

# Cyclothone example

This is an example of how to use KRMr.

This is not an official release and is still in experimental phase.

## Reading the shape information

Cyclothone used here are non-swimbladdered fish. We have to shape files, one for the fish body and one for the swimbladder.

The shape file for the fish body contains one column named **x\_fb** the fishbody (fb) along the x axis (Length), **w\_fb** the width (seen from top) of the fish body at each position x (along the fish body length), **z\_fbL** the lower height of the fish body along the x axis (lower (L) fish body (fb) extend in direction z) and **z\_fbU** the upper height along the x axis (upper (U) fish body (fb) extend in direction z).

Analogously, the swimbladder (sb) shape file contains **x\_sb** - x axis (Length direction), **w\_sb** - Width of the swimbladder along the x axis, **z\_sbL** - lower swimbladder extend and **z\_sbU** - upper swimbladder extend.

```
library(dplyr)

##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##   filter, lag

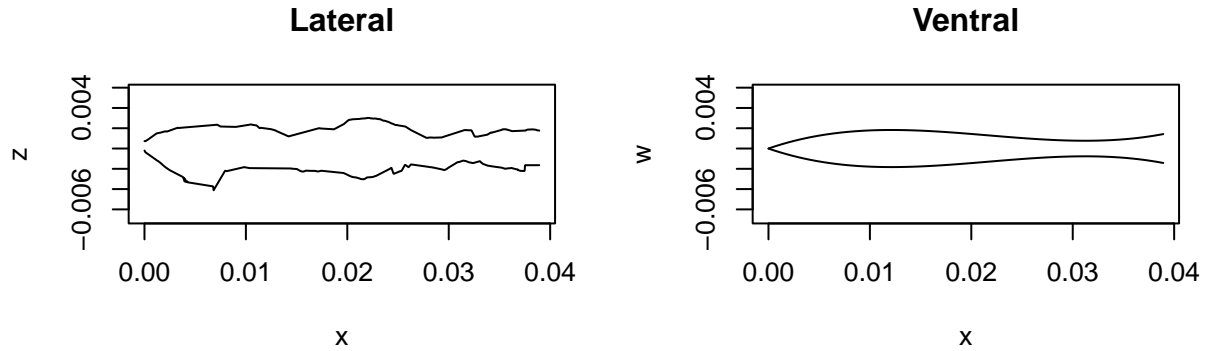
## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union

library(KRMr)
stb_fb <- read.csv(paste0(dirname(getwd()), '/data/fb1contour.csv'))
```

Let's have a look at the shapes we loaded.

To plot the loaded shapes, we can use the `shplot(x_fb, w_fb, x_sb, w_sb, z_fbU, z_fbL, z_sbU, z_sbL)` function:

```
fb =stb_fb
layout(matrix(c(1,2,3,3), 2, 2, byrow = TRUE))
KRMr::shplot(x_fb = fb$x_fb, w_fb = fb$w_fb,
             x_sb = NA, w_sb = NA,
             z_fbU = fb$z_fbU, z_fbL = fb$z_fbL,
             z_sbU = NA, z_sbL = NA)
```



## Running a KRM simulation

Now that we have our shapes sorted, let's have a go at a KRM simulation.

We want to get TS at 38, 70, 120 and 200 kHz, with the below defined settings:

- c.w = ambient water sound speed
- rho.w = density ambient water
  
- theta = orientation in degrees
- c.fb = sound speed inside fish body
- c.sb = sound speed inside swimbladder
- rho.sb = density inside swimbladder
- rho.fb = density inside fish body = L = length of the fishbody in m
- x\_fb = Fish body coordinates along x axis
- x\_sb = Swimbladder coordinates along x axis
- w\_fb = Fish body width along x axis
- w\_sb = Swimbladder width coordinates along x axis
- z\_fbU = Upper height of the fish body along the x axis
- z\_fbL = Lower height of the fish body along the x axis
- z\_sbU = Upper height of the swimbladder along the x axis
- z\_sbL = Lower height of the swimbladder along the x axis

A description of the parameters is also available in the help files `?KRMr::krm` (runs a single krm simulation) or `?KRMr::krm.sim` (runs multiple simulations if any input parameter contains more than one value (except for coordinates))

```

TS = KRMr::krm.sim(frequency =c(38,70,120,200) * 1000,
                  c.w = 1490,
                  rho.w = 1030,
                  theta=90,
                  c.fb = 1570,
                  c.sb = 345,
                  rho.sb = 1.24,
                  rho.fb = 1070,
                  L=0.25,
                  x_fb = fb$x_fb,
                  x_sb = NULL,
                  w_fb = fb$w_fb,
                  w_sb = NULL,
                  z_fbU = fb$z_fbU,
                  z_fbL = fb$z_fbL,
                  z_sbU = NULL,
                  z_sbL = NULL)

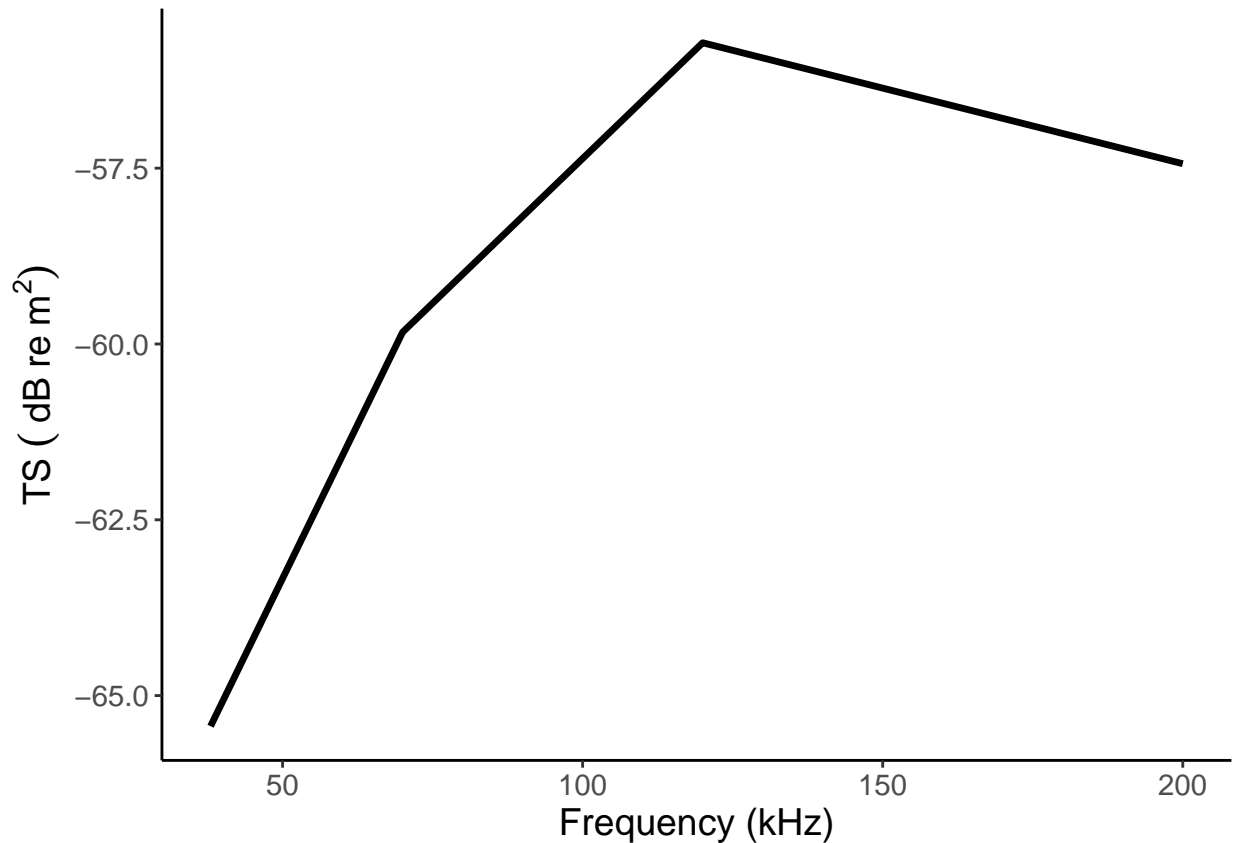
```

Let's have a look at the outcome:

```

library(ggplot2)
ggplot(data=TS,
       aes(x=frequency/1000, y=TS))+
  geom_line(size=1.2)+
  ylab(expression(TS~(~dB~re~m^2)))+
  xlab('Frequency (kHz)')+
  theme_classic()+
  theme(text=element_text(size=14),
        legend.position='top')

```



We can also compute slightly more sophisticated simulations of a range of frequencies, lengths and orientations (remember the KRM model is only valid for orientations, close to 90 degrees ~65-115 degrees):

```
TS = krm.sim(frequency = c(38,70,120,200) * 1000,
             c.w = 1490,
             rho.w = 1030,
             theta=seq(65,115),
             c.fb = 1570,
             c.sb = 345,
             rho.sb = 1.24,
             rho.fb = 1070,
             L=seq(0.015,0.065,by=0.005),
             x_fb = fb$x_fb,
             x_sb = NULL,
             w_fb = fb$w_fb,
             w_sb = NULL,
             z_fbU = fb$z_fbU,
             z_fbL = fb$z_fbL,
             z_sbU = NULL,
             z_sbL = NULL)
```

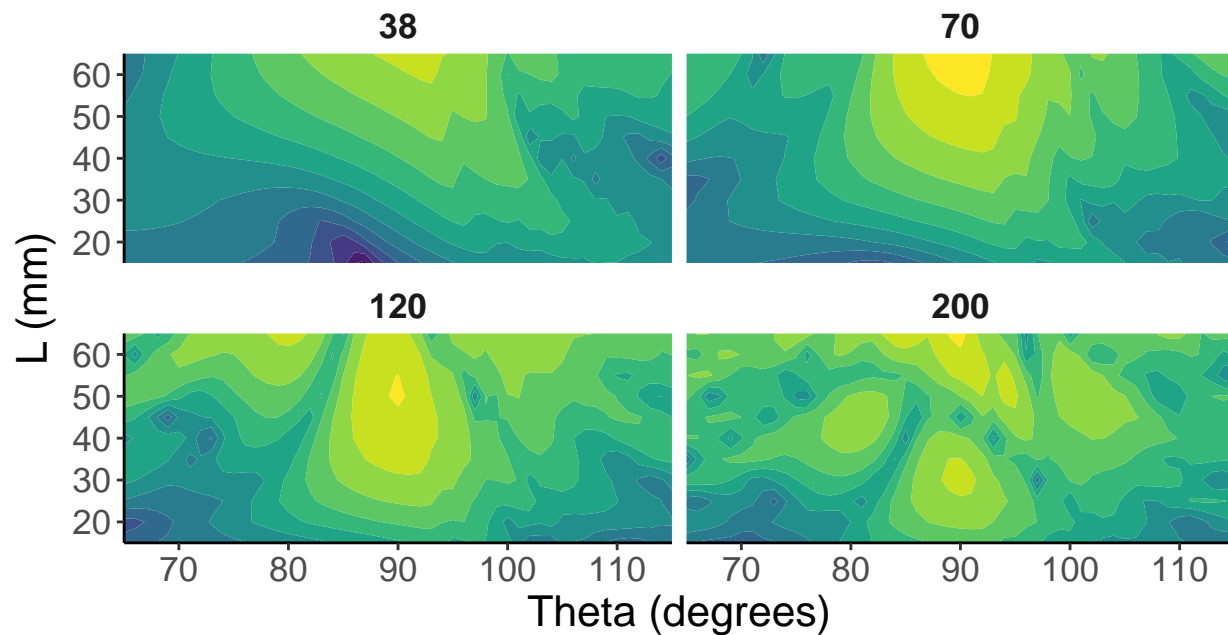
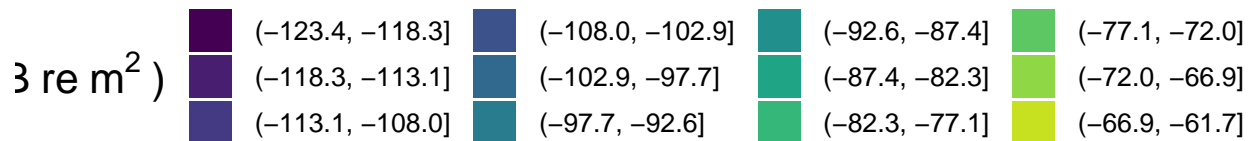
Let's plot the results:

```
ggplot2::ggplot(data=TS, ggplot2::aes(x= theta, y=L*1000 , z=TS))+
  facet_wrap(~frequency/1000)+
  geom_contour_filled(bins=15)+
```

```

ggplot2::scale_fill_viridis_d(expression(TS~('~-dB~re~ m^2~')))+
ggplot2::xlab('Theta (degrees)')+
ggplot2::ylab(expression(L~'(mm)'))+
ggplot2::scale_x_continuous(expand=c(0,0))+
ggplot2::scale_y_continuous(expand=c(0,0))+
ggplot2::theme_classic()+
ggplot2::theme(legend.position='top',
               legend.text=element_text(size=10),
               text=element_text(size=16),
               strip.background = element_blank(),
               strip.text = element_text(face='bold'))

```



Or produce another plot:

```

TS%>%
  mutate(frequency=frequency/1000)%>%
  mutate(sigma=10^(TS/10))%>%
  group_by(frequency,theta) %>%
  summarise(mean.sigma = mean(sigma, na.rm=T),
            sd.sigma = sd(sigma, na.rm=T),
            n.TS = n()) %>%
  mutate(se.sigma = sd.sigma / sqrt(n.TS),
         lower.ci.sigma = mean.sigma - qt(1 - (0.05 / 2), n.TS - 1) * se.sigma,
         upper.ci.sigma = mean.sigma + qt(1 - (0.05 / 2), n.TS - 1) * se.sigma,
         mean.TS=10*log10(mean.sigma),
         sd.TS=10*log10(sd.sigma),

```

```
se.TS=10*log10(se.sigma),
lower.ci.TS=10*log10(lower.ci.sigma),
upper.ci.TS=10*log10(upper.ci.sigma))>TS.ci
```

## 'summarise()' has grouped output by 'frequency'. You can override using the '.groups' argument.

## Warning in mask\$eval\_all\_mutate(quo): NaNs produced

## Warning in mask\$eval\_all\_mutate(quo): NaNs produced

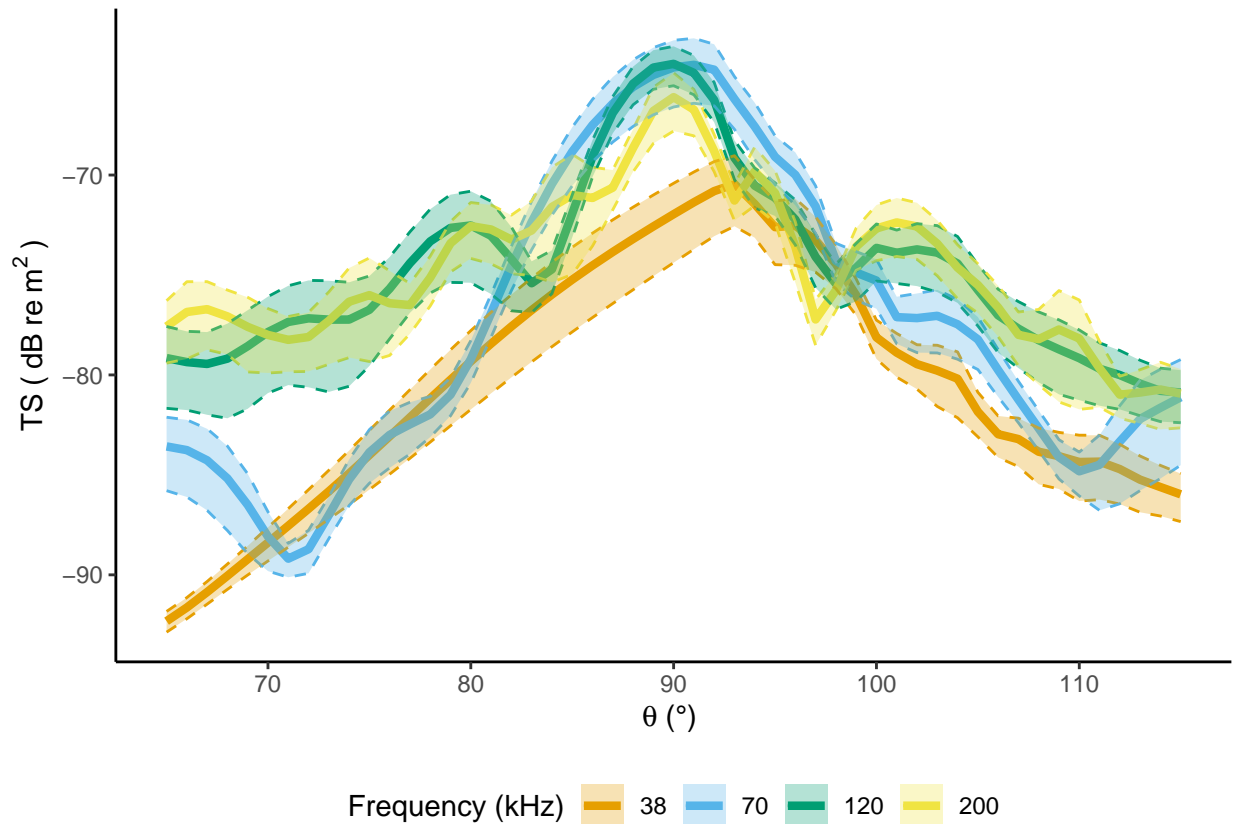
## Warning in mask\$eval\_all\_mutate(quo): NaNs produced

## Warning in mask\$eval\_all\_mutate(quo): NaNs produced

```
#TS.ci=TS
#TS.ci$sigma=10^(TS.ci$TS/10)
okabe <- c("#E69F00", "#56B4E9", "#009E73", "#F0E442", "#0072B2", "#D55E00", "#CC79A7")

ggplot(data=TS.ci, aes(x=theta, group=factor(frequency)))+
  #geom_smooth(aes(x= theta, y=sigma, group=frequency, col=factor(frequency),
  #               method = lm, formula=y~x,
  #               method.args = list(family = gaussian(link = 'log'))), lwd=1.5)

  geom_line(aes(x= theta, y=mean.TS, group=frequency, col=factor(frequency)), lwd=1.5)+
  geom_line(aes(x= theta, y=10*log10(mean.sigma+se.sigma), group=frequency, col=factor(frequency)), lty=2)+
  geom_line(aes(x= theta, y=10*log10(mean.sigma-se.sigma), group=frequency, col=factor(frequency)), lty=2)+
  geom_ribbon(
    aes(ymin=10*log10(mean.sigma-se.sigma),
        ymax=10*log10(mean.sigma+se.sigma), fill=factor(frequency)), alpha=0.3)+
  theme_classic()+
  scale_color_manual(values=okabe, name='Frequency (kHz)')+
  scale_fill_manual(values=okabe, name='Frequency (kHz)')+
  theme(legend.position = 'bottom')+
  ylab(expression(TS~'('~dB~re~m^2~')' ))+
  xlab(expression(paste(theta, ' (\u00B0) ')))
```



We can also look at orientation plots for a fish of 5 cm:

```
TSrot = krm.sim(frequency = c(38,70,120,200) * 1000,
               c.w = 1490,
               rho.w = 1030,
               theta=65:115,
               c.fb = 1570,
               c.sb = 345,
               rho.sb = 1.24,
               rho.fb = 1070,
               L=0.05,
               x_fb = fb$x_fb,
               x_sb = NA,#sb$x_sb,
               w_fb = fb$w_fb,
               w_sb = NA,#sb$w_sb,
               z_fbU = fb$z_fbU,
               z_fbL = fb$z_fbL,
               z_sbU = NA,#sb$z_sbU,
               z_sbL = NA)#sb$z_sbL)
ggplot(TSrot, aes(x = theta, y = TS, group=frequency/1000, col=TS)) +
  geom_path(size=1.2) +
  facet_wrap(~frequency/1000)+
  scale_x_continuous(limits=c(0,360), breaks=c(0,90,180))+
  coord_polar(start=-pi/2,direction=1)+
  scale_colour_viridis_c(name='', limits=c(-80,-70), oob=scales::squish)+
```

```

ylab(expression(TS~'('~dB~re~m^2~')' ))+
xlab(expression(paste(theta,' (\u00B0) ')))+

geom_vline(xintercept=90, lty=2)+
geom_vline(xintercept=0,size=1)+
geom_vline(xintercept=180,size=1)+

theme_classic()+
theme(strip.background = element_blank(),
      strip.text=element_text(size=18),
      legend.position='top',
      legend.text=element_text(angle=-15),
      legend.key.width = unit(2,'cm'),
      axis.line.y = element_blank(),
      axis.text.y = element_blank(),
      axis.ticks.y = element_blank(),
      axis.line.x = element_blank(),
      panel.spacing.y = unit(-5,'lines'),
      axis.title.x = element_text(vjust=27, size=16),
      axis.title.y = element_text(hjust=0.75, size=16),
      text=element_text(size=18))

```



