	9099
## 1157	USP32P3	2.064825e-03	3.834303e-02	2.513311	347716
## 1158	USP6NL	1.371863e-06	8.697441e-05	-2.298400	9712
## 1159	UTS2R	9.514596e-05	3.067781e-03	3.040311	2837
## 1160	VAX1	7.412510e-05	2.486196e-03	5.398244	11023
## 1161	VGF	6.023291e-04	1.377095e-02	2.908115	7425
## 1162	VIT	3.082910e-07	2.404437e-05	-4.231284	5212
## 1163	VPS37D	1.816291e-05	7.614423e-04	2.890571	155382
## 1164	VPS4B	1.605215e-04	4.746016e-03	-2.031815	9525
## 1165	VSIG10L	1.561286e-10	3.132880e-08	-4.112878	147645
## 1166	VSIG2	1.063380e-04	3.370850e-03	-3.334456	23584
## 1167	WVA5B2	1.326630e-06	8.462587e-05	6.115367	90113
## 1168	WDR54	1.173555e-05	5.421798e-04	2.446509	84058
## 1169	WDR72	3.973141e-10	7.266981e-08	4.036126	256764
## 1170	WNK4	3.207310e-12	9.758363e-10	-5.772009	65266
## 1171	WNT11	4.607905e-10	8.070864e-08	5.019165	7481
## 1172	WNT2	2.528722e-03	4.501953e-02	4.186985	7472
## 1173	XCL1	4.156820e-06	2.266838e-04	3.932025	6375
## 1174	XIRP1	2.044729e-07	1.704051e-05	7.117296	165904
## 1175	XIST	9.408175e-16	6.272521e-13	-24.898870	7503
## 1176	XPNPEP2	2.837938e-04	7.510180e-03	-3.535088	7512
## 1177	ZBTB16	5.470011e-07	3.911909e-05	-2.903755	7704
## 1178	ZDHHC15	4.850272e-05	1.737356e-03	-2.477272	158866
## 1179	ZDHHC21	1.449244e-03	2.888425e-02	-1.877833	340481
## 1180	ZIC1	2.511976e-03	4.483377e-02	5.559280	7545
## 1181	ZIC2	2.272580e-06	1.349704e-04	4.318575	7546
## 1182	ZIC5	1.713194e-09	2.657141e-07	7.867597	85416
## 1183	ZNF114	5.551843e-05	1.942742e-03	3.289053	163071
## 1184	ZNF185	3.989142e-06	2.192755e-04	-3.092249	7739
## 1185	ZNF185_1	1.828335e-08	2.134917e-06	-3.513000	<NA>
## 1186	ZNF365	7.262496e-05	2.463529e-03	-3.589701	22891
## 1187	ZNF431	1.966448e-03	3.684729e-02	-2.045058	170959
## 1188	ZNF469	4.976252e-06	2.650752e-04	3.830037	84627
## 1189	ZNF536	3.270465e-04	8.438698e-03	-3.547536	9745
## 1190	ZNF584-DT	4.081939e-04	1.014400e-02	2.528976	123706526
## 1191	ZNF677	1.159294e-04	3.603050e-03	-2.738006	342926
## 1192	ZNF750	2.602564e-03	4.625090e-02	-2.333524	79755
## 1193	ZNF823	2.401370e-03	4.323302e-02	-1.767326	55552

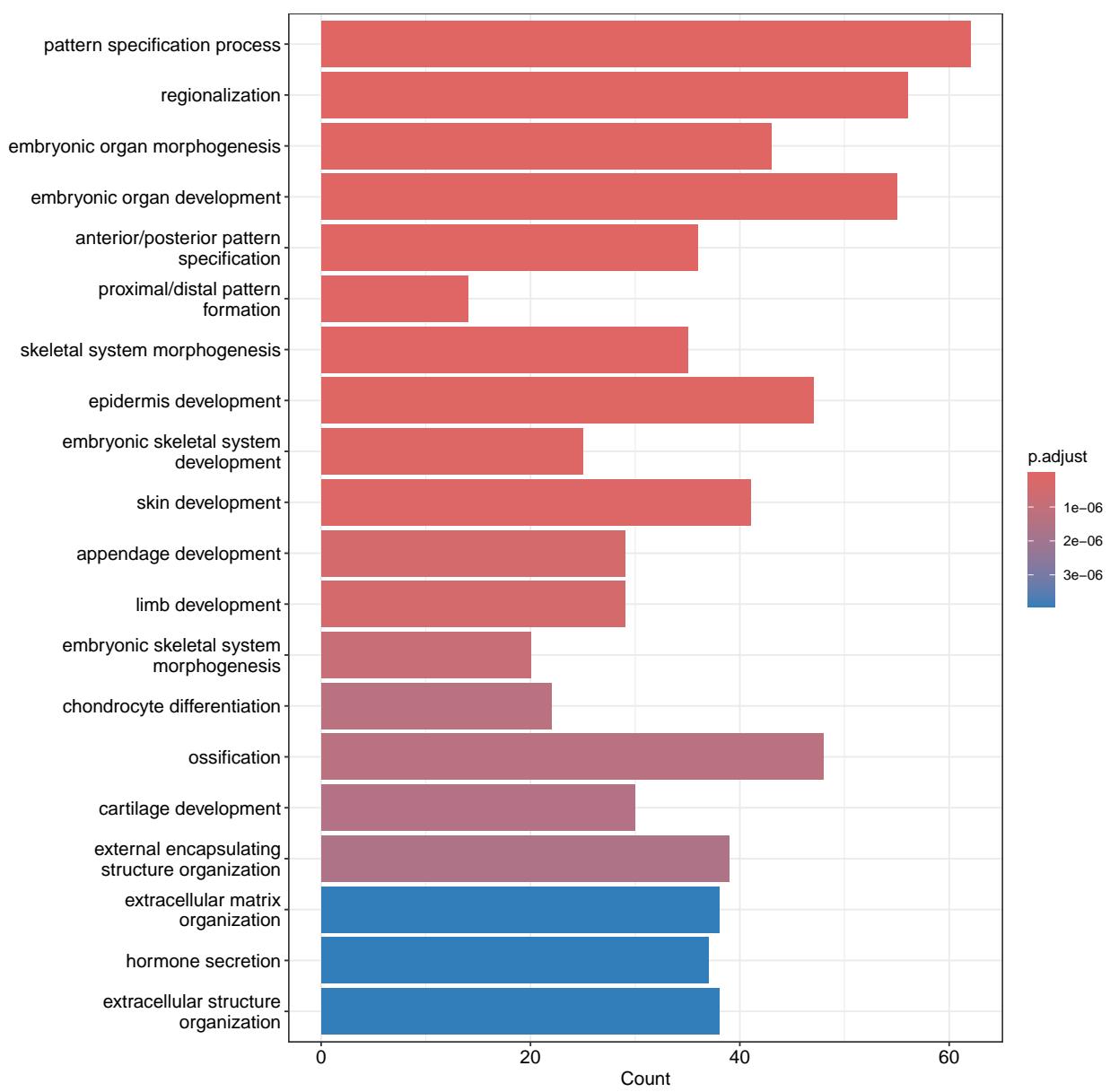
Let's take a look at some of the biological processes, molecular functions and cellular components that are enriched in this DEG profile.

```
# GO enrichment
library(clusterProfiler)
library(org.Hs.eg.db)

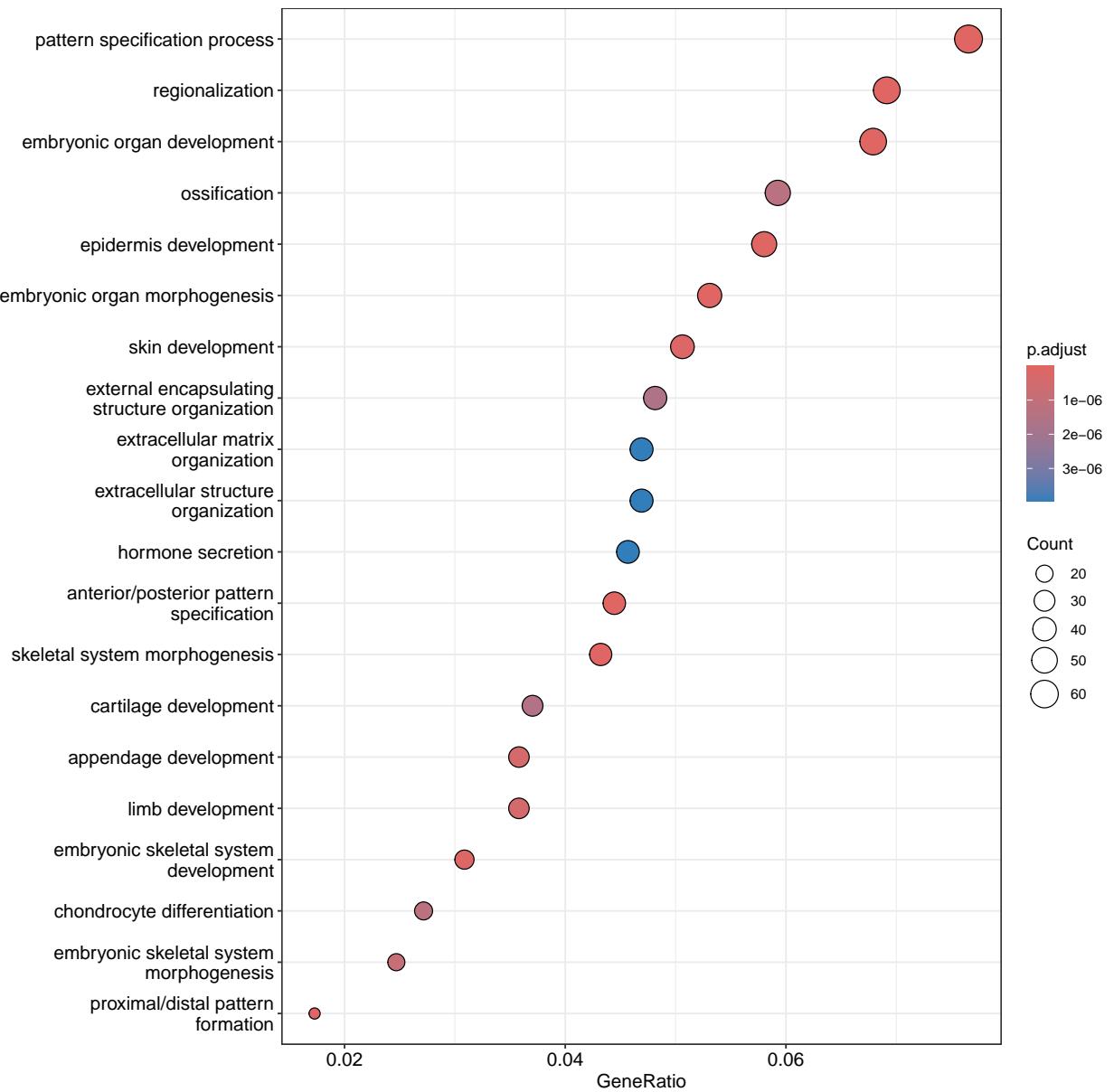
# For BP (Biological Process)
rs_ego <- enrichGO(rownames(res_lfc_padj_0.05), OrgDb = "org.Hs.eg.db", keyType = "SYMBOL",
                     ont ="BP", pAdjustMethod = "BH", pvalueCutoff = 0.05)
head(rs_ego)

##                                     ID          Description GeneRatio
## GO:0007389 GO:0007389      pattern specification process 62/810
## GO:0003002 GO:0003002      regionalization        56/810
## GO:0048562 GO:0048562      embryonic organ morphogenesis 43/810
## GO:0048568 GO:0048568      embryonic organ development 55/810
## GO:0009952 GO:0009952 anterior/posterior pattern specification 36/810
## GO:0009954 GO:0009954      proximal/distal pattern formation 14/810
##           BgRatio      pvalue     p.adjust      qvalue
## GO:0007389 475/18870 4.549951e-15 2.177151e-11 1.836743e-11
## GO:0003002 430/18870 1.114470e-13 2.666369e-10 2.249469e-10
## GO:0048562 295/18870 1.977370e-12 3.153905e-09 2.660777e-09
## GO:0048568 453/18870 3.274792e-12 3.505226e-09 2.957167e-09
## GO:0009952 219/18870 3.662723e-12 3.505226e-09 2.957167e-09
## GO:0009954 34/18870 4.036119e-11 3.218805e-08 2.715529e-08
##
## GO:0007389 AURKA/BARX1/BMPR1B/CDX2/CFAP53/CHRD1/CITED2/DAAM2/DKK1/DLL3/DLX1/DNAH5/DOP1B/EN1/FE
## GO:0003002
## GO:0048562
## GO:0048568
## GO:0009952
## GO:0009954
##           Count
## GO:0007389    62
## GO:0003002    56
## GO:0048562    43
## GO:0048568    55
## GO:0009952    36
## GO:0009954    14

barplot(rs_ego, showCategory = 20)
```



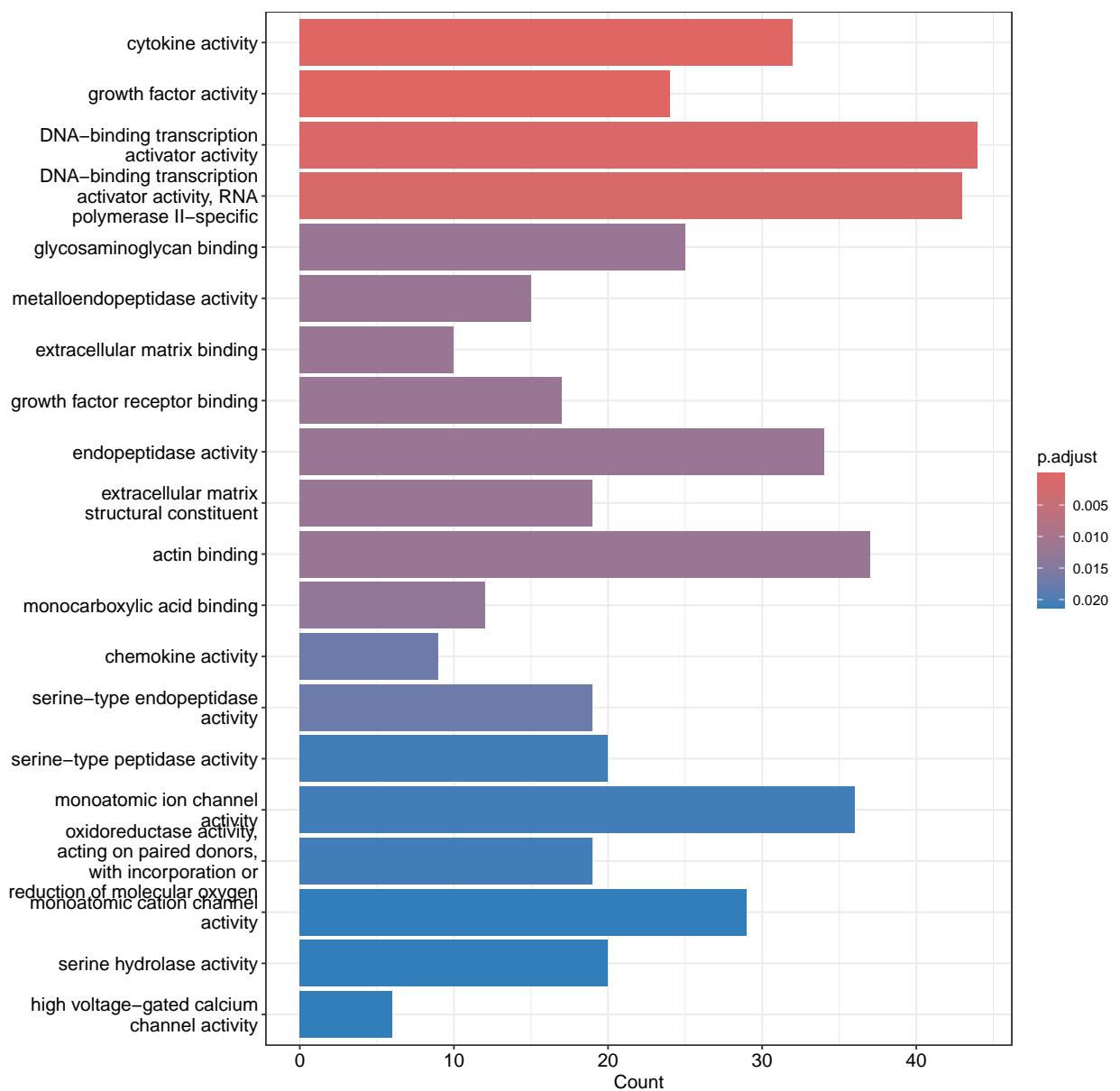
```
dotplot(rs_ego, showCategory = 20)
```



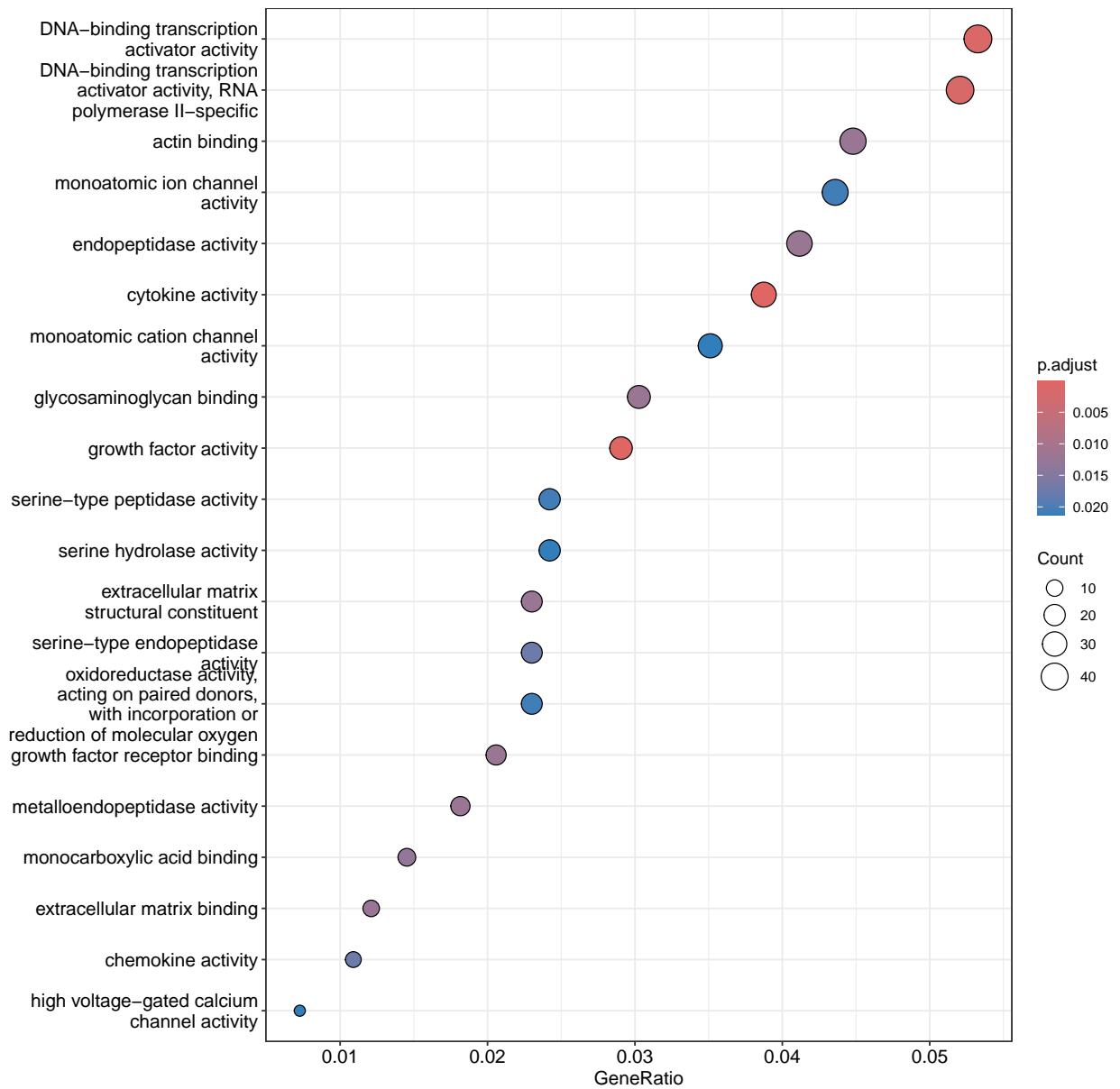
```
# For MF (molecular function)

mf_ego <- enrichGO(rownames(res_lfc_padj_0.05), OrgDb = "org.Hs.eg.db", keyType = "SYMBOL",
                     ont = "MF", pAdjustMethod = "BH", pvalueCutoff = 0.05)

barplot(mf_ego, showCategory = 20)
```

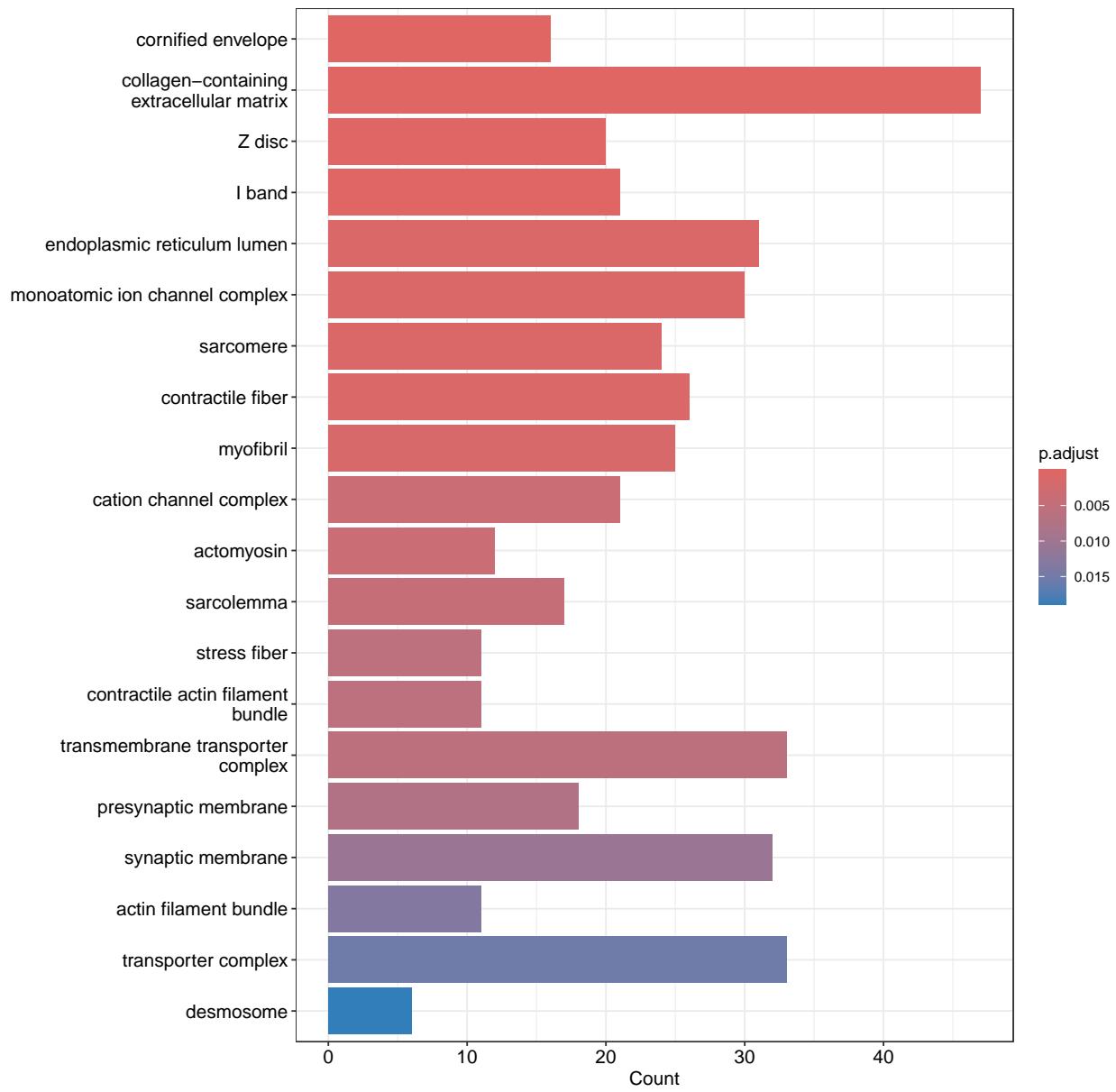


```
dotplot(mf_ego, showCategory = 20)
```

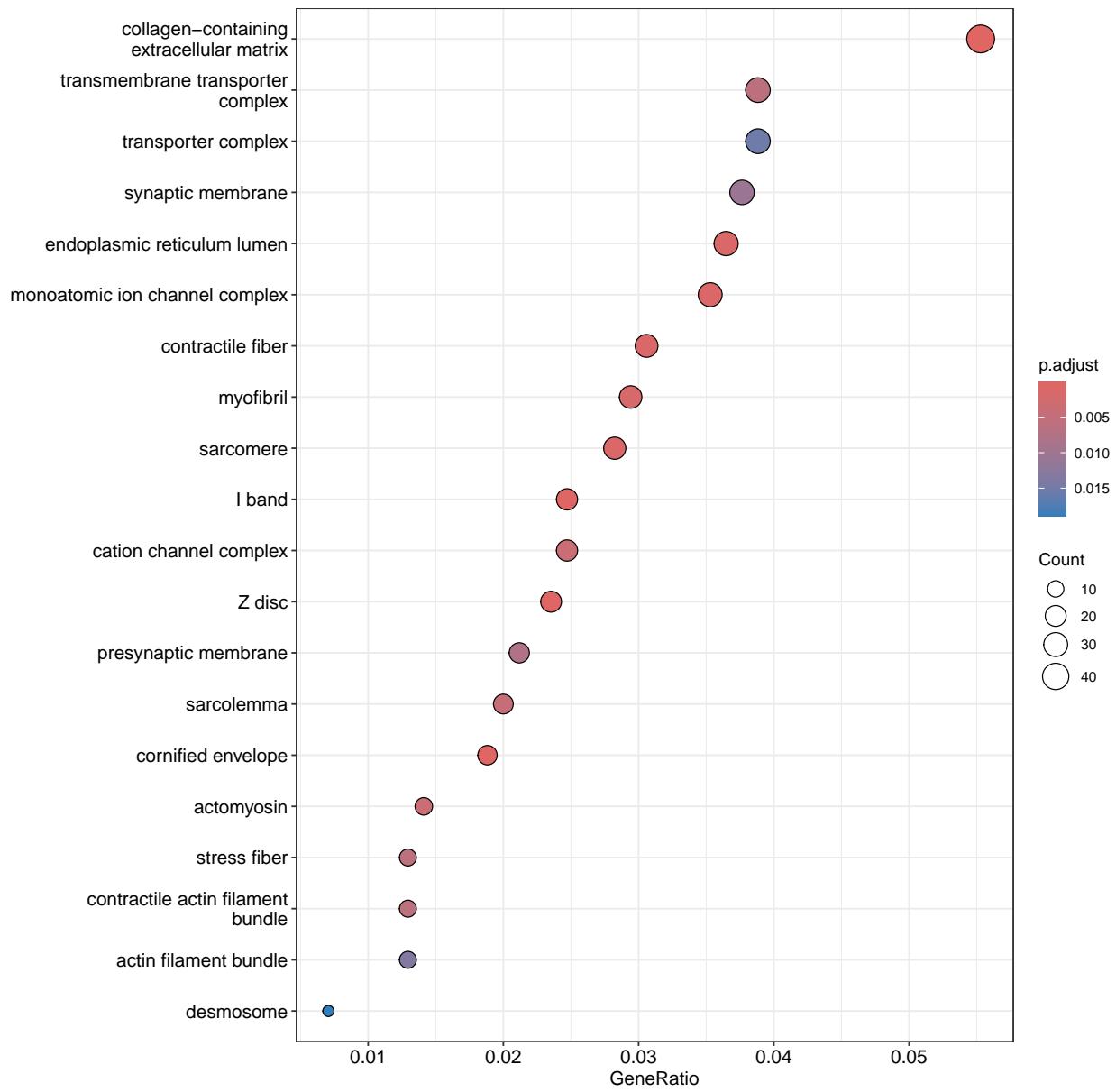


```
# For CC (cellular component)
cc_ego <- enrichGO(rownames(res_lfc_padj_0.05), OrgDb = "org.Hs.eg.db", keyType = "SYMBOL",
                     ont = "CC", pAdjustMethod = "BH", pvalueCutoff = 0.05)

barplot(cc_ego, showCategory = 20)
```



```
dotplot(cc_ego, showCategory = 20)
```



deg\_list

```
## [1] "A2ML1"          "ABCA13"
## [3] "ABCC2"          "ABCG4"
## [5] "ABLIM1"          "ABLIM3"
## [7] "ACER1"           "ACKR1"
## [9] "ACKR3"           "ACOT11"
## [11] "ACOX1"          "ACOX2"
## [13] "ACOX3"          "ACP3"
## [15] "ACTG2"           "ACVR1C"
## [17] "ADAM12"          "ADAM23"
## [19] "ADAM32_1"        "ADAM33"
## [21] "ADAMTS12"        "ADAMTS14"
```

```

## [23] "ADAMTS7"          "ADAMTS8"
## [25] "ADAMTS9-AS1"       "ADAMTS9-AS2"
## [27] "ADCY5"             "ADD2"
## [29] "ADGRF2P"           "ADGRG6"
## [31] "ADH1B"              "AFDN"
## [33] "AFF3"               "AGFG2"
## [35] "AGTR1"              "AIF1L"
## [37] "AKR1B1"             "AKR1C1"
## [39] "AKR1C2"             "AKR1C3"
## [41] "ALDH9A1"            "ALG1L1P"
## [43] "ALOX12"              "ALOX12P2"
## [45] "ALOX15"              "ALX1"
## [47] "AMOTL2"              "ANGPTL1"
## [49] "ANKRD19P"            "ANKRD20A11P"
## [51] "ANKRD20A5P"          "ANKRD20A9P"
## [53] "ANKRD20A9P_1"        "ANKRD22"
## [55] "ANKRD34B"            "ANO1"
## [57] "ANXA10"              "ANXA9"
## [59] "AO3"                 "AOX1"
## [61] "APLN"                "APOC1"
## [63] "AQP5"                "AQP7P1"
## [65] "ARHGAP10"            "ARHGAP27"
## [67] "ARHGAP27_2"          "ARHGEF10L"
## [69] "ARSF"                 "ARTN"
## [71] "ASB5"                 "ASCL2"
## [73] "ASPA"                 "ASPG"
## [75] "ASPN_1"               "ATG9B"
## [77] "ATOH8"                "ATP13A4"
## [79] "ATP1A2"               "AUNIP"
## [81] "AURKA"                "AURKB"
## [83] "B3GNT8"               "B4GALNT1"
## [85] "B4GALNT4"              "BAMBI"
## [87] "BARX1"                 "BARX2"
## [89] "BBOX1-AS1"             "BCAN"
## [91] "BCAS1"                 "BICD1"
## [93] "BIRC5"                 "BLNK"
## [95] "BMP8A"                 "BMPR1B"
## [97] "BMS1P14"               "BMS1P23"
## [99] "BNIPL"                 "BOC"
## [101] "BPIFB2"               "C15orf40"
## [103] "C15orf62"              "C16orf74"
## [105] "C1orf116"              "C1QTNF6"
## [107] "C3orf70"               "C6orf132"
## [109] "C6orf15"               "C6orf15_2"
## [111] "C6orf15_3"              "C6orf15_4"
## [113] "C6orf15_5"              "C7"
## [115] "C7orf57"               "CA13"
## [117] "CA9"                   "CAB39L"
## [119] "CABYR"                  "CACNA1B"
## [121] "CACNB2"                  "CACNG4"
## [123] "CADM3"                  "CALB1"
## [125] "CAMK2N2"                  "CAPN14"
## [127] "CASC9"                  "CASQ2"
## [129] "CBS"                     "CCDC162P"

```

```

## [131] "CCDC187"          "CCL14"
## [133] "CCL14_1"           "CCL20"
## [135] "CCN4"                "CCN5"
## [137] "CCNA1"               "CCNP"
## [139] "CD1E"                 "CD207"
## [141] "CD276"               "CD300LG"
## [143] "CD83"                 "CDC45"
## [145] "CDC6"                 "CDCA3"
## [147] "CDCA5"               "CDH11"
## [149] "CDH19"               "CDKN3"
## [151] "CDSN_3"              "CDSN_4"
## [153] "CDX2"                 "CEACAM1"
## [155] "CEACAM7"              "CECR2"
## [157] "CENPA"                "CENPF"
## [159] "CENPH"               "CENPM"
## [161] "CENPV"                "CEP55"
## [163] "CES2"                  "CFAP251"
## [165] "CFAP47"               "CFAP53"
## [167] "CFD"                   "CFD_1"
## [169] "CGNL1"                "CHGB"
## [171] "CHODL"                "CHP2"
## [173] "CHRDL1"               "CHRNA1"
## [175] "CHST2"                 "CILP"
## [177] "CILP2"                 "CIMAP2"
## [179] "CIP2A"                 "CITED2"
## [181] "CKS1B"                 "CKS2"
## [183] "CLCA4"                 "CLDN17"
## [185] "CLEC12A-AS1"           "CLEC3B"
## [187] "CLIC3"                 "CLSPN"
## [189] "CNFN"                  "CNGB1"
## [191] "CNIH2"                 "CNN1"
## [193] "CNTN5"                 "CNTNAP2"
## [195] "CNTNAP2_1"              "COCH"
## [197] "COL10A1"               "COL11A1"
## [199] "COL24A1"               "COL27A1"
## [201] "COL2A1"                 "COL7A1"
## [203] "CPEB3"                 "CPLX2"
## [205] "CPNE6"                 "CPNE6_1"
## [207] "CPXM1"                 "CRABP1"
## [209] "CRCT1"                 "CRISP2"
## [211] "CRISP3"               "CRMA"
## [213] "CRNN"                  "CRTAC1"
## [215] "CRYBG1"                "CRYBG2"
## [217] "CSAG2_1"               "CSAG3"
## [219] "CSAG3_1"               "CSF2"
## [221] "CSR P1"                "CST1"
## [223] "CST2"                  "CSTB"
## [225] "CT45A5"                 "CTHRC1"
## [227] "CTNNA3"                 "CTSC"
## [229] "CTTN-DT"                "CTTN-DT_1"
## [231] "CTTNBP2"                "CXCL1"
## [233] "CXCL11"                 "CXCL12"
## [235] "CXCL14"                 "CXCL5"
## [237] "CXCL6"                  "CXCR2"

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## [239] "CYP11A1"          "CYP11A1_1"
## [241] "CYP1A1"           "CYP26A1"
## [243] "CYP2C18"          "CYP2J2"
## [245] "CYP4B1"           "CYP4F12"
## [247] "CYP4F22"          "CYP4F29P"
## [249] "CYP4F35P"         "CYSRT1"
## [251] "DAAM2"            "DAPP1"
## [253] "DEFB4B_2"         "DEGS2"
## [255] "DENND2C"          "DES"
## [257] "DHCR24"           "DHCR7-DT"
## [259] "DHRS2"             "DIP2C-AS1"
## [261] "DKK1"              "DLGAP1-AS1"
## [263] "DLGAP5"            "DLL3"
## [265] "DLX1"              "DLX1_1"
## [267] "DLX2"              "DLX6"
## [269] "DLX6-AS1"         "DMP1"
## [271] "DMRT1"            "DNAH14"
## [273] "DNAH17"           "DNAH5"
## [275] "DNMT3B"           "DOCK9"
## [277] "DOP1B"             "DPF1"
## [279] "DPYSL5"           "DRAXIN"
## [281] "DSG3"               "DTL"
## [283] "DUOX1"             "DUSP9"
## [285] "DUXAP10"           "DUXAP8"
## [287] "DUXAP9"            "DYNAP"
## [289] "E2F1"               "E2F7"
## [291] "ECM1"               "ECRG4"
## [293] "ECT2"               "EGF"
## [295] "EHD3"               "EHF"
## [297] "EIF5A2"            "ELOVL4"
## [299] "EMID1"              "EMP1"
## [301] "EMSY-DT"           "EN1"
## [303] "EN2"                "ENDOU"
## [305] "ENTREP1"            "EPB41L4A"
## [307] "EPCAM"              "EPGN"
## [309] "EPHA1"              "EPHA1_1"
## [311] "EPHB2"              "EPHX2"
## [313] "EPHX3"              "EPS8L1"
## [315] "EPS8L2"            "EREG"
## [317] "ERFE"               "ESM1"
## [319] "ESPL1_1"            "ETFDH"
## [321] "EVA1A"              "EVPL"
## [323] "F10"                 "FABP4"
## [325] "FADD"               "FADS1"
## [327] "FADS2"              "FADS3"
## [329] "FALEC"              "FAM107A"
## [331] "FAM135A"            "FAM25A"
## [333] "FAM3B"              "FAM3D"
## [335] "FAR2P1"              "FBN2"
## [337] "FCER1A"              "FCER1A_1"
## [339] "FCHO2"              "FEZF1"
## [341] "FEZF1-AS1"          "FGF12"
## [343] "FGF19"              "FGF5"
## [345] "FHAD1"              "FHL1"

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## [347] "FIBCD1"                      "FIRRE"
## [349] "FLG"                          "FLNC"
## [351] "FLOT1_1"                      "FNDC4"
## [353] "FOXD2-AS1"                    "FOXD3-AS1"
## [355] "FOXI2"                        "FOXI3"
## [357] "FOXL1"                        "FOXL2"
## [359] "FOXM1"                        "FOXO6_1"
## [361] "FREM2"                        "FRMD4B"
## [363] "FUT3_1"                       "FUT6"
## [365] "FXYD1"                        "GAB1"
## [367] "GABRA4"                       "GABRP"
## [369] "GAL"                           "GALNT17"
## [371] "GAP43"                         "GAST"
## [373] "GATA4"                        "GBP6"
## [375] "GBX2"                          "GCKR"
## [377] "GCNT4"                        "GCOM1"
## [379] "GDA"                           "GDF15"
## [381] "GDF7"                          "GDPD2"
## [383] "GDPD3"                        "GFOD2"
## [385] "GFRA1"                        "GGT6"
## [387] "GINS1"                        "GINS2"
## [389] "GJB7"                          "GLI1"
## [391] "GLI2"                           "GLTP"
## [393] "GNG7"                          "GPC2"
## [395] "GPD1L"                         "GPIHBP1"
## [397] "GPR149"                        "GPR158"
## [399] "GPR19"                         "GPR50"
## [401] "GPT"                            "GPT2"
## [403] "GPX3"                          "GREM2"
## [405] "GRHL1"                         "GRHL3"
## [407] "GRIN2D"                        "GRM4"
## [409] "GRP"                            "GSC"
## [411] "GTF2H2"                        "GTF3C2-AS2"
## [413] "GUCY1B2"                       "GYS2"
## [415] "H2AC19"                        "H2BC9"
## [417] "H3C8"                          "HAGHL"
## [419] "HAGLR"                         "HAGLROS"
## [421] "HAP1"                           "HCG22_1"
## [423] "HCG22_2"                       "HCG22_3"
## [425] "HCG22_6"                       "HCG22_7"
## [427] "HES6"                           "HEY1"
## [429] "HHIP"                           "HHIPL2"
## [431] "HJURP"                         "HLA-DOA_5"
## [433] "HLA-DQB2_3"                     "HLA-DQB2_5"
## [435] "HMCN2"                         "HMGA2"
## [437] "HMGB3"                         "HMGCS2"
## [439] "HMMR"                           "HMX1"
## [441] "HMX1_1"                        "HOMER3"
## [443] "HOTAIR"                        "HOXA10"
## [445] "HOXA10-AS"                     "HOXA11"
## [447] "HOXA13"                        "HOXA9"
## [449] "HOXB6"                         "HOXB7"
## [451] "HOXB8"                         "HOXB9"
## [453] "HOXC10"                        "HOXC11"

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## [461] "HOXD8"                        "HPGD"
## [463] "HPSE2"                         "HS3ST5"
## [465] "HSPA6"                         "HSPB6"
## [467] "HSPB7"                         "HTR2C"
## [469] "HTR3B"                         "IBSP"
## [471] "IFI6"                           "IFIT1"
## [473] "IGF2BP1"                      "IGF2BP2"
## [475] "IGFBP3"                        "IGFL1"
## [477] "IL11"                           "IL12A"
## [479] "IL12A-AS1"                     "IL13RA2"
## [481] "IL18"                           "IL1RN"
## [483] "IL24"                           "IL31RA"
## [485] "IL34"                           "IL36A"
## [487] "IL36B"                          "IL36G"
## [489] "INHBA"                         "IP6K3"
## [491] "IRS4"                           "ISG15"
## [493] "IVL"                            "JAKMIP3"
## [495] "JPH2"                           "KALRN"
## [497] "KANK1"                          "KAT2B"
## [499] "KAZN"                           "KCNA5"
## [501] "KCNB1"                          "KCNC3"
## [503] "KCNG3"                         "KCNK3"
## [505] "KCNMA1"                        "KCNMB1"
## [507] "KCNMB2-AS1"                     "KCNMB3"
## [509] "KCNS3"                          "KIAA1549"
## [511] "KIF14"                          "KIF1A"
## [513] "KIF26B"                        "KIF4A"
## [515] "KILH"                           "KLHDC7B"
## [517] "KLHDC7B-DT"                     "KLHDC8A"
## [519] "KLK11"                          "KLK12"
## [521] "KLK13"                          "KPRP"
## [523] "KREMEN2"                       "KRT13"
## [525] "KRT24"                          "KRT3"
## [527] "KRT32"                          "KRT4"
## [529] "KRT42P"                         "KRT75"
## [531] "KRT78"                          "KRT8"
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## [535] "LAMB4"                          "LAMC2"
## [537] "LAMP3"                          "LAYN"
## [539] "LCAL1"                          "LCN1"
## [541] "LCN10"                          "LDB3"
## [543] "LGI1"                           "LGI3"
## [545] "LHX2"                           "LHX5"
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## [549] "LINC00519"                     "LINC00622"
## [551] "LINC00649"                     "LINC00942"
## [553] "LINC01088"                     "LINC01234"
## [555] "LINC01269"                     "LINC01297-DUXAP10-NBEAP6"
## [557] "LINC01305"                     "LINC01614"
## [559] "LINC01615"                     "LINC01873"
## [561] "LINC02041_1"                    "LINC02043"

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## [563] "LINC02086"          "LINC02137"
## [565] "LINC02154"          "LINC02159"
## [567] "LINC02253"          "LINC02321"
## [569] "LINC02457"          "LINC02487"
## [571] "LINC02538"          "LINC02560"
## [573] "LINC02561"          "LINC02577"
## [575] "LINC02609"          "LINC02623"
## [577] "LINC02636"          "LINC02827"
## [579] "LINC02884"          "LINC02893"
## [581] "LINC02994"          "LINC02999"
## [583] "LINC03011"          "LINC03016"
## [585] "LINC03040"          "LINC03050"
## [587] "LINC03057"          "LIPH"
## [589] "LM07"                "LMOD1"
## [591] "LMX1B"               "LOC100287072"
## [593] "LOC101927293"        "LOC101927469"
## [595] "LOC101928051"        "LOC101928391"
## [597] "LOC101928844"        "LOC101928844_1"
## [599] "LOC101929007"        "LOC101929594"
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## [603] "LOC102723686"        "LOC102723825"
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## [613] "LOC105369526"        "LOC105369601"
## [615] "LOC105369887"        "LOC105369975"
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## [623] "LOC105371207"        "LOC105371216"
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## [627] "LOC105371956"        "LOC105372130"
## [629] "LOC105373421"        "LOC105373582"
## [631] "LOC105373949"        "LOC105374122"
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## [663] "LOC107987365"        "LOC107987401"
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## [667] "LOC112267868"        "LOC112268114"
## [669] "LOC112268153"        "LOC122513141"

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## [675] "LOC124901002" "LOC124901321"
## [677] "LOC124901380" "LOC124901465"
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## [707] "LOC124906047" "LOC124906245"
## [709] "LOC124906252" "LOC124906274"
## [711] "LOC124906315" "LOC124907726"
## [713] "LOC124907737" "LOC124907854"
## [715] "LOC124907910" "LOC124907946"
## [717] "LOC408186" "LOC440910"
## [719] "LOC730101" "LOXL2"
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## [723] "LRP12" "LRRC15"
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## [727] "LYPD3" "LYVE1"
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## [731] "MACC1" "MAFA"
## [733] "MAGEA1" "MAGEA11"
## [735] "MAGEA3" "MAGEA3_1"
## [737] "MAGEA4" "MAGEA6_1"
## [739] "MAL" "MALL"
## [741] "MALLP2" "MAPK12"
## [743] "MAPT" "MAPT_1"
## [745] "MAPT_2" "MARCKSL1"
## [747] "MC1R" "MCCC2_1"
## [749] "MCIDAS" "MCM2"
## [751] "MCOLN3" "MDK"
## [753] "MECOM-AS1" "MEGF10"
## [755] "MELTF" "MELTF-AS1"
## [757] "MEST" "METTL21A"
## [759] "MFAP2" "MFAP2_1"
## [761] "MGLL" "MINDY1"
## [763] "MIR9-3HG" "MISP"
## [765] "MLLT11" "MME"
## [767] "MMP1" "MMP10"
## [769] "MMP11" "MMP11_1"
## [771] "MMP12" "MMP13"
## [773] "MMP3" "MND1"
## [775] "MPZL3" "MS4A4E"
## [777] "MSANTD3" "MSC"

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## [779] "MSI1"                      "MSX2"
## [781] "MUC21"                      "MUC21_1"
## [783] "MUC21_2"                    "MUC21_3"
## [785] "MUC22"                      "MUC22_1"
## [787] "MUC22_2"                    "MUC22_7"
## [789] "MUC5B"                      "MUCL1"
## [791] "MUCL3"                      "MUCL3_1"
## [793] "MUCL3_2"                    "MUSTN1"
## [795] "MYBL2"                      "MYH11"
## [797] "MYH11_1"                    "MYL9"
## [799] "MYOC"                       "MYOCD"
## [801] "MYOZ1"                      "MYRIP"
## [803] "MYZAP"                      "N4BP3"
## [805] "NAT14"                      "NAT8L"
## [807] "NCCRP1"                     "NCK1-DT"
## [809] "NDC80"                      "NDRG2"
## [811] "NEB"                         "NECAB2"
## [813] "NEK10"                      "NEK2"
## [815] "NELL2"                      "NEURL3"
## [817] "NEXN"                       "NGB"
## [819] "NIBAN2"                     "NICOL1"
## [821] "NIPAL1"                     "NKAIN2"
## [823] "NKX2-1"                    "NLRX1"
## [825] "NMB"                        "NMB_1"
## [827] "NMRAL2P"                   "NOS2"
## [829] "NOTUM"                      "NPAS2"
## [831] "NPBWR1"                     "NPY1R"
## [833] "NROB1"                      "NR CAM"
## [835] "NREP"                        "NRIP3"
## [837] "NT5C2"                      "NTRK3"
## [839] "NTS"                         "NUF2"
## [841] "NXPH4"                      "OCLN_1"
## [843] "ODC1"                       "OGDHL"
## [845] "OGN"                        "OGN_1"
## [847] "ONECUT2"                    "OPRK1"
## [849] "ORC6"                       "OTOP3"
## [851] "OTX1"                        "P2RX1"
## [853] "P2RX2"                      "P2RY14"
## [855] "P3H4"                        "P4HA3"
## [857] "PABIR3"                     "PADI1"
## [859] "PADI1_1"                    "PAQR5"
## [861] "PAX3"                        "PAX6"
## [863] "PAX7"                        "PAX9"
## [865] "PBK"                         "PBX2_2"
## [867] "PCAT7"                      "PCDH1"
## [869] "PCDHB9_1"                   "PCLAF"
## [871] "PCP4"                        "PCP4_1"
## [873] "PCSK1N"                     "PCSK2"
## [875] "PDE11A"                     "PDGFA"
## [877] "PDK4"                        "PDLIM2"
## [879] "PDZRN3"                     "PDZRN4"
## [881] "PFN2"                        "PGM2L1"
## [883] "PGM5"                        "PGM5-AS1"
## [885] "PGM5P4"                     "PHACTR2"

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## [887] "PHACTR4"          "PHLDB3"
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## [891] "PIMREG"             "PITX1"
## [893] "PKD1L1"             "PKHD1L1"
## [895] "PKP3"                "PLA2G2A"
## [897] "PLA2G2C"             "PLA2G4B"
## [899] "PLAAT2"              "PLAU"
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## [905] "PLEKHM1"             "PLEKHM1_2"
## [907] "PLIN4"                "PLN"
## [909] "PLP1"                  "PLPP4"
## [911] "PMM1"                  "POMC"
## [913] "POPDC3"              "POSTN"
## [915] "POTEF_1"              "POU6F2-AS2"
## [917] "PPL"                   "PPP1R12B"
## [919] "PPP1R1A"              "PPP1R1C"
## [921] "PRAME"                 "PRAME_1"
## [923] "PRDM13"               "PRELID3A"
## [925] "PRELP"                  "PROC"
## [927] "PRR15"                  "PRR15-DT"
## [929] "PRR20G"                 "PRR4"
## [931] "PRR4_1"                  "PRR4_2"
## [933] "PRR5L"                  "PRSS21"
## [935] "PRSS27"                 "PRSS3"
## [937] "PRUNE2"                 "PSCA"
## [939] "PTGIS"                  "PTGS2"
## [941] "PTH2R"                  "PTHLH"
## [943] "PTK6"                   "PTPRN"
## [945] "PWWP3B"                 "PXDN"
## [947] "PYGM"                   "PYY2"
## [949] "RAB15"                  "RAB25"
## [951] "RAB32"                  "RAB3B"
## [953] "RAB40A"                 "RAB9B"
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## [957] "RAD51AP1"                "RAET1E"
## [959] "RAG1"                   "RANBP9"
## [961] "RARG"                   "RASL11B"
## [963] "RBFOX3"                 "RBM47"
## [965] "RBP1"                   "RBP4"
## [967] "RBPM2S"                 "RDH12"
## [969] "RELCH"                  "RFC4"
## [971] "RHCg"                   "RHEBL1"
## [973] "RIMS2"                  "RMND5B"
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## [979] "RNFT2"                  "ROR1"
## [981] "ROS1"                   "RPL39L"
## [983] "RRAD"                   "SALL4"
## [985] "SAMD5"                  "SASH1"
## [987] "SCARA5"                 "SCAT2"
## [989] "SCAT8"                  "SCEL"
## [991] "SCG2"                   "SCGB1D2"
## [993] "SCGB2A2"                "SCGB3A1"

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## [999] "SCN9A"                       "SCNN1B"
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## [1009] "SERPINH1"                   "SEZ6L2"
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## [1013] "SH3BGRL2"                  "SHH"
## [1015] "SHISA2"                    "SHOX2"
## [1017] "SIM2"                       "SIX1"
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## [1029] "SLC2A4"                    "SLC2A4_1"
## [1031] "SLC30A3"                   "SLC34A3"
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## [1035] "SLC51A"                    "SLC52A1"
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## [1039] "SLC6A2"                    "SLC7A5P2_1"
## [1041] "SLC8A1-AS1"                "SLC01A2"
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## [1051] "SNCB"                      "SNHG33"
## [1053] "SNORD3A"                  "SNX10"
## [1055] "SORBS1"                   "SORCS1"
## [1057] "SORT1"                     "SOST"
## [1059] "SOX10"                     "SOX11"
## [1061] "SOX4"                      "SPC24"
## [1063] "SPC25_1"                  "SPINDOC"
## [1065] "SPINK5"                   "SPNS2"
## [1067] "SPOCD1"                   "SPP1"
## [1069] "SPRR2G"                   "SPRR3"
## [1071] "SPSB4"                     "SPTLC3"
## [1073] "SRL"                       "ST6GALNAC1"
## [1075] "ST8SIA2"                  "STIM1"
## [1077] "STK40"                     "STMN1"
## [1079] "STRA6"                     "STRA6_1"
## [1081] "STRN"                      "STUM"
## [1083] "STX1A"                     "SULT2B1"
## [1085] "SYCP2"                     "SYNDIG1"
## [1087] "SYNM"                      "SYNPO2"
## [1089] "SYNP02L"                  "SYT1"
## [1091] "SYT14"                     "TACR1"
## [1093] "TACR2"                     "TAGLN"
## [1095] "TBX15"                     "TBX6"
## [1097] "TCF15"                     "TDO2"
## [1099] "TD02_1"                   "TEC"
## [1101] "TENT5B"                   "TESMIN"

```

```

## [1103] "TFAP2B"                      "TGFBI"
## [1105] "TGFBR3"                       "TGM1"
## [1107] "TGM1_1"                        "TGM3"
## [1109] "TIAM1"                         "TJP1_1"
## [1111] "TJP1_2"                        "TJP3"
## [1113] "TK1"                           "TM4SF19"
## [1115] "TMEFF1"                        "TMEM132C"
## [1117] "TMEM154"                       "TMEM35A"
## [1119] "TMEM44-AS1"                   "TMEM79"
## [1121] "TMOD3"                         "TMPRSS11B"
## [1123] "TNFAIP6"                       "TNFRSF18"
## [1125] "TNXB_7"                        "TOLLIP"
## [1127] "TOLLIP_1"                      "TOM1"
## [1129] "TOM1L2"                        "TOP2A"
## [1131] "TPM2"                          "TPRG1"
## [1133] "TREH"                           "TREM1"
## [1135] "TREM2"                         "TRIM29"
## [1137] "TRIM46"                        "TRIP13"
## [1139] "TRPM2-AS"                     "TTC39A"
## [1141] "TTC9"                           "TTYH3"
## [1143] "TUBB3"                         "UBE2C"
## [1145] "UBE2T"                         "UBL3"
## [1147] "UCHL1"                         "UGT1A8"
## [1149] "UNC13B"                        "UNC5B-AS1"
## [1151] "UNC93A"                        "UPK1A"
## [1153] "UPK3B"                          "UPK3B_1"
## [1155] "USP2"                           "USP32P3"
## [1157] "USP6NL"                        "UTS2R"
## [1159] "VAX1"                           "VGF"
## [1161] "VIT"                            "VPS37D"
## [1163] "VPS4B"                          "VSIG10L"
## [1165] "VSIG2"                          "VWA5B2"
## [1167] "WDR54"                         "WDR72"
## [1169] "WNK4"                           "WNT11"
## [1171] "WNT2"                           "XCL1"
## [1173] "XIRP1"                         "XIST"
## [1175] "XPNPEP2"                       "ZBTB16"
## [1177] "ZDHHC15"                       "ZDHHC21"
## [1179] "ZIC1"                           "ZIC2"
## [1181] "ZIC5"                           "ZNF114"
## [1183] "ZNF185"                         "ZNF185_1"
## [1185] "ZNF365"                         "ZNF431"
## [1187] "ZNF469"                         "ZNF536"
## [1189] "ZNF584-DT"                     "ZNF677"
## [1191] "ZNF750"                         "ZNF823"

```

```
length(deg_list)
```

```
## [1] 1192
```

```
write.table(degs, sep = "\t", file = "rna_seq_degs.tsv",
           row.names = F, col.names = T)
```