

Mængde

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```
library(devtools)
```

```
## Loading required package: usethis
```

```
install_github('legendenomgeorg/VitalDBR/VitalDBR', force=TRUE)
```

```
## Downloading GitHub repo legendenomgeorg/VitalDBR@HEAD
```

```
## * checking for file '/private/var/folders/81/jf2m8tyj02q6bhybvrfzfw_c0000gn/T/Rtmpj0Z8m0/remotes10bf
```

```
## * preparing 'VitalDBR':
```

```
## * checking DESCRIPTION meta-information ... OK
```

```
## * checking for LF line-endings in source and make files and shell scripts
```

```
## * checking for empty or unneeded directories
```

```
## Omitted 'LazyData' from DESCRIPTION
```

```
## * building 'VitalDBR_0.1.0.tar.gz'
```

```
library(VitalDBR)
```

```
library("tidyverse")
```

```
## -- Attaching packages ----- tidyverse 1.3.1 --
```

```
## v ggplot2 3.3.5      v purrr  0.3.4
```

```
## v tibble  3.1.6      v dplyr  1.0.8
```

```
## v tidyr   1.2.0      v stringr 1.4.0
```

```
## v readr   2.1.2      v forcats 0.5.1
```

```
## -- Conflicts ----- tidyverse_conflicts() --
```

```
## x dplyr::filter() masks stats::filter()
```

```
## x dplyr::lag()     masks stats::lag()
```

```
library(readr)
```

Først har vi alle cases hvor de generelle ting er opfyldt:

```
cases <- VitalDBR::load_VDB("https://api.vitaldb.net/cases") %>%
```

```
  dplyr::filter(approach=="Open",
```

```
                department=="General surgery",
```

```
                ane_type=="General") %>%
```

```
  dplyr::select(caseid, death_inhosp, icu_days, age, sex, asa, emop, bmi)
```

```
head(cases)
```

```
##   caseid death_inhosp icu_days age sex asa emop  bmi
## 1      1           0         0  77  M   2   0  26.3
## 2      2           0         0  54  M   2   0  19.6
## 3      5           0        13  66  M   3   1  20.4
## 4      8           0         0  81  F   2   0  27.4
## 5     12           0         4  46  F   2   0  28.4
```

```
## 6      16      0      0 57  M   2      0 26.2
```

```
nrow(cases)
```

```
## [1] 2805
```

Der er cirka 2800 rows

```
death <- cases %>% dplyr::filter(death_inhosp==1)
head(death)
```

```
##   caseid death_inhosp icu_days age sex asa emop  bmi
## 1    410           1         0  64  M   2    0 21.2
## 2    465           1         0  42  M   1    0 20.3
## 3    494           1         0  54  F   1    0 17.2
## 4   1927           1         0  45  F   2    0 22.5
## 5   2035           1         0  71  M   3    0 21.4
## 6   2571           1         0  27  M   2    0 22.6
```

```
nrow(death)
```

```
## [1] 33
```

Og kun cirka 33 der er døde

Derefter har vi alle hvor det kun er art

```
only_art <- VitalDBR::load_VDB("https://api.vitaldb.net/trks") %>%
  dplyr::filter(tname == "SNUADC/ART")
head(only_art)
```

```
##   caseid      tname      tid
## 1      1 SNUADC/ART 724cdd7184d7886b8f7de091c5b135bd01949959
## 2      3 SNUADC/ART 54be86b8ba1eb9c42a572b5824310eb6c9f19aef
## 3      4 SNUADC/ART e28777c4706fe3a5e714bf2d91821d22d782d802
## 4      7 SNUADC/ART a693647dfcc9b1ba19042b05de10276bd4f3d839
## 5     10 SNUADC/ART 58d78f57ffad2ce71fea9cb8e9df2e8fdeeb7556
## 6     12 SNUADC/ART 548e82b021822840d9af99893651cfcfea198510
```

```
nrow(only_art)
```

```
## [1] 3645
```

Og her jointet på det originale table

```
merged <- merge(x=only_art,y=cases,by="caseid")
death_merged <- merged %>% dplyr::filter(death_inhosp==1)
head(death_merged)
```

```
##   caseid      tname      tid death_inhosp
## 1   3101 SNUADC/ART 0a6f12aed42abad3b0f3600e482fe11f2f6ea23f          1
## 2   3136 SNUADC/ART c63fd71b67646570bc700a6c0acb65779adc1ba3          1
## 3   3230 SNUADC/ART b0fb3dca1d79917454edcb5e0bf6b4c241ac6c22          1
## 4   3407 SNUADC/ART 1c6005de5881d6b88a4023e69ff8c3af3faf4d9b          1
## 5   3486 SNUADC/ART 8e56dd60d77c97dbc72580a3fd43404648c614be          1
## 6   3527 SNUADC/ART 75a72a4ec66dbd1d6e7c2767aff7c92ca1651d7c          1
##   icu_days  age sex  asa emop  bmi
## 1      1 22.0  M   NA    1 29.5
## 2      5 81.0  F    3    1 20.5
## 3     38  0.7  F    4    1 21.7
## 4      0 60.0  F    2    1 21.4
```

```
## 5      38 0.7 F 4 1 21.7
## 6      25 82.0 F 4 1 26.7
```

```
nrow(death_merged)
```

```
## [1] 23
```

Hvor der både er awp og art

```
both <- VitalDBR::load_VDB("https://api.vitaldb.net/trks") %>%
  dplyr::filter(tname == "Primus/AWP" | tname == "SNUADC/ART") %>%
  count(caseid) %>%
  dplyr::filter(n == 2)
head(both)
```

```
##   caseid n
## 1      1 2
## 2      3 2
## 3      4 2
## 4      7 2
## 5     10 2
## 6     12 2
```

```
nrow(both)
```

```
## [1] 3629
```

Joinet på det originale og filtreret udfra om de er døde

```
merged <- merge(x=both,y=cases,by="caseid") %>% dplyr::select(-one_of("n"))
merged_both <- merged %>% dplyr::filter(death_inhosp==1)
head(merged_both)
```

```
##   caseid death_inhosp icu_days age sex asa emop bmi
## 1   3101           1         1 22.0 M  NA   1 29.5
## 2   3136           1         5 81.0 F   3   1 20.5
## 3   3230           1        38 0.7 F   4   1 21.7
## 4   3407           1         0 60.0 F   2   1 21.4
## 5   3486           1        38 0.7 F   4   1 21.7
## 6   3527           1        25 82.0 F   4   1 26.7
```

```
nrow(merged_both)
```

```
## [1] 23
```