

# Venn diagrams

Required packages

```
library(dplyr)

##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
##   filter, lag
## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union
library(tibble)
library(grid)
library(stringr)
library(gplots)

##
## Attaching package: 'gplots'
## The following object is masked from 'package:stats':
##
##   lowess
library(VennDiagram)

## Loading required package: futile.logger
library(extrafont)

## Registering fonts with R
```

## Purpose:

To determine the overlap in differentially expressed genes between timepoints d8 and d28 post-infection for the various infection conditions.

Upload files

```
data_dir <- "Human DGEs_donortreatment"
sampleFiles <- basename(Sys.glob(file.path(data_dir, "*.txt")))
sampleNames <- str_replace(sampleFiles, "[0-9]*-[0-9]*-[0-9]*", "") %>%
  str_replace("_analysis_results.txt", "")
sampleNames

## [1] "human_donor_treatmentHumanHBVgenes-HBV_vs_mock_d28"
## [2] "human_donor_treatmentHumanHBVgenes-HBV_vs_mock_d8"
## [3] "human_donor_treatmentHumanHBVgenes-coinf_vs_HBV_d28"
## [4] "human_donor_treatmentHumanHBVgenes-coinf_vs_HBV_d8"
## [5] "human_donor_treatmentHumanHBVgenes-coinf_vs_mock_d28"
## [6] "human_donor_treatmentHumanHBVgenes-coinf_vs_mock_d8"
```

```

read_in <- function(files) {
  d <- read.delim(files, header = TRUE)
  dd <- dplyr::select(d, log2FoldChange, padj, X) %>%
    na.omit() %>%
    dplyr::filter(padj <= 0.05) %>%
    dplyr::filter(abs(log2FoldChange) >= 0.5) %>%
    select(X, log2FoldChange)
  dd
}

subset <- lapply(file.path(data_dir, sampleFiles), read_in)
names(subset) <- sampleNames

for(i in names(subset)) {
  filename <- paste(i, "Venn_subset.txt", sep = "")
  write.table(subset[i], file = file.path("Venn diagrams", filename),
    col.names = FALSE, row.names=FALSE, sep="\t", quote=FALSE)
}

subset_up <- lapply(subset, function(x) dplyr::filter(x, log2FoldChange > 0)) %>%
  sapply(nrow)
subset_up

##   human_donor_treatmentHumanHBVgenes-HBV_vs_mock_d28
##                                     4
##   human_donor_treatmentHumanHBVgenes-HBV_vs_mock_d8
##                                     61
##   human_donor_treatmentHumanHBVgenes-coinf_vs_HBV_d28
##                                     0
##   human_donor_treatmentHumanHBVgenes-coinf_vs_HBV_d8
##                                     67
##   human_donor_treatmentHumanHBVgenes-coinf_vs_mock_d28
##                                     4
##   human_donor_treatmentHumanHBVgenes-coinf_vs_mock_d8
##                                     53

write.table(subset_up, file = "Venn diagrams/Venn genes upregulated.txt",
  col.names = FALSE, row.names=TRUE, sep="\t", quote=FALSE)

subset_down <- lapply(subset, function(x) dplyr::filter(x, log2FoldChange < 0)) %>%
  sapply(nrow)
subset_down

##   human_donor_treatmentHumanHBVgenes-HBV_vs_mock_d28
##                                     1
##   human_donor_treatmentHumanHBVgenes-HBV_vs_mock_d8
##                                     102
##   human_donor_treatmentHumanHBVgenes-coinf_vs_HBV_d28
##                                     10
##   human_donor_treatmentHumanHBVgenes-coinf_vs_HBV_d8
##                                     58
##   human_donor_treatmentHumanHBVgenes-coinf_vs_mock_d28
##                                     18

```

```
## human_donor_treatmentHumanHBVgenes-coinf_vs_mock_d8
## 95

write.table(subset_down, file = "Venn diagrams/Venn genes downregulated.txt",
            col.names = FALSE, row.names=TRUE, sep="\t", quote=FALSE)

genes_only <- lapply(subset, function(x) dplyr::select(x, X))

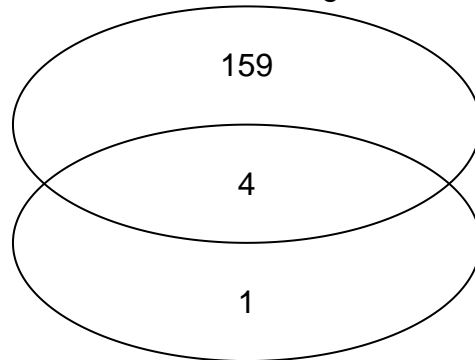
##To keep track of the names of the different elements within the list
names(genes_only)

## [1] "human_donor_treatmentHumanHBVgenes-HBV_vs_mock_d28"
## [2] "human_donor_treatmentHumanHBVgenes-HBV_vs_mock_d8"
## [3] "human_donor_treatmentHumanHBVgenes-coinf_vs_HBV_d28"
## [4] "human_donor_treatmentHumanHBVgenes-coinf_vs_HBV_d8"
## [5] "human_donor_treatmentHumanHBVgenes-coinf_vs_mock_d28"
## [6] "human_donor_treatmentHumanHBVgenes-coinf_vs_mock_d8"

##For determining overlap areas, use gplots venn function
venn1 <- venn(genes_only[1:2])
```

```
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
## Warning in getVennCounts.list(l, universe, intersections = TRUE, ...): List element [[2]] has dimens
## Warning in getVennCounts.list(l, universe, intersections = TRUE, ...): List element [[1]] has dimens
## Warning in getVennCounts.list(l, universe, intersections = TRUE, ...): List element [[1]] has dimens
## Warning in getVennCounts.list(l, universe, intersections = TRUE, ...): List element [[2]] has dimens
## Warning in getVennCounts.list(l, universe, intersections = TRUE, ...): List element [[1]] has dimens
## Warning in getVennCounts.list(l, universe, intersections = TRUE, ...): List element [[2]] has dimens

human_donor_treatmentHumanHBVgenes-HBV_vs_mock_d8
```



```
human_donor_treatmentHumanHBVgenes-HBV_vs_mock_d28
```

```
attr(venn1, "intersections")
```

```

## $`human_donor_treatmentHumanHBVgenes-HBV_vs_mock_d28:human_donor_treatmentHumanHBVgenes-HBV_vs_mock_
## [1] "AAB59969.1" "AAB59972.1" "AAB59971.1" "AAB59970.1"
##
## $`human_donor_treatmentHumanHBVgenes-HBV_vs_mock_d8`
## [1] "ENSG00000130508" "ENSG00000179776" "ENSG00000187498"
## [4] "ENSG00000261371" "ENSG00000125810" "ENSG00000136872"
## [7] "ENSG00000138207" "ENSG00000087303" "ENSG00000091879"
## [10] "ENSG00000110245" "ENSG00000087245" "ENSG00000134871"
## [13] "ENSG00000118523" "ENSG00000158104" "ENSG00000134013"
## [16] "ENSG00000255974" "ENSG00000142798" "ENSG00000100652"
## [19] "ENSG00000163975" "ENSG00000150938" "ENSG00000143819"
## [22] "ENSG00000011028" "ENSG00000196616" "ENSG00000163453"
## [25] "ENSG00000154133" "ENSG00000169744" "ENSG00000164692"
## [28] "ENSG00000124762" "ENSG00000165092" "ENSG00000118137"
## [31] "ENSG00000198848" "ENSG00000106991" "ENSG00000196569"
## [34] "ENSG00000172831" "ENSG00000166592" "ENSG00000248144"
## [37] "ENSG00000204301" "ENSG00000151790" "ENSG00000108821"
## [40] "ENSG00000244067" "ENSG00000196502" "ENSG00000151365"
## [43] "ENSG00000161638" "ENSG00000116962" "ENSG00000145623"
## [46] "ENSG00000118257" "ENSG00000175003" "ENSG00000166086"
## [49] "ENSG00000140945" "ENSG00000100031" "ENSG00000175899"
## [52] "ENSG00000244474" "ENSG00000116661" "ENSG00000091136"
## [55] "ENSG00000010327" "ENSG00000186642" "ENSG00000137801"
## [58] "ENSG00000196628" "ENSG00000130300" "ENSG00000161940"
## [61] "ENSG00000160282" "ENSG00000164125" "ENSG00000171724"
## [64] "ENSG00000175274" "ENSG00000113140" "ENSG00000184557"
## [67] "ENSG00000256612" "ENSG00000214548" "ENSG00000171747"
## [70] "ENSG00000257335" "ENSG00000164161" "ENSG00000021826"
## [73] "ENSG00000004399" "ENSG00000143772" "ENSG00000017427"
## [76] "ENSG00000134954" "ENSG00000164283" "ENSG00000143845"
## [79] "ENSG00000166341" "ENSG00000188338" "ENSG00000064989"
## [82] "ENSG00000115380" "ENSG00000100889" "ENSG00000124875"
## [85] "ENSG00000149591" "ENSG00000162460" "ENSG00000131747"
## [88] "ENSG00000156515" "ENSG00000145321" "ENSG00000162482"
## [91] "ENSG00000196600" "ENSG00000270504" "ENSG00000153936"
## [94] "ENSG00000169738" "ENSG00000143627" "ENSG00000157510"
## [97] "ENSG00000112902" "ENSG00000147862" "ENSG00000142910"
## [100] "ENSG00000182704" "ENSG00000196083" "ENSG00000141756"
## [103] "ENSG00000187193" "ENSG00000145555" "ENSG00000151388"
## [106] "ENSG00000125144" "ENSG00000143416" "ENSG00000066056"
## [109] "ENSG00000100003" "ENSG00000153822" "ENSG00000106804"
## [112] "ENSG00000071246" "ENSG00000171388" "ENSG00000170439"
## [115] "ENSG00000087116" "ENSG00000134817" "ENSG00000118785"
## [118] "ENSG00000128052" "ENSG00000158125" "ENSG00000198077"
## [121] "ENSG00000164266" "ENSG00000072080" "ENSG00000132693"
## [124] "ENSG00000026025" "ENSG00000177556" "ENSG00000189334"
## [127] "ENSG00000169242" "ENSG00000135424" "ENSG00000148803"
## [130] "ENSG00000129422" "ENSG00000133048" "ENSG00000140505"
## [133] "ENSG00000186910" "ENSG00000144908" "ENSG00000162618"
## [136] "ENSG00000142173" "ENSG00000168306" "ENSG00000135114"
## [139] "ENSG00000147113" "ENSG00000198099" "ENSG00000129657"
## [142] "ENSG00000035862" "ENSG00000166391" "ENSG00000111348"
## [145] "ENSG00000105825" "ENSG00000170921" "ENSG00000136856"
## [148] "ENSG00000239887" "ENSG00000113600" "ENSG00000120899"

```

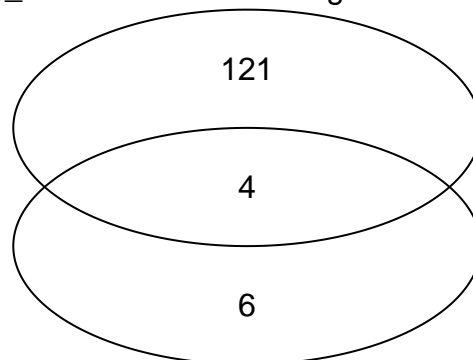
```
## [151] "ENSG00000123243" "ENSG00000112419" "ENSG00000260314"
## [154] "ENSG00000171603" "ENSG00000186908" "ENSG00000131495"
## [157] "ENSG00000110799" "ENSG00000129538" "ENSG00000107957"
##
## $`human_donor_treatmentHumanHBVgenes-HBV_vs_mock_d28`
## [1] "ENSG00000126838"
```

```
##Four in common (all HBV); otherwise d8 has 159; d28 has 1.
```

```
venn2 <- venn(genes_only[3:4])
```

```
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
## Warning in getVennCounts.list(1, universe, intersections = TRUE, ...): List element [[2]] has dimens
## Warning in getVennCounts.list(1, universe, intersections = TRUE, ...): List element [[1]] has dimens
## Warning in getVennCounts.list(1, universe, intersections = TRUE, ...): List element [[1]] has dimens
## Warning in getVennCounts.list(1, universe, intersections = TRUE, ...): List element [[2]] has dimens
## Warning in getVennCounts.list(1, universe, intersections = TRUE, ...): List element [[1]] has dimens
## Warning in getVennCounts.list(1, universe, intersections = TRUE, ...): List element [[2]] has dimens
```

human\_donor\_treatmentHumanHBVgenes-coinf\_vs\_HBV\_d8



human\_donor\_treatmentHumanHBVgenes-coinf\_vs\_HBV\_d28

```
attr(venn2, "intersections")
```

```
## $`human_donor_treatmentHumanHBVgenes-coinf_vs_HBV_d28:human_donor_treatmentHumanHBVgenes-coinf_vs_HBV`
## [1] "ENSG00000130300" "ENSG00000128052" "ENSG00000261371" "ENSG00000154133"
##
## $`human_donor_treatmentHumanHBVgenes-coinf_vs_HBV_d8`
## [1] "ENSG00000011465" "ENSG00000168542" "ENSG00000118785"
## [4] "ENSG00000142748" "ENSG00000091879" "ENSG00000125810"
## [7] "ENSG00000102755" "ENSG00000196569" "ENSG00000145321"
## [10] "ENSG00000010327" "ENSG00000054654" "ENSG00000039537"
## [13] "ENSG00000108821" "ENSG00000164692" "ENSG00000162618"
```

```
## [16] "ENSG00000166391" "ENSG00000152377" "ENSG00000151388"
## [19] "ENSG00000118946" "ENSG00000164188" "ENSG00000142156"
## [22] "ENSG00000188257" "ENSG00000125144" "ENSG00000164283"
## [25] "ENSG00000165092" "ENSG00000146072" "ENSG00000164850"
## [28] "ENSG00000129538" "ENSG00000132693" "ENSG00000163687"
## [31] "ENSG00000125966" "ENSG00000177575" "ENSG00000138315"
## [34] "ENSG00000058085" "ENSG00000164120" "ENSG00000124839"
## [37] "ENSG00000179776" "ENSG00000090238" "ENSG00000184374"
## [40] "ENSG00000095303" "ENSG00000136235" "ENSG00000038427"
## [43] "ENSG00000116690" "ENSG00000166592" "ENSG00000140450"
## [46] "ENSG00000106351" "ENSG00000147027" "ENSG00000134363"
## [49] "ENSG00000087245" "ENSG00000119946" "ENSG00000134853"
## [52] "ENSG00000119922" "ENSG00000168631" "ENSG00000163581"
## [55] "ENSG00000186340" "ENSG00000184557" "ENSG00000124145"
## [58] "ENSG00000011028" "ENSG00000115232" "ENSG00000110925"
## [61] "ENSG00000161011" "ENSG00000111799" "ENSG00000142173"
## [64] "ENSG00000253368" "ENSG00000140416" "ENSG00000228278"
## [67] "ENSG00000187955" "ENSG00000087116" "ENSG00000136881"
## [70] "ENSG00000104635" "ENSG00000198840" "ENSG00000276980"
## [73] "ENSG00000239887" "ENSG00000144481" "ENSG00000156968"
## [76] "ENSG00000125845" "ENSG00000205060" "ENSG00000244414"
## [79] "ENSG00000163050" "ENSG00000160285" "ENSG00000102760"
## [82] "ENSG00000106327" "ENSG00000164266" "ENSG00000137054"
## [85] "ENSG00000025423" "ENSG00000115380" "ENSG00000244067"
## [88] "ENSG00000111181" "ENSG00000198121" "ENSG00000152779"
## [91] "ENSG00000032444" "ENSG00000135069" "ENSG00000165169"
## [94] "ENSG00000110693" "ENSG00000139194" "ENSG00000188338"
## [97] "ENSG00000162460" "ENSG00000254166" "ENSG00000165475"
## [100] "ENSG00000136720" "ENSG00000175538" "ENSG00000066739"
## [103] "ENSG00000254607" "ENSG00000198610" "ENSG00000188488"
## [106] "ENSG00000196975" "ENSG00000076351" "ENSG00000123384"
## [109] "ENSG00000188707" "ENSG00000127954" "ENSG00000115884"
## [112] "ENSG00000170458" "ENSG00000144852" "ENSG00000214548"
## [115] "ENSG00000110799" "ENSG00000091436" "ENSG00000135744"
## [118] "ENSG00000204262" "ENSG00000196878" "ENSG00000134871"
## [121] "ENSG00000160282"
##
## $`human_donor_treatmentHumanHBVgenes-coinf_vs_HBV_d28`
## [1] "ENSG00000113555" "ENSG00000107719" "ENSG00000164161" "ENSG00000176435"
## [5] "ENSG00000163762" "ENSG00000133574"
```

```
##Four in common; d8 has 121, d28 has 6
```

```
venn3 <- venn(genes_only[5:6])
```

```
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
```

```
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
```

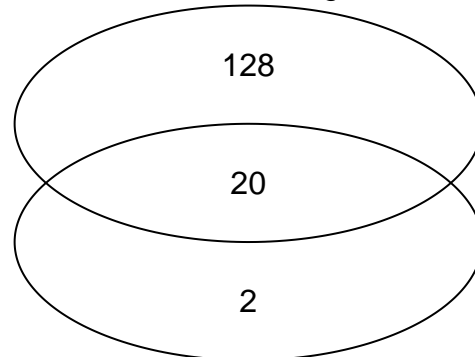
```
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
```

```
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
```

```
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
```

```
## Warning in getVennCounts.list(data, universe = universe, intersections = intersections): List element
```

```
## Warning in getVennCounts.list(l, universe, intersections = TRUE, ...): List element [[2]] has dimens
## Warning in getVennCounts.list(l, universe, intersections = TRUE, ...): List element [[1]] has dimens
## Warning in getVennCounts.list(l, universe, intersections = TRUE, ...): List element [[1]] has dimens
## Warning in getVennCounts.list(l, universe, intersections = TRUE, ...): List element [[2]] has dimens
## Warning in getVennCounts.list(l, universe, intersections = TRUE, ...): List element [[1]] has dimens
## Warning in getVennCounts.list(l, universe, intersections = TRUE, ...): List element [[2]] has dimens
human_donor_treatmentHumanHBVgenes-coinf_vs_mock_d8
```



human\_donor\_treatmentHumanHBVgenes-coinf\_vs\_mock\_d28

```
attr(veenn3, "intersections")
```

```
## $`human_donor_treatmentHumanHBVgenes-coinf_vs_mock_d28:human_donor_treatmentHumanHBVgenes-coinf_vs_m
## [1] "AAB59970.1"      "AAB59969.1"      "AAB59971.1"
## [4] "AAB59972.1"      "ENSG000000136235" "ENSG000000261371"
## [7] "ENSG000000118785" "ENSG000000091879" "ENSG000000069122"
## [10] "ENSG00000010327" "ENSG000000163687" "ENSG000000204301"
## [13] "ENSG000000260314" "ENSG000000176435" "ENSG000000124126"
## [16] "ENSG000000133574" "ENSG000000118946" "ENSG000000135218"
## [19] "ENSG000000164161" "ENSG000000169744"
##
## $`human_donor_treatmentHumanHBVgenes-coinf_vs_mock_d8`
## [1] "ENSG000000125810" "ENSG000000179776" "ENSG000000154133"
## [4] "ENSG000000110799" "ENSG000000142748" "ENSG000000162618"
## [7] "ENSG000000177575" "ENSG000000102755" "ENSG000000129538"
## [10] "ENSG000000164283" "ENSG000000128052" "ENSG000000146038"
## [13] "ENSG000000130300" "ENSG000000064989" "ENSG000000127399"
## [16] "ENSG000000127329" "ENSG000000122254" "ENSG000000118271"
## [19] "ENSG000000188488" "ENSG000000161267" "ENSG000000091583"
## [22] "ENSG000000102760" "ENSG000000106351" "ENSG000000175538"
## [25] "ENSG000000123700" "ENSG000000119922" "ENSG000000105825"
## [28] "ENSG000000038945" "ENSG000000155659" "ENSG000000139567"
## [31] "ENSG000000113555" "ENSG000000091436" "ENSG000000196628"
## [34] "ENSG000000152779" "ENSG000000198408" "ENSG000000111181"
## [37] "ENSG000000108582" "ENSG000000170525" "ENSG000000089220"
## [40] "ENSG000000086548" "ENSG000000054654" "ENSG000000066056"
## [43] "ENSG000000163050" "ENSG000000147113" "ENSG000000214530"
## [46] "ENSG000000001617" "ENSG000000118257" "ENSG000000157554"
## [49] "ENSG000000138722" "ENSG000000128917" "ENSG000000160862"
## [52] "ENSG000000173269" "ENSG000000265972" "ENSG000000123243"
## [55] "ENSG000000110169" "ENSG000000124839" "ENSG000000167191"
```

```

## [58] "ENSG00000135744" "ENSG00000120885" "ENSG00000162511"
## [61] "ENSG00000197629" "ENSG00000058085" "ENSG00000138207"
## [64] "ENSG00000151790" "ENSG00000171345" "ENSG00000137801"
## [67] "ENSG00000116690" "ENSG00000145321" "ENSG00000101445"
## [70] "ENSG00000165124" "ENSG00000196975" "ENSG00000171115"
## [73] "ENSG00000161940" "ENSG00000153162" "ENSG00000166173"
## [76] "ENSG00000117601" "ENSG00000125734" "ENSG00000176046"
## [79] "ENSG00000138079" "ENSG00000121858" "ENSG00000170315"
## [82] "ENSG00000144481" "ENSG00000124882" "ENSG00000231574"
## [85] "ENSG00000125968" "ENSG00000158352" "ENSG00000106927"
## [88] "ENSG00000188707" "ENSG00000184374" "ENSG00000171793"
## [91] "ENSG00000142168" "ENSG00000145012" "ENSG00000138315"
## [94] "ENSG00000071246" "ENSG00000073849" "ENSG00000233276"
## [97] "ENSG00000185950" "ENSG00000143418" "ENSG00000182551"
## [100] "ENSG00000105939" "ENSG00000156096" "ENSG00000132821"
## [103] "ENSG00000213694" "ENSG00000104635" "ENSG00000107719"
## [106] "ENSG00000163735" "ENSG00000170458" "ENSG00000106541"
## [109] "ENSG00000105835" "ENSG00000124145" "ENSG00000125703"
## [112] "ENSG00000164543" "ENSG00000198712" "ENSG00000037280"
## [115] "ENSG00000196611" "ENSG00000173372" "ENSG00000168631"
## [118] "ENSG00000104938" "ENSG00000106991" "ENSG00000153048"
## [121] "ENSG00000211459" "ENSG00000116661" "ENSG00000169903"
## [124] "ENSG00000135678" "ENSG00000131386" "ENSG00000151726"
## [127] "ENSG00000155368" "ENSG00000165092"
##
## $`human_donor_treatmentHumanHBVgenes-coinf_vs_mock_d28`
## [1] "ENSG00000126838" "ENSG00000144668"

venn3_attr <- attr(venn3, "intersections")
## In common 20 total including 4 HBV genes; otherwise d8 has 128; d28 has 2.

##As the only comparison with overlapping genes besides the HBV genes (which we
##know all are > 1), I want to see how the expression of these common genes compare
##(i.e. upregulated versus downregulated at each timepoint).
venn3_intersect <- as.data.frame(venn3_attr[1], stringsAsFactors = FALSE)
colnames(venn3_intersect) <- c("X")

coinfvmockd28_df <- as.data.frame(subset[5]) %>%
  droplevels()
colnames(coinfvmockd28_df) <- c("X", "d28log2FoldChange")
coinfvmockd8_df <- as.data.frame(subset[6]) %>%
  droplevels()
colnames(coinfvmockd8_df) <- c("X", "d8log2FoldChange")
venn3_intersect_l2FC <- inner_join(venn3_intersect, coinfvmockd8_df, by = "X") %>%
  inner_join(coinfvmockd28_df, by = "X")

## Warning: Column `X` joining character vector and factor, coercing into
## character vector

## Warning: Column `X` joining character vector and factor, coercing into
## character vector

write.csv(venn3_intersect_l2FC, file = file.path("Venn diagrams",
  filename = "coinfvmock_d8_d28_intersection.csv"))

```

Session info



## sessionInfo()

```
## R version 3.3.3 (2017-03-06)
## Platform: x86_64-apple-darwin13.4.0 (64-bit)
## Running under: macOS Sierra 10.12.6
##
## locale:
## [1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8
##
## attached base packages:
## [1] grid      stats      graphics  grDevices  utils      datasets  methods
## [8] base
##
## other attached packages:
## [1] bindrcpp_0.2      extrafont_0.17    VennDiagram_1.6.17
## [4] futile.logger_1.4.3 gplots_3.0.1      stringr_1.2.0
## [7] tibble_1.3.3      dplyr_0.7.3
##
## loaded via a namespace (and not attached):
## [1] Rcpp_0.12.10      Rttf2pt1_1.3.5    knitr_1.16
## [4] bindr_0.1         magrittr_1.5      R6_2.2.0
## [7] rlang_0.1.2       caTools_1.17.1    tools_3.3.3
## [10] KernSmooth_2.23-15 extrafontdb_1.0    lambda.r_1.1.9
## [13] htmltools_0.3.5   gtools_3.5.0      yaml_2.1.14
## [16] assertthat_0.2.0  rprojroot_1.2     digest_0.6.12
## [19] futile.options_1.0.0 bitops_1.0-6      glue_1.1.1
## [22] evaluate_0.10     rmarkdown_1.4     gdata_2.17.0
## [25] stringi_1.1.5     backports_1.0.5    pkgconfig_2.0.1
```