

# title: Commercial catch and survey data for use in MSY proxies estimation

Razors

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## MYDAS

Task 5 of the MYDAS project <https://github.com/laurieKell/mydas> potentially requires commercial/Survey inputs to estimate the current status of the case-study stocks. The available data sources within the EU are either country specific, or from collective sources such as DATRAS <http://www.ices.dk/marine-data/data-portals/Pages/DATRAS.aspx> for survey data and the STECF <https://stecf.jrc.ec.europa.eu/data-dissemination> for commercial data. As part of tasks 1-2 these data have already been acquired and subset on a stock basis for the MYDAS stocks of interest. Also included are estimates obtained from life-history experiments.

First load the required libraries

```
library(plyr)
library(dplyr)
library(ggplot2)
library(DBI)
library(RPostgreSQL)
library(reshape)
```

## Database connection

```
drv  = dbDriver("PostgreSQL")

con  = dbConnect(drv, host = 'postgresql-seascope.csffkpr9jjjn.eu-west-2.rds.amazonaws.com',
                  dbname = 'mydasDev',
                  port = 5432,
                  user = 'MydasApplication',
                  password = 'gmit2017!')
```

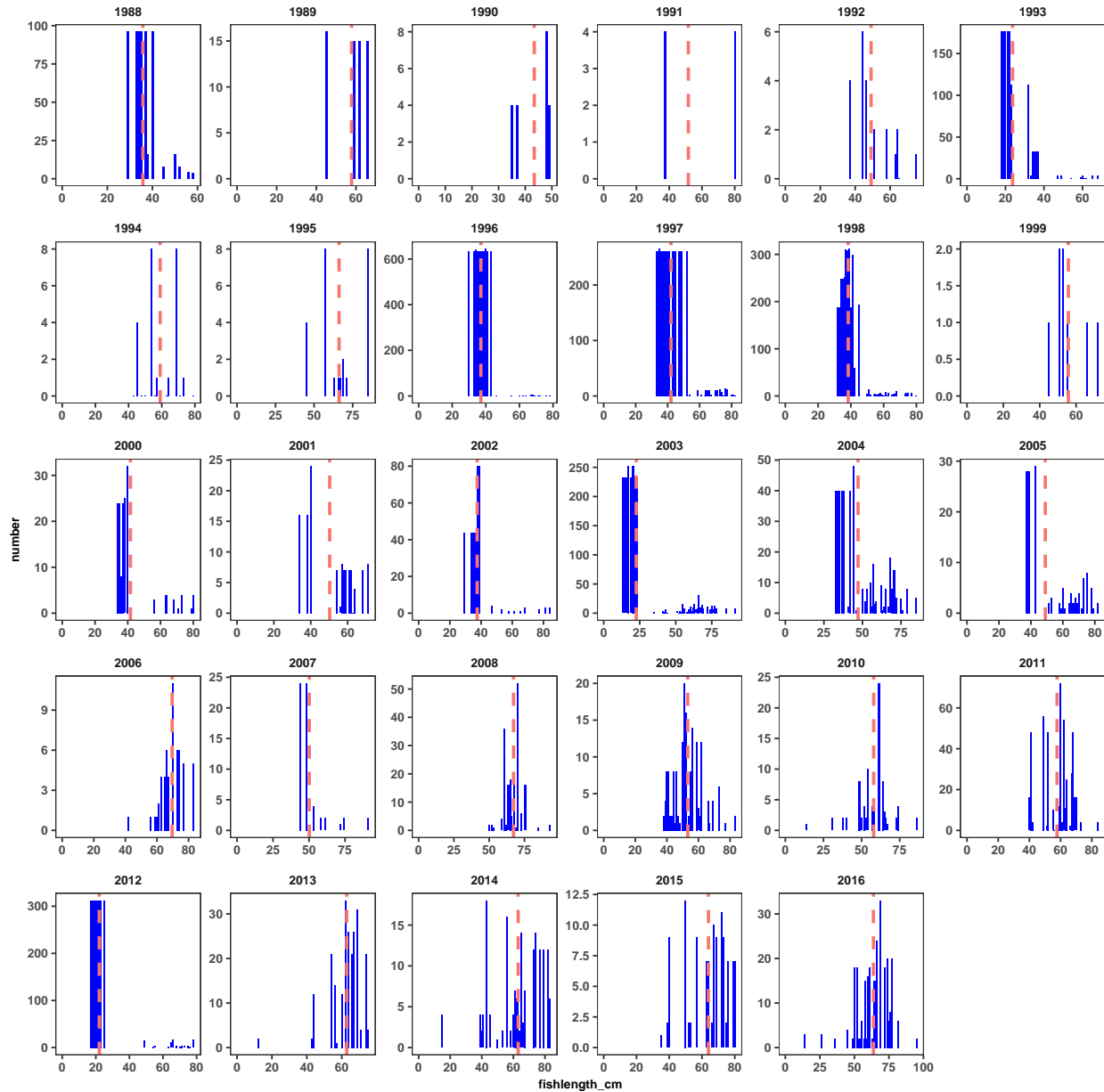
## Survey data extraction

Extract via sql the case study stocks from Mydas survey database and get numbers at length

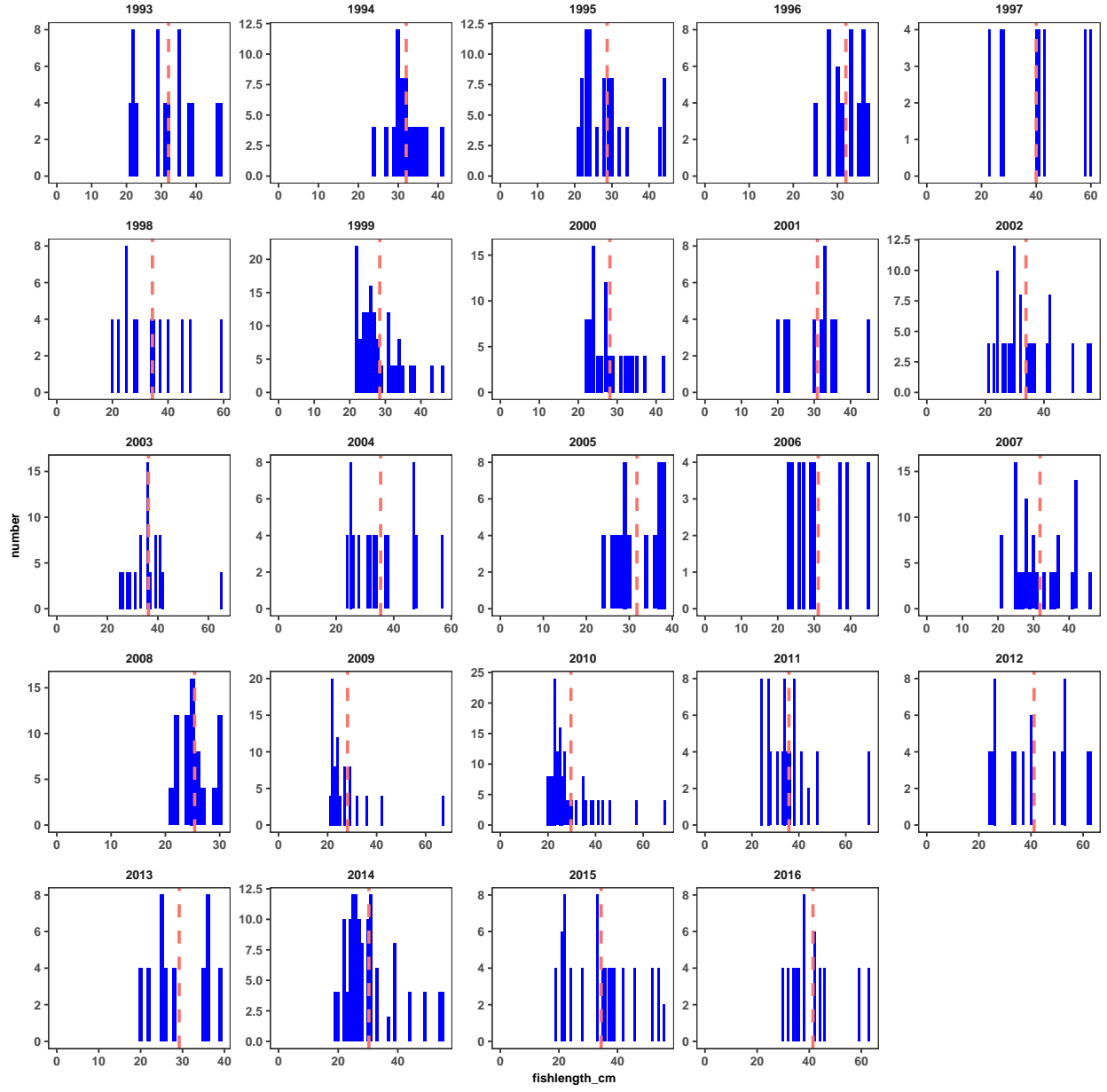
```
bio = dbGetQuery(con, paste0("SELECT year, speciesgp, fishlength_cm, SUM(number*(60/hauldur_min)) AS n
                              FROM data_surveybio
                              WHERE speciesgp in ('TUR','BLL','POL','SKA','SPR')
                              GROUP BY year, speciesgp, fishlength_cm;"))
```

For more in depth detail of the data see <https://data.marine.gov.scot/sites/default/files//SMFS%200816.pdf>  
 Calculate weighted mean length for case study species

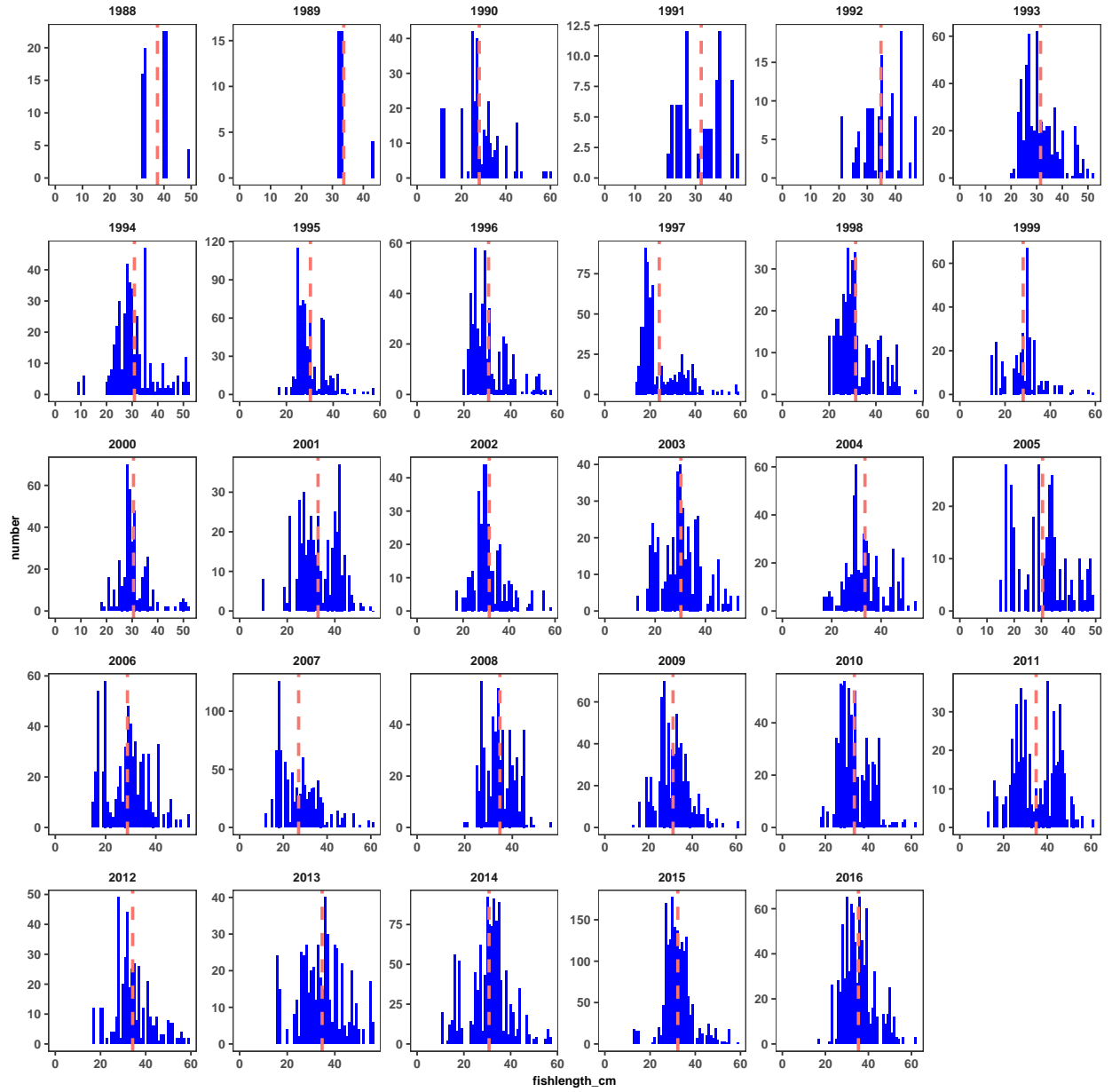
```
lenn=bio%>%
  group_by(year, speciesgp)%>%
  summarise(mnlen= weighted.mean(fishlength_cm,number), totnum=sum(number))
```



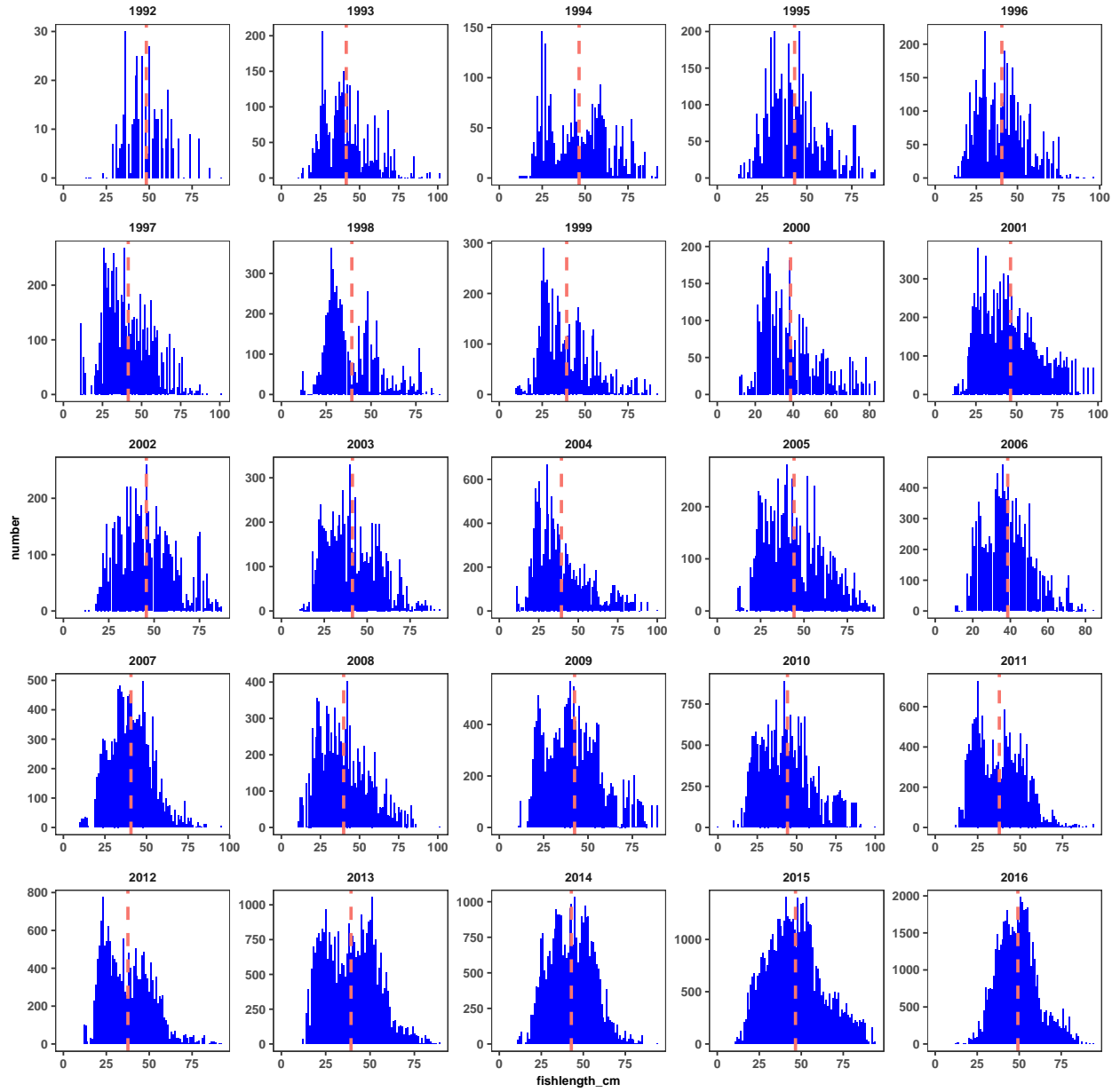
**Figure 1** Plot of numbers at length for pollock with weighted mean length represented by dashed red line.



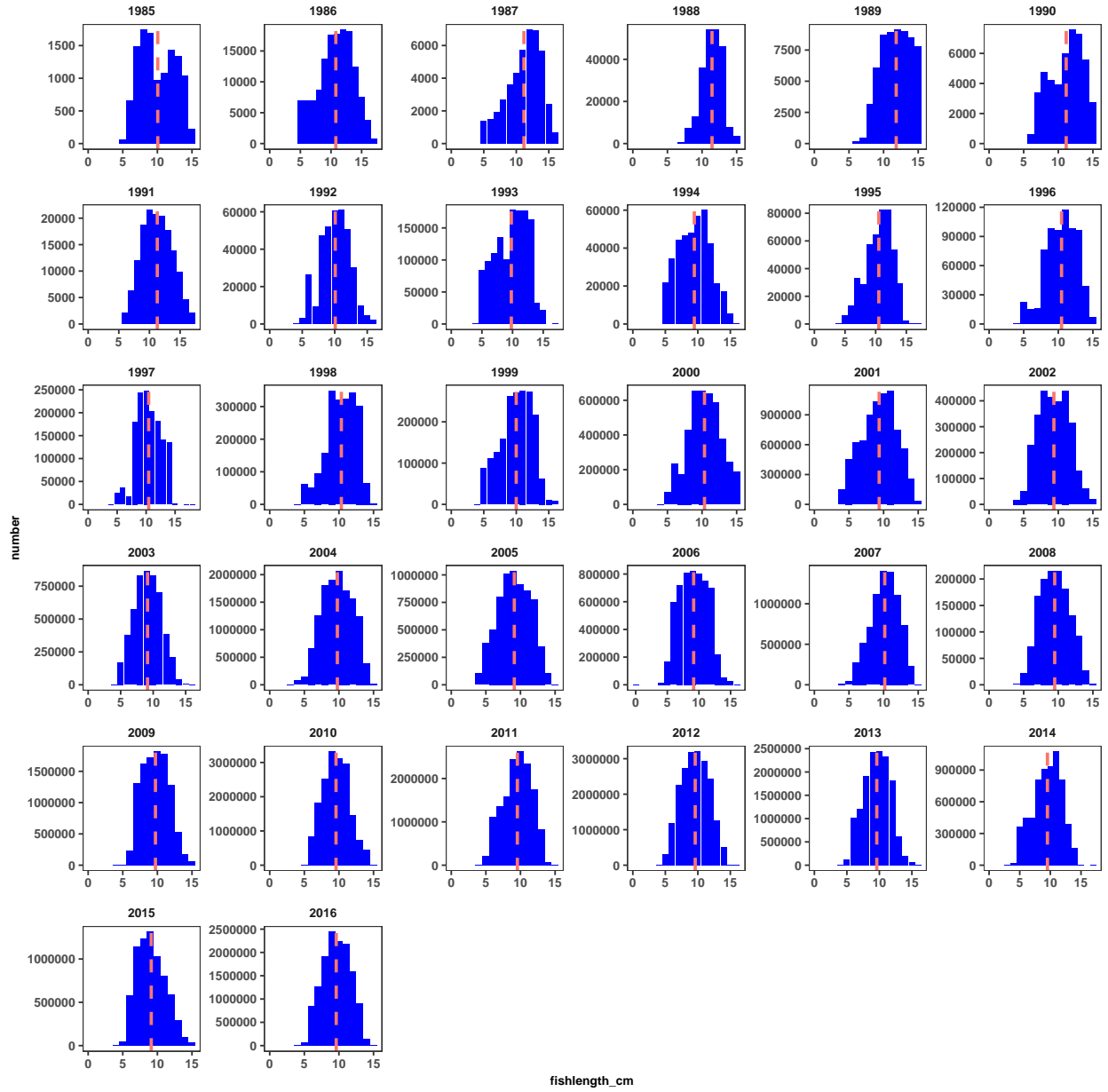
**Figure 2** Plot of numbers at length for turbot with weighted mean length represented by dashed red line.



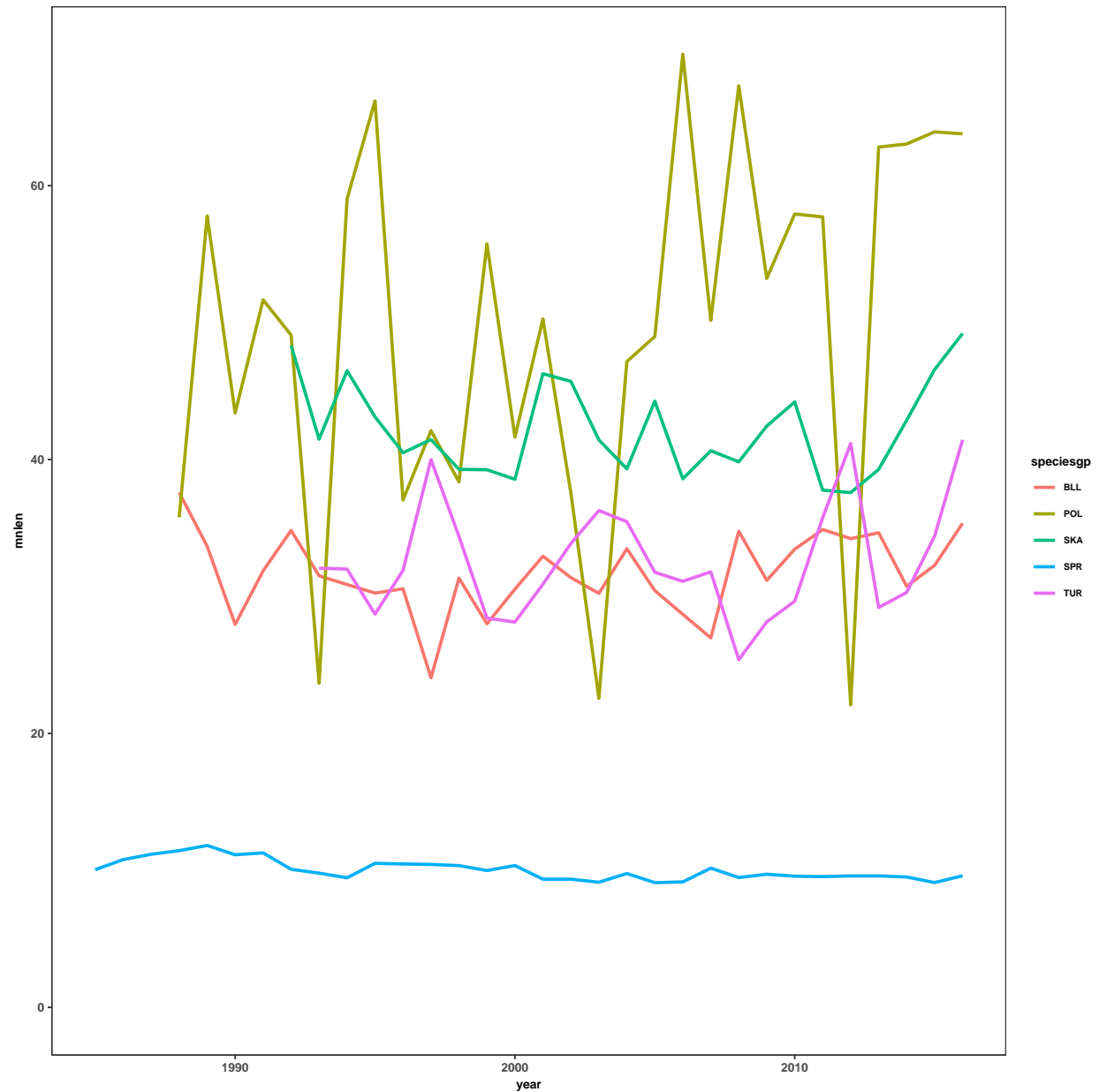
**Figure 3** Plot of numbers at length for brill with weighted mean length represented by dashed red line.



**Figure 4** Plot of numbers at length for skate with weighted mean length represented by dashed red line.



**Figure 5** Plot of numbers at length for sprat with weighted mean length represented by dashed red line.



**Figure 6** Time series of weighted mean length.

Get all survey data from sql query

```
bio1 = dbGetQuery(con, paste0("SELECT *
                                FROM data_surveybio
                                WHERE speciesgp in ('TUR','BLL','POL','SKA','SPR');"))
```

Estimate fish weight in kgs

```
bio1$weight = ((bio1$lwra*bio1$fishlength_cm**bio1$lwrb)/1000)*bio1$number
```

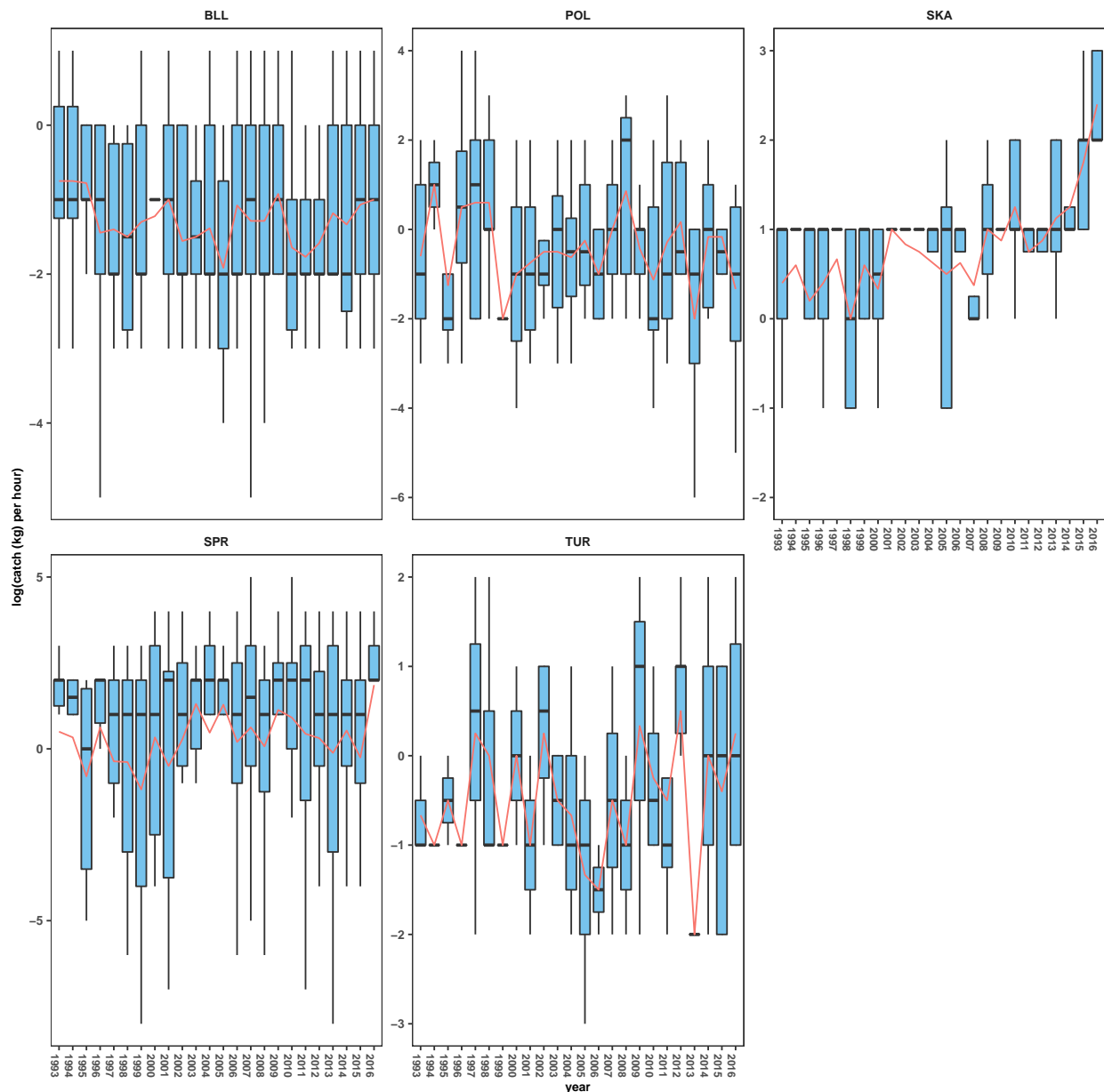
```

bio1$n      = bio1$hauldur_min/60
effort = bio1 %>% group_by(survey_acronym, ices_division, year) %>% summarise(hours=sum(n))

catch=bio1%>%group_by(speciesgp, survey_acronym,ices_division, year) %>% summarise(catchwt=sum(weight))
cpue_surv = effort %>% inner_join(catch) %>% mutate(cpue=round(log(catchwt/hours, 4)))

cpuemn= ddpdy(subset(cpue_surv,year>1992) , .(year, speciesgp), summarise, mncpue=mean(cpue))

```



**Figure 7** Catch per unit effort in kgs (CPUE) time series with red line depicting mean cpue.

Fish numbers

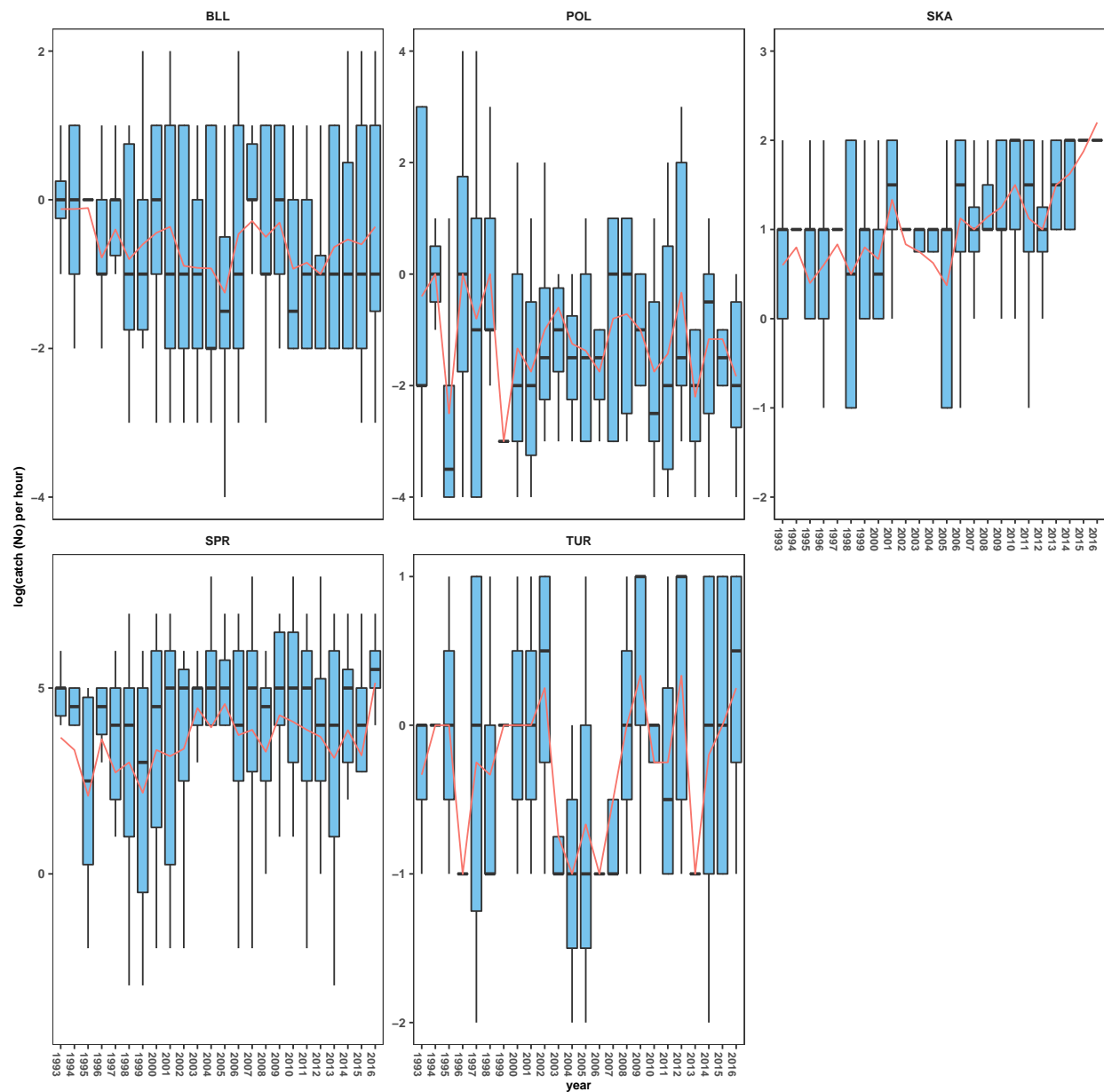
```

catchn=bio1%>%group_by(speciesgp, survey_acronym,ices_division, year) %>% summarise(catchn=sum(number))
cpue_survn = effort %>% inner_join(catchn) %>% mutate(cpue=round(log(catchn/hours, 4)))

```



```
cpuemmn= ddply(subset(cpue_survn, year>1992), .(year, speciesgp), summarise, mncpue=mean(cpue))#
```



**Figure 8** Catch per unit effort in numbers (CPUE) time series with red line depicting mean cpue.

## Commercial data

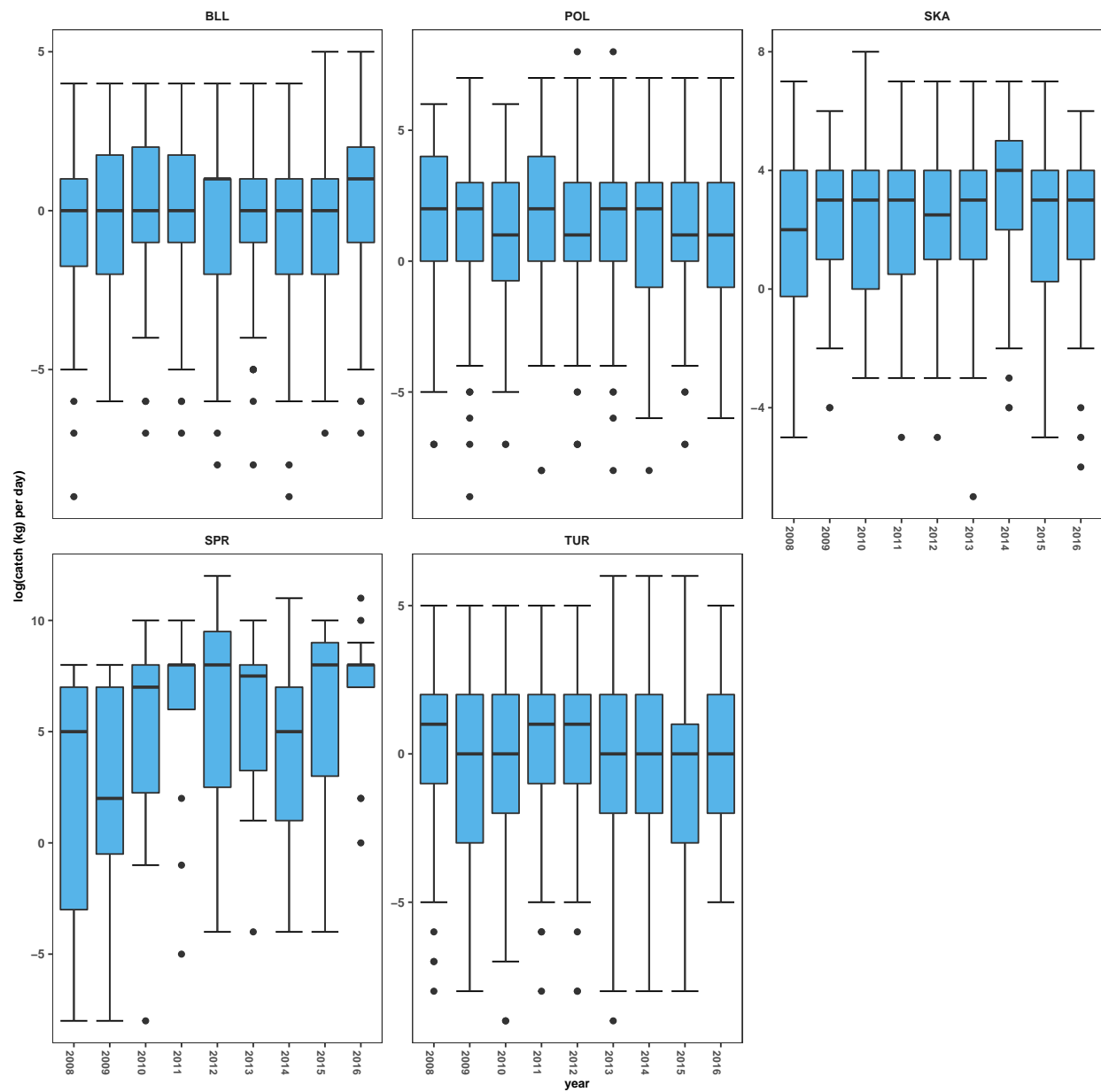
Part of the issues with the main catch data in the stecf database is that the effort does not match up by gear and area with the effort estimations. Here the STECF Annual Economic Report database is used.

Extract the data via sql

```
ct_ef = dbGetQuery(con, paste0("SELECT * FROM data_stecf_aer_cpuedays
                                WHERE speciesgp in ('TUR','BLL','POL','SKA','SPR'); "))
```

Estimate catch in kgs per day

```
ct_ef1 = ct_ef %>% group_by(year, speciesgp)%>%
  mutate(cpue = round(log(totctch/totdays)))
```



**Figure 9** Time series of commercial cpue.

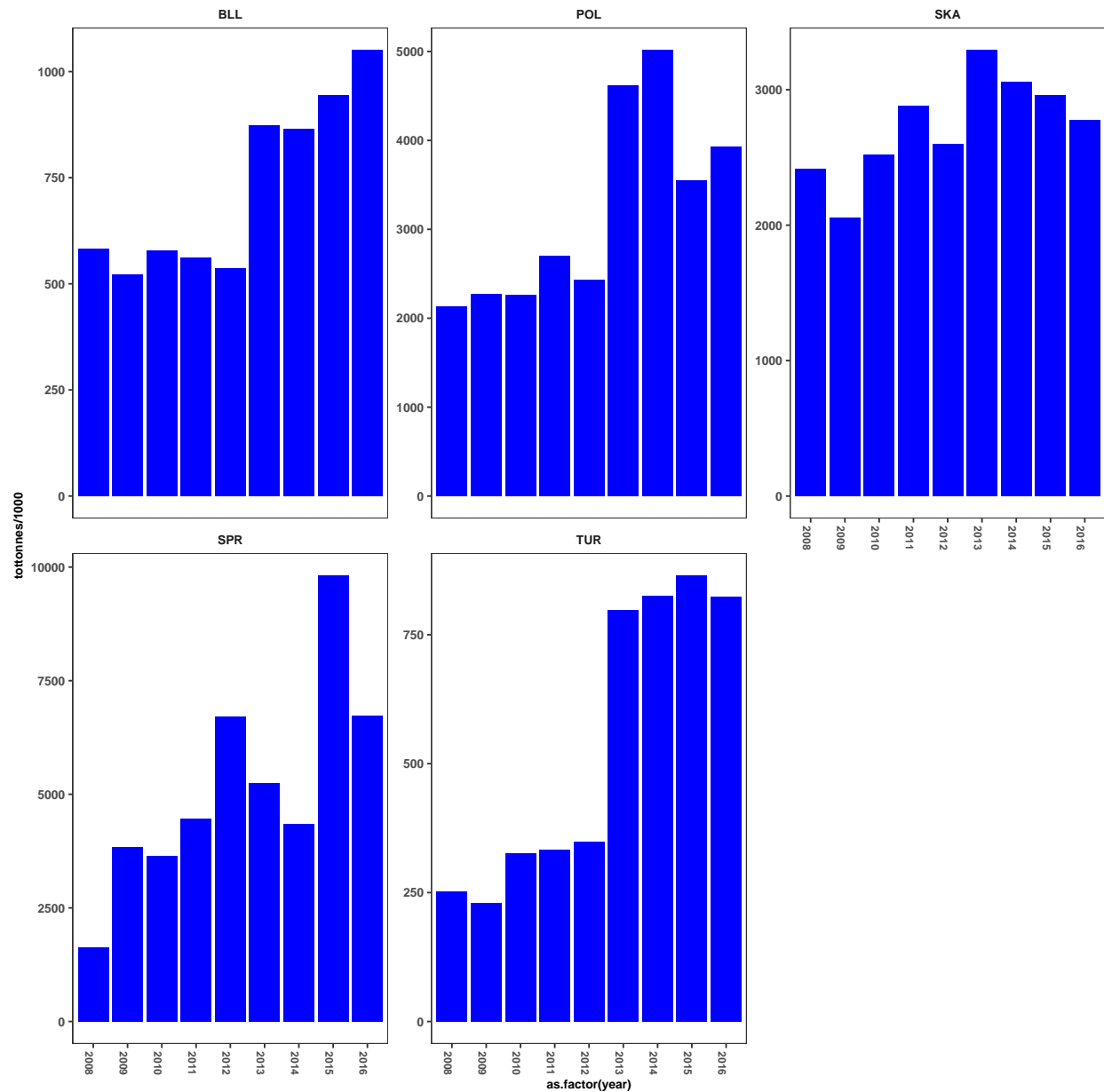


Figure 10 Time series of total catch in tonnes.

## Estimate abundance from survey data

See <http://www.fao.org/docrep/w5449e/w5449e0f.htm> section 13.7

```
rects      = dbGetQuery(con, paste0("select * from data_icesrects"))
surdense   = bio1 %>% group_by( year, speciesgp, ices_rectangle, ices_division) %>%
  summarise(cabund= sum(densbiom_kg_sqkm/1000)) #in tonnes per km2
```

Get survey stations

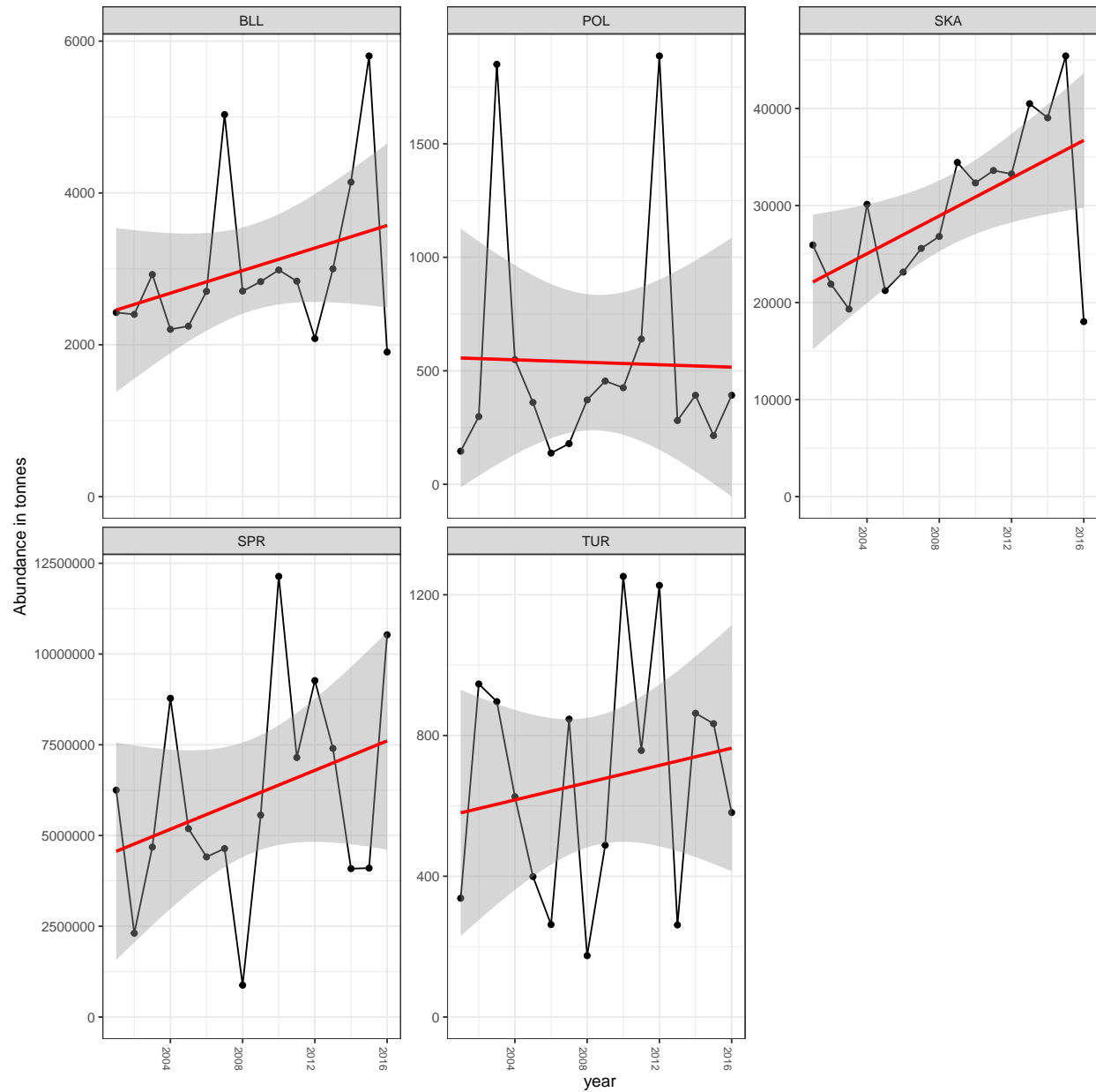
```
surv_eff = dbGetQuery(con, paste0("SELECT *FROM data_surveystns"))
```

Calculate total area/total number of stations the survey covered by division and merge with catch tonnes perkm2

```
nstations = surv_eff %>% group_by(year, ices_rectangle, ices_division) %>%
  summarise(totn=length(unique(haulid))) %>% inner_join(rects)%>%
  group_by(year, ices_division)%>%summarise(totarea=sum(area_km2), totstn=sum(totn)) %>% inner_jo
```

Biomass = totalsurveyarea x 1/total number of stations within the survey area x sum of catch

```
nstations$relabund = ((nstations$totarea/1))*(1/nstations$totstn)*nstations$cabund
nstations2 = nstations %>%group_by(year, speciesgp, ices_division) %>% summarise(totabund= sum(relabund))
nstations3 = nstations2 %>% group_by(year, speciesgp) %>% summarise(totabun= sum(totabund) )
```



## Life history parameters from the literature

```
lh = dbGetQuery(con, paste0("SELECT * FROM data_lhistories
                             WHERE speciesgp in ('TUR','BLL','POL','SKA','SPR'); "))
library(knitr)
kable(lh, caption="life history paramters")
```

Table 1: life history paramters

id	species	speciesgp	t.0	linf	k	lmat	tm	sex	area	source
1	Amblyraja radiata	SKA		66.00	0.2330	46.0	4.0	U	North Sea	[12]

id	species	speciesgp	t.0	linf	k	lmat	tm	sex	area	source
7	Leucoraja circularis	SKA		98.80	0.1210	51.2	5.3	U	North Sea	[12]
8	Leucoraja fullonica	SKA		98.80	0.1210	51.2	5.3	U	North Sea	[12]
17	Pollachius pollachius	POL		85.60	0.1900	NA	NA	U	North Sea	[18]
18	Pollachius pollachius	POL		85.60	0.1860	44.8	3.7	U	North Sea	[12]
23	Psetta maxima	TUR	-0.05	64.80	0.2600	46.0	NA	F	North Sea	[15]
24	Psetta maxima	TUR	-0.51	49.20	0.3700	NA	NA	M	North Sea	[15]
25	Psetta maxima	TUR		65.20	0.3240	49.0	NA	M	Bay of Biscay	[9], [7]
26	Psetta maxima	TUR		73.60	0.2770	49.0	NA	F	Bay of Biscay	[9], [7]
27	Psetta maxima	TUR	-1.79	69.60	0.2497	49.0	4.5	U	North Sea	[19], [8], [16], [17]
28	Psetta maxima	TUR		57.00	0.3200	46.0	4.5	U	North Sea	[12]
29	Psetta maxima	TUR		52.50	0.3200	NA	NA	M	North Sea	[17]
30	Psetta maxima	TUR	-1.79	70.00	0.1480	NA	NA	F	North Sea	[17]
31	Raja batis	SKA	-1.63	254.00	0.0600	130.0	11.0	U	North Sea	[11]
32	Raja batis	SKA		253.70	0.0570	155.0	15.0	U	North Sea	[12]
33	Raja brachyura	SKA		139.00	0.1200	100.0	9.3	U	North Sea	[12]
34	Raja brachyura	SKA	-0.8	118.00	0.1900	NA	NA	M	Irish Sea	[13]
35	Raja brachyura	SKA	-1.52	139.00	0.1200	NA	NA	F	Irish Sea	[13]
36	Raja clavata	SKA	-1.32	128.00	0.0900	NA	NA	F	Irish Sea	[13]
37	Raja clavata	SKA	-0.6	85.60	0.2100	NA	NA	M	Irish Sea	[13]
38	Raja clavata	SKA		105.00	0.2200	65.0	5.0	U	North Sea	[12]
39	Raja montagui	SKA	-0.37	72.80	0.1800	NA	NA	F	Irish Sea	[13]
40	Raja montagui	SKA	-0.56	68.70	0.1900	NA	NA	M	Irish Sea	[13]
41	Raja montagui	SKA		97.80	0.1480	67.0	6.0	U	North Sea	[12]
42	Raja naevus	SKA	-0.465	92.00	0.1100	59.0	9.0	U	North Sea	[11]
43	Raja naevus	SKA		91.60	0.1090	59.0	9.0	U	North Sea	[12]
44	Raja radiata	SKA		66.00	0.2300	46.0	4.0	U	North Sea	[22]
45	Raja undulata	SKA		112.00	0.1000	57.5	6.3	U	North Sea	[12]
46	Scophthalmus rhombus	BLL		74.88	0.1400	37.0	NA	M	Bay of Biscay	[9], [7]
47	Scophthalmus rhombus	BLL		85.23	0.1470	37.0	NA	F	Bay of Biscay	[9], [7]
48	Scophthalmus rhombus	BLL		50.00	0.2700	37.0	4.5	U	North Sea	[12]
49	Sprattus sprattus	SPR	*	16.00	0.6500	13.0	NA	U	North Sea	[20]

## Conclusion from the above analysis for 5 of the 7 case studies

- Pollack and turbot have poor coverage in terms of numbers at length from the survey data and hence potentially poor abundance estimates.
- Brill, sprat and skates have long time series of numbers at length. Brill has adequate numbers from 1993 to 2016 as has skate, while sprat contains a longer time series from 1985-to 2016.
- Observations from the commercial time-series show that for all species there are 9 years of data.
- There are enough life history parameters from the literature to condition an operating model for all case-study species apart from sprat.

## The plan - simulation test the below

- Condition the operating model using life histories (from table above) from brill, turbot, skate and Pollack as per the Mydas project (see the quick start). Will have to use MI estimates for sprat. Use estimates of length (brill, sprat, skate). Obtain length frequencies for turbot and pollack from MI.
- With further investigation look at predicted abundance estimates for brill, sprat (although on high side) and skate (abundance was for all species of skates and rays, will have to make consistent with commercial

species... maybe use thornback only). Pollack and turbot catches are greater than abundance so cannot use these.

- With such good time series of survey length data for brill, sprat and skate it would good to sim test the mean size by using the mlz package <https://cran.r-project.org/web/packages/MLZ/vignettes/MLZ.html#introduction>. This would give an estimate of  $Z$  (assuming constant  $M$ ) and thus look at changes in  $F$ .
- It may also be worth pursuing the LBSR for length-based composition for the above (and compare) with life-history parameters  $M/K$  ratio and  $L_{inf}$  to estimate  $F/M$  and  $F/FMSY$ . With SPR being the biological reference point. <https://cran.r-project.org/web/packages/LBSPR/vignettes/LBSPR.html>
- For turbot and pollack it maybe beneficially to use an empirical approach (as survey info pretty poor) based on the commercial cpue (biomass index from MI lengths or estimated by weights) such as the ICES 2/3 rule (as per Simon Fischer).

For the observation model commercial cpues could be used.

Further: Need to discuss with MI what data we can get for Lobsters and Razors in terms of lengths. We have 1 set of priors from Iyves.

## Software Versions

- R version 3.5.1 (2018-07-02)
- plyr: 1.8.4
- dplyr: 0.7.8
- ggplot2: 3.1.0
- DBI: 1.0.0
- RPostgreSQL: 0.6.2
- reshape 0.8.8
- **Compiled:** Thu Dec 13 13:59:37 2018

## Author information

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## Acknowledgements

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## References

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## Session Info

```
sessionInfo()
```

```
R version 3.5.1 (2018-07-02)
```

```
Platform: x86_64-w64-mingw32/x64 (64-bit)
```

```
Running under: Windows 10 x64 (build 17134)
```

```
Matrix products: default
```

```
locale:
```

```
[1] LC_COLLATE=English_United Kingdom.1252  
[2] LC_CTYPE=English_United Kingdom.1252  
[3] LC_MONETARY=English_United Kingdom.1252  
[4] LC_NUMERIC=C  
[5] LC_TIME=English_United Kingdom.1252
```

```
attached base packages:
```

```
[1] stats      graphics  grDevices  utils      datasets  methods    base
```

```
other attached packages:
```

```
[1] bindrcpp_0.2.2  reshape_0.8.8    RPostgreSQL_0.6-2 DBI_1.0.0  
[5] ggplot2_3.1.0   dplyr_0.7.8      plyr_1.8.4        knitr_1.20
```

```
loaded via a namespace (and not attached):
```

```
[1] tidyselect_0.2.5  purrr_0.2.5      splines_3.5.1  
[4] lattice_0.20-38  colorspace_1.3-2 htmltools_0.3.6  
[7] yaml_2.2.0        base64enc_0.1-3  survival_2.43-3  
[10] rlang_0.3.0.1     pillar_1.3.0     foreign_0.8-71  
[13] glue_1.3.0        withr_2.1.2      RColorBrewer_1.1-2  
[16] bindr_0.1.1       stringr_1.3.1    munsell_0.5.0  
[19] gtable_0.2.0      htmlwidgets_1.3  evaluate_0.12  
[22] labeling_0.3      latticeExtra_0.6-28 highr_0.7  
[25] htmlTable_1.12    Rcpp_1.0.0       acepack_1.4.1  
[28] scales_1.0.0      backports_1.1.2  checkmate_1.8.5  
[31] Hmisc_4.1-1       gridExtra_2.3    digest_0.6.18  
[34] stringi_1.2.4     grid_3.5.1       tools_3.5.1  
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[52] compiler_3.5.1
```