

# Cre-IHC-colocalization

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```
setwd("D:/Microscopy/IHC images/20221020 - Opn4-cre-EGFP/analysis/processing/CSVs")
```

```
#um 5 and 7 could have more GFP cells  
wdir <- getwd()
```

```
count_list <- list.files("D:/Microscopy/IHC images/20221020 - Opn4-cre-EGFP/analysis/processing/CSVs",  
length(count_list)
```

```
## [1] 10
```

```
coloc_list <- list.files("D:/Microscopy/IHC images/20221020 - Opn4-cre-EGFP/analysis/processing/CSVs",  
length(coloc_list)
```

```
## [1] 10
```

```
img_sumstats <- data.frame()  
mouse_sumstats <- data.frame()  
dapi_names <- c()  
gfp_names <- c()  
cre_names <- c()  
  
for (file in count_list){  
  counts <- read.csv(paste0(wdir,"/",file))  
  counts <- counts %>%  
    rename(VolUnit = Vol..unit.,  
           VolPix = Vol..pix.) %>%  
    select(c(Nb, Label, Name, VolUnit, VolPix))  
  counts <- counts %>% mutate(Name = str_replace(Name, "obj325-val3", "obj325-val1"),  
                             Name = str_replace(Name, "obj532-val2", "obj532-val1"),  
                             Name = str_replace(Name, "obj802-val2", "obj801-val1"))  
  cutoffs <- counts %>% filter(str_detect(Name, ".*(val1)$"))  
  counts <- counts %>% mutate(type = case_when(  
    between(Nb+1, cutoffs[1,1], cutoffs[2,1]) ~ "DAPI",  
    between(Nb+1, cutoffs[2,1], cutoffs[3,1]) ~ "GFP",  
    Nb+1 > cutoffs[3,1] ~ "cre"  
  ))  
  dapi_names[file] <- counts %>% filter(type == "DAPI") %>% select(Name)  
  gfp_names[file] <- counts %>% filter(type == "GFP") %>% select(Name)  
  cre_names[file] <- counts %>% filter(type == "cre") %>% select(Name)  
  #print(head(counts))
```

```

imgsumstats <- counts %>%
  group_by(type) %>%
  summarise(filename = file,
            n = n(),
            minVol = min(VolUnit),
            medVol = median(VolUnit),
            meanVol = mean(VolUnit),
            maxVol = max(VolUnit))
img_sumstats <- rbind(img_sumstats, imgsumstats)
mousesumstats <- counts %>% filter(type == "GFP") %>%
  mutate(mouse = as.factor(str_extract(file, "(?<=_).*(?=-[:digit:]_)))
mouse_sumstats <- rbind(mouse_sumstats, mousesumstats)
}
img_sumstats

```

```

## # A tibble: 30 x 7
##   type filename                n minVol medVol meanVol maxVol
##   <chr> <chr>                <int> <dbl> <dbl> <dbl> <dbl>
## 1 cre   M_M-6-40x-8avg-1_quant.csv 4744  0.0243  0.292  1.07  505.
## 2 DAPI  M_M-6-40x-8avg-1_quant.csv  375  0.243   1.78   3.41  52.2
## 3 GFP   M_M-6-40x-8avg-1_quant.csv    2 261.    290.   290.   319.
## 4 cre   M_M-6-40x-8avg-2_quant-fixed.csv 664  0.0243  0.243  1.03  205.
## 5 DAPI  M_M-6-40x-8avg-2_quant-fixed.csv 322  0.243   1.96   3.98  154.
## 6 GFP   M_M-6-40x-8avg-2_quant-fixed.csv  4 34.1    244.   224.   373.
## 7 cre   M_M-6-40x-8avg-3_quant-fixed.csv 1066  0.0243  0.219  0.788  199.
## 8 DAPI  M_M-6-40x-8avg-3_quant-fixed.csv 1063  0.243   1.27   3.24  101.
## 9 GFP   M_M-6-40x-8avg-3_quant-fixed.csv  3 139.    155.   164.   198.
## 10 cre  M_M-6-40x-8avg-4_quant-fixed.csv 1933  0.0243  0.268  0.819  121.
## # ... with 20 more rows

```

```

mouse_gfp_sumstats <- mouse_sumstats %>% group_by(mouse) %>%
  summarise(minVol = min(VolUnit),
            medVol = median(VolUnit),
            meanVol = mean(VolUnit),
            maxVol = max(VolUnit))
mouse_gfp_sumstats

```

```

## # A tibble: 2 x 5
##   mouse minVol medVol meanVol maxVol
##   <fct> <dbl> <dbl> <dbl> <dbl>
## 1 M-6-40x-8avg 34.1 211. 204. 373.
## 2 wm-40x-8avg 48.4 210. 222. 424.

```

```

overall_sumstats <- mouse_sumstats %>%
  summarise(minVol = min(VolUnit),
            medVol = median(VolUnit),
            meanVol = mean(VolUnit),
            maxVol = max(VolUnit))
overall_sumstats

```

```

##   minVol medVol meanVol maxVol
## 1 34.11097 210.2982 212.8922 424.2459

```

```
names(dapi_names) <- sub("_quant.*", "", names(dapi_names))
names(dapi_names) <- sub("M_", "", names(dapi_names))
names(gfp_names) <- sub("_quant.*", "", names(gfp_names))
names(gfp_names) <- sub("M_", "", names(gfp_names))
names(cre_names) <- sub("_quant.*", "", names(cre_names))
names(cre_names) <- sub("M_", "", names(cre_names))
```

```
#gfp_names
```

```
coloc_sumstats <- data.frame()

for (file in coloc_list){
  coloc_count <- read.csv(paste0(wdir,"/",file))
  filename <- paste0(file)
  filename <- str_replace(filename, "C_", "")
  filename <- str_replace(filename, "_cre.*", "")
  coloc_count <- coloc_count %>% mutate(Label1 = str_replace(Label1, "obj325-val3", "obj325-val1"),
    Label1 = str_replace(Label1, "obj532-val2", "obj532-val1"),
    Label1 = str_replace(Label1, "obj802-val2", "obj801-val1"))
  coloc_count <- coloc_count %>%
    select(c(Label1, Label2, coloc, PcColoc)) %>%
    filter(Label1 %in% gfp_names[filename][[1]] & Label2 %in% cre_names[filename][[1]])
  #coloc_count <- full_join(coloc_count, cell_types, by=c("Label1"="Name"))
  #coloc_count <- full_join(coloc_count, cell_types, by=c("Label2"="Name"), suffix=c("_1", "_2"))
  #coloc_count <- coloc_count %>% filter(type_1 == "GFP" & type_2 == "cre", na.rm = TRUE)
  sumstats <- coloc_count %>% group_by(Label1) %>%
    summarise(filename = file,
      n = n(),
      colocPerc = sum(PcColoc),
      trueColoc1 = colocPerc >= 1,
      trueColoc5 = colocPerc >= 5,
      posObjs = sum(coloc > 1),
      truePos1 = posObjs >= 1,
      truePos5 = posObjs >= 5,
    )
  coloc_sumstats <- rbind(coloc_sumstats, sumstats)
}

head(coloc_sumstats)
```

```
## # A tibble: 6 x 9
##   Label1      filename      n coloc~1 trueC~2 trueC~3 posObjs trueP~4 trueP~5
##   <chr>      <chr>      <int> <dbl> <lgl> <lgl> <int> <lgl> <lgl>
## 1 obj376-val1 C_M-6-40x-8~ 4744 6.17 TRUE TRUE 23 TRUE TRUE
## 2 obj377-val2 C_M-6-40x-8~ 4744 3.58 TRUE FALSE 12 TRUE TRUE
## 3 obj325-val1 C_M-6-40x-8~ 663 0.637 FALSE FALSE 3 TRUE FALSE
## 4 obj326-val4 C_M-6-40x-8~ 663 78.7 TRUE TRUE 1 TRUE FALSE
## 5 obj327-val5 C_M-6-40x-8~ 663 1.08 TRUE FALSE 11 TRUE TRUE
## 6 obj329-val7 C_M-6-40x-8~ 663 12.3 TRUE TRUE 4 TRUE FALSE
## # ... with abbreviated variable names 1: colocPerc, 2: trueColoc1,
## # 3: trueColoc5, 4: truePos1, 5: truePos5
```

```
coloc_stats <- coloc_sumstats %>% group_by(filename) %>% summarise(n = n(),
  ">1% coloc" = sum(trueColoc1)/n*100,
  ">5% coloc" = sum(trueColoc5)/n*100,
  minPerc = min(colocPerc),
  meanPerc = mean(colocPerc),
  maxPerc = max(colocPerc),
  ">1dot" = sum(truePos1)/n*100,
  ">5dots" = sum(truePos5)/n*100,)
kable(coloc_stats)
```

filename	n	>1% coloc	>5% coloc	minPerc	meanPerc	maxPerc	>1dot	>5dots
C_M-6-40x-8avg-1_cre-gfp-colocalization.csv	2	100.00000	50.00000	3.5785137	4.8746999	6.1708861	100	100.00000
C_M-6-40x-8avg-2_cre-gfp-colocalization-fixed.csv	4	75.00000	50.00000	0.6374881	23.163909	78.6729363	100	25.00000
C_M-6-40x-8avg-3_cre-gfp-colocalization-fixed.csv	3	66.66667	66.66667	0.1099592	33.907244	87.3190302	100	33.33333
C_M-6-40x-8avg-4_cre-gfp-colocalization-fixed.csv	3	100.00000	100.00000	12.348668	331.444039	652.9235382	100	33.33333
C_M-6-40x-8avg-5_cre-gfp-colocalization-fixed.csv	1	0.00000	0.00000	0.0000000	0.0000000	0.0000000	0	0.00000
C_wm-40x-8avg-1_cre-gfp-colocalization.csv	5	100.00000	100.00000	9.8334474	20.004917	832.8857101	100	100.00000
C_wm-40x-8avg-5_cre-gfp-colocalization-fixed.csv	2	100.00000	50.00000	3.0524605	9.7003751	16.3482896	100	50.00000
C_wm-40x-8avg-6_cre-gfp-colocalization.csv	3	100.00000	66.66667	3.0672556	5.6041485	7.6546925	100	100.00000
C_wm-40x-8avg-7_cre-gfp-colocalization.csv	2	100.00000	100.00000	31.374281	1945.260005	359.1457286	100	50.00000
C_wm-40x-8avg-8_cre-gfp-colocalization.csv	1	0.00000	0.00000	0.0770416	0.0770416	0.0770416	100	0.00000

```
coloc5_stats <- coloc_sumstats %>% group_by(filename) %>%
  filter(colocPerc > 5) %>%
  summarise(n = n(),
    ">5% coloc" = sum(trueColoc5)/n*100,
    minPerc = min(colocPerc),
    meanPerc = mean(colocPerc),
    maxPerc = max(colocPerc))
kable(coloc5_stats)
```

filename	n	>5% coloc	minPerc	meanPerc	maxPerc
C_M-6-40x-8avg-1_cre-gfp-colocalization.csv	1	100	6.170886	6.170886	6.170886
C_M-6-40x-8avg-2_cre-gfp-colocalization-fixed.csv	2	100	12.268188	45.470562	78.672936
C_M-6-40x-8avg-3_cre-gfp-colocalization-fixed.csv	2	100	14.292743	50.805887	87.319030

filename	n	>5% coloc	minPerc	meanPerc	maxPerc
C_M-6-40x-8avg-4_cre-gfp-colocalization-fixed.csv	3	100	12.348668	31.444040	52.923538
C_wm-40x-8avg-1_cre-gfp-colocalization.csv	5	100	9.833447	20.004918	32.885710
C_wm-40x-8avg-5_cre-gfp-colocalization-fixed.csv	1	100	16.348290	16.348290	16.348290
C_wm-40x-8avg-6_cre-gfp-colocalization.csv	2	100	6.090497	6.872595	7.654693
C_wm-40x-8avg-7_cre-gfp-colocalization.csv	2	100	31.374282	45.260005	59.145729

```

mouse_coloc_sumstats <- coloc_sumstats %>%
  mutate(mouse = str_extract(filename, "(?<=C_).*(?=-[:digit:]_)")) %>%
  filter(colocPerc > 5) %>%
  group_by(mouse) %>%
  summarise(
    n = n(),
    minPerc = min(colocPerc),
    meanPerc = mean(colocPerc),
    maxPerc = max(colocPerc)
  )
mouse_coloc_sumstats

```

```

## # A tibble: 2 x 5
##   mouse          n minPerc meanPerc maxPerc
##   <chr>      <int>   <dbl>   <dbl>   <dbl>
## 1 M-6-40x-8avg     8    6.17    36.6    87.3
## 2 wm-40x-8avg    10    6.09    22.1    59.1

```

```

overall_coloc_sumstats <- coloc_sumstats %>%
  filter(colocPerc > 5) %>%
  summarise(
    n = n(),
    minPerc = min(colocPerc),
    meanPerc = mean(colocPerc),
    maxPerc = max(colocPerc)
  )
overall_coloc_sumstats

```

```

## # A tibble: 1 x 4
##       n minPerc meanPerc maxPerc
##   <int>   <dbl>   <dbl>   <dbl>
## 1    18    6.09    28.5    87.3

```