Project tutoré 2 - Murat Simsek et Yuquan Dai

R pour la question 1 et Python pour la question 2

```
# Importer les données nécessaires
dm1 <- read.csv2("dm1.csv", header = TRUE, sep = ",")</pre>
dm2 <- read.csv2("dm2.csv", header = TRUE, sep = ",")</pre>
ie1 <- read.csv2("ie1.csv", header = TRUE, sep = ",")</pre>
ie2 <- read.csv2("ie2.csv", header = TRUE, sep = ",")</pre>
lb1 <- read.csv2("lb1.csv", header = TRUE, sep = ",")</pre>
1b2 <- read.csv2("lb2.csv", header = TRUE, sep = ",")</pre>
# Jointure les données
alldm <- full_join(dm1,dm2)</pre>
## Joining, by = c("STUDYID", "DOMAIN", "USUBJID", "EPOCH", "VISIT", "VISITNUM",
## "RFSTDTC", "RFENDTC", "SITEID", "BRTHDTC", "AGE", "AGEU", "SEX", "RACE",
## "ETHNIC", "ARMCD", "ARM", "COUNTRY", "DMDTC", "DMDY")
allie <- full_join(ie1,ie2)
## Joining, by = c("STUDYID", "DOMAIN", "USUBJID", "EPOCH", "IESEQ", "IETESTCD",
## "IETEST", "IECAT", "IEORRES", "IESTRESC", "VISIT", "VISITNUM", "IEDTC", "IEDY")
alllb <- full_join(lb1,lb2)
## Joining, by = c("STUDYID", "DOMAIN", "USUBJID", "EPOCH", "LBSEQ", "LBTESTCD",
## "LBTEST", "LBCAT", "LBORRES", "LBORRESU", "LBORNRLO", "LBORNRHI", "LBSTRESC",
## "LBSTRESN", "LBSTRESU", "LBSTNRLO", "LBSTNRHI", "LBNRIND", "LBSTAT",
## "LBREASND", "LBSPEC", "LBMETHOD", "LBBLFL", "VISIT", "VISITNUM", "LBDTC",
## "LBDY")
# On a choisit dm pour travailler
n_distinct(alldm$USUBJID)
## [1] 411
# Faire une exclusion spéciale
allusrdm <- alldm$USUBJID[!(alldm$USUBJID %in% allie$USUBJID[allie$IECAT=="EXCLUSION"])]
summary(allusrdm)
##
      Length
                 Class
##
         369 character character
```

```
# Séparer BUPRENORPHINE/NALOXONE et CLONIDINE
allusrBN <- filter(alldm,
                   alldm$ARM=="BUPRENORPHINE/NALOXONE" &
                   alldm$USUBJID %in% allusrdm)
allusrC <- filter(alldm,</pre>
                  alldm$ARM=="CLONIDINE" &
                  alldm$USUBJID %in% allusrdm)
n_distinct(allusrBN$USUBJID)
## [1] 233
n_distinct(allusrC$USUBJID)
## [1] 110
\# On considérera les patients qui ont des résultats au tests urinaires à J13 ou J14
usrBNfin <- filter(alllb,
                   (alllb$VISITNUM==13 | alllb$VISITNUM==14)&
                     (alllb$LBSTAT!="NOT DONE")&
                     (alllb$USUBJID %in% allusrBN$USUBJID))
usrCfin <- filter(all1b,
                  (alllb$VISITNUM==13 | alllb$VISITNUM==14)&
                    (alllb$LBSTAT!="NOT DONE")&
                    (alllb$USUBJID %in% allusrC$USUBJID))
n_distinct(usrBNfin$USUBJID)
## [1] 150
n_distinct(usrCfin$USUBJID)
```

[1] 28