# Reproducible code for manuscript figure 03 – Input data

### J.A. Torres-Matallana

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```
# organization 1: Luxembourg Institute of Science and Technology (LIST), Belvaux, Luxembourg
# organization 2: Wagenigen University and Research Centre (WUR), Wageningen, The Netherlands
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```

### Compile Rmarkdown file

```
library("rmarkdown")
rmarkdown::render("Main_reproducible_input_data.R")
```

#### Setup

```
library(knitr)
library(lmom)
library(tikzDevice)

Sys.setenv("LANGUAGE"="En")
Sys.setlocale("LC_ALL", "en_GB.UTF-8")
```

```
## [1] "LC_CTYPE=en_GB.UTF-8;LC_NUMERIC=C;LC_TIME=en_GB.UTF-8;LC_COLLATE=en_GB.UTF-8;LC_MONETARY=en_GB.Ttiming.ini <- Sys.time()
```

#### Plot functions

```
## final plot CODs
load("data/data_DWF.RData")
data <- sort(log(data[,"CODs"])) #log(CODs)

lmom <- samlmu(data, nmom=5)
parnor <- pelnor(lmom)

xmin <- min(data)
xmax <- max(data)

sfin <- seq(xmin, xmax, length.out=91)

# pdf(width = 8, height = 15, file= "output/fig03_notikz.pdf")
tikz(width = 3.5, height = 9, file= "output/fig03_canvas.tex")</pre>
```

```
par(mfrow = c(3,1))
par(mar = c(5, 5, 7, 3), cex.axis=1.4, cex.lab=1.4, cex.main=1.4) # bottom, left, top and right margins
xlab1 <- "CODs [g/(PE d)]"</pre>
xlab <- "log(CODs [g/(PE d)])"</pre>
hist(data, breaks=50, freq = F, ylim=c(0,1.5), main = "", font.main=1,
     xlab = xlab)
lines(density(data),col="red",lwd=2, lty="dashed")
cdf <- cdfnor(x =sfin, para = parnor)</pre>
diff_cdf <- diff(cdf)</pre>
diff_sfin <- diff(sfin)</pre>
df.CODs <- diff_cdf/diff_sfin</pre>
lines(sfin[1:90], df.CODs,col="blue",lwd=2)
xmin <- round(xmin, digits=0); xmax <- round(xmax, digits=0)</pre>
axis(side=3, at=seq(xmin,xmax,1), labels=round(exp(seq(xmin,xmax,1)), digits=0))
mtext(xlab1, side=3, line=3, cex.lab=1,las=1, col="black")
## final plot NH4s
load("data/data DWF.RData")
data <- sort(log(data[,"NH4s"])) #log(NH4s)</pre>
lmom <- samlmu(data, nmom=5)</pre>
parnor <- pelnor(lmom)</pre>
xmin <- min(data)</pre>
xmax <- max(data)</pre>
sfin <- seq(xmin, xmax, length.out=91)</pre>
par(mar = c(5, 5, 9, 3), cex.axis=1.4, cex.lab=1.4, cex.main=1.4) # bottom, left, top and right margins
xlab1 \leftarrow "NH4s [g/(PE d)]"
xlab <- "log(NH4s [g/(PE d)])"</pre>
hist(data, breaks=50, freq = F, ylim=c(0,2.0), xlim=c(0.5,2.5), main = "",
     xlab = xlab)
lines(density(data),col="red",lwd=2, lty="dashed")
cdf <- cdfnor(x =sfin, para = parnor)</pre>
diff_cdf <- diff(cdf)</pre>
diff_sfin <- diff(sfin)</pre>
df.NH4s <- diff_cdf/diff_sfin</pre>
lines(sfin[1:90], df.NH4s,col="blue",lwd=2)
xmin <- round(xmin, digits=1); xmax <- round(xmax+.1, digits=1)</pre>
axis(side=3, at=seq(xmin,xmax,.5), labels=round(exp(seq(xmin,xmax,.5)), digits=1))
mtext(xlab1, side=3, line=3, cex.lab=1,las=1, col="black")
## final plot CODr
## CODr values literature (A+A) Mean: 71 mg/l; Std dev.: 150 mg/l
logCODr \leftarrow rnorm(1e7, 3.40, 1.31)
```

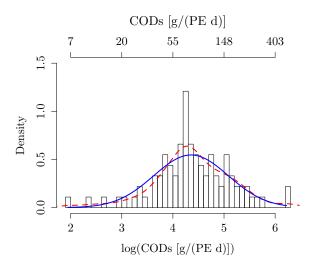
```
CODr <- exp(logCODr)
mean(CODr)
## [1] 70.72822
sd(CODr)
## [1] 151.322
data <- log(CODr)</pre>
lmom <- samlmu(data, nmom=5)</pre>
parnor <- pelnor(lmom)</pre>
xmin <- min(data)</pre>
xmax <- max(data)</pre>
sfin <- seq(xmin, xmax, length.out=91)
par(mar = c(5, 5, 9, 3), cex.axis=1.4, cex.lab=1.4, cex.main=1.4) # bottom, left, top and right margins
xlab1 <- "CODr [mg/1]"</pre>
xlab <- "log(CODr [mg/l])"</pre>
hist(data, breaks=50, freq = F, ylim=c(0,.7), xlim=c(0,6.5),
     main = "", xlab = xlab, border = "white")
# lines(density(data),col="red",lwd=2, lty="dashed")
cdf <- cdfnor(x =sfin, para = parnor)</pre>
diff_cdf <- diff(cdf)</pre>
diff_sfin <- diff(sfin)</pre>
df.CODr <- diff_cdf/diff_sfin</pre>
lines(sfin[1:length(df.CODr)], df.CODr,col="blue",lwd=2)
xmin <- round(xmin, digits=0); xmax <- round(xmax, digits=0)</pre>
x.at <- seq(xmin,xmax,1)</pre>
axis(side=3, at=x.at, labels=round(exp(x.at), digits=0))
mtext(xlab1, side=3, line=3, cex.lab=1,las=1, col="black")
dev.off()
## pdf
##
Render latex file to pdf
```

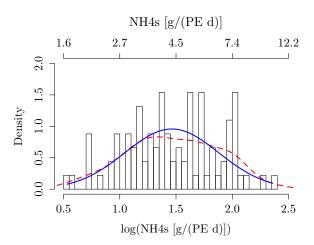
## Include pdf

### **Timing**

```
timing.end <- Sys.time()
(timing.elapsed <- timing.end - timing.ini)</pre>
```

## Time difference of 4.106522 secs





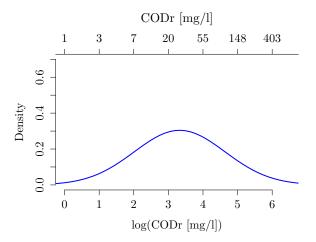


Figure 1: Figure 03.