

Elaborazioni Gambero2025

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```
# -----  
# dir.create('input', showWarnings = FALSE)  
# dir.create('output', showWarnings = FALSE)  
# dir.create('plots', showWarnings = FALSE)  
# dir.create('R-script', showWarnings = FALSE)  
library(pacman)  
p_load(tidyverse, stargazer, skimr, janitor, lubridate,  
       ggpubr, rstatix, openxlsx, patchwork, knitr, zoo)  
  
options(scipen = 9999)  
theme_set(theme_minimal(base_size = 18))  
# -----
```

Gambero 2025

Serie storica triennale su *Procambarus clarkii* del lago Trasimeno. Anni di riferimento: luglio 2018 - maggio 2021

```
# import data  
pc <- read.xlsx('input/Gamberi morfometria 3 anni_.xlsx',  
               detectDates = T)  
glimpse(pc)
```

Rows: 3,737

Columns: 13

\$ data	<date> 2018-07-04, 2018-07-04, 2018-07-04, 2018-07-04, 2018-~
\$ n	<dbl> 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16,~
\$ sesso	<chr> "M", "M", "M", "M", "M", "M", "M", "M", "M", "M", "M", "M",~
\$ lt.carapace	<dbl> 43.28, 54.11, 52.73, 81.55, 72.00, 52.21, 54.89, 59.21~

```

$ peso.tot      <dbl> 16.90, 34.50, 30.20, 35.40, 28.50, 33.60, 37.00, 39.10~
$ peso.addome   <dbl> 5.0, 8.2, 7.0, 8.2, 6.0, 7.5, 7.7, 7.8, 8.2, 6.0, 6.4,~
$ peso.epato    <dbl> 0.7, 1.7, 1.8, 1.9, 0.9, 2.1, 1.3, 1.7, 1.6, 1.4, 1.1,~
$ Hiw           <dbl> 4.142012, 4.927536, 5.960265, 5.367232, 3.157895, 6.25~
$ TwB           <dbl> 29.58580, 23.76812, 23.17881, 23.16384, 21.05263, 22.3~
$ SA            <chr> "SI", "SI", "SI", "SI", "SI", "SI", "SI", "SI", "SI", ~
$ muta          <chr> "NO", "NO", "NO", "NO", "NO", "NO", "NO", "NO", "NO", ~
$ maturità.gonadi <chr> NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA~
$ peso.gonadi    <dbl> NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA~

```

controllo qualità dei dati

```

pc %>%
  select(where(is.numeric), sesso, -n) %>%
  group_by(sesso) %>%
  get_summary_stats(type = 'five_number') %>%
  kable(digits = 2)

```

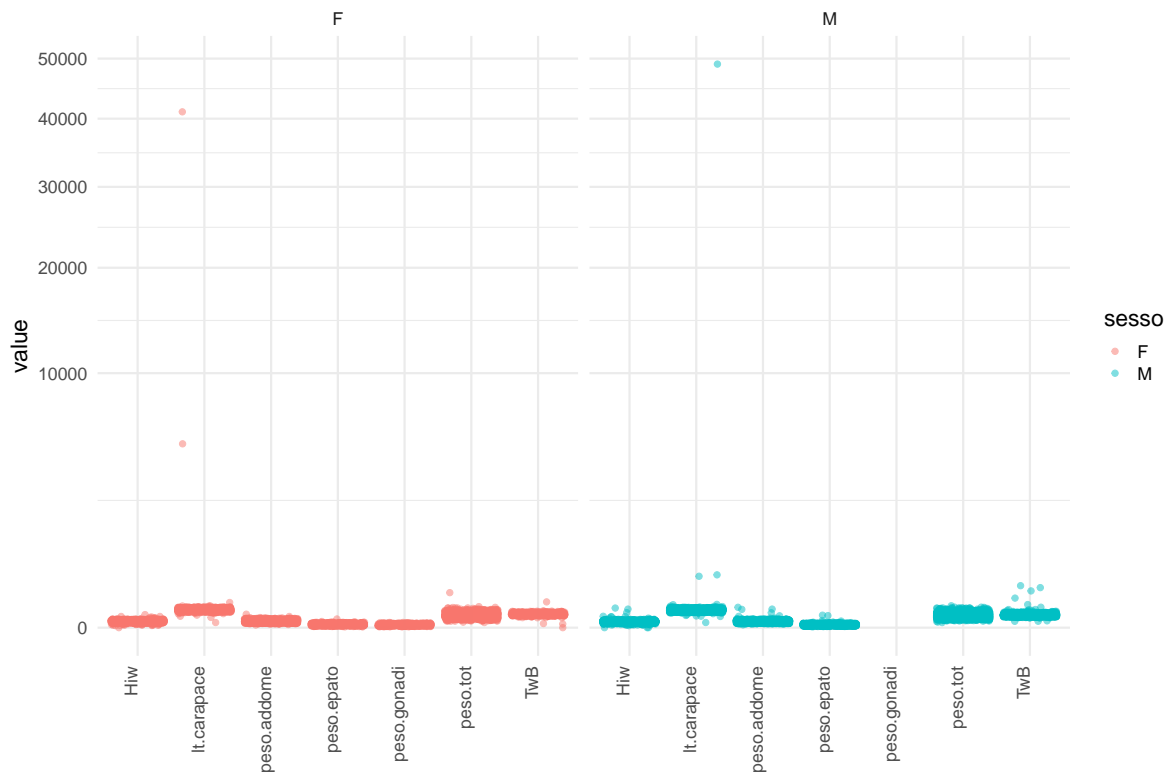
	sezzo	variable	n	min	max	q1	median	q3
	F	lt.carapace	1243	4.06	41081.00	45.41	50.04	54.83
	F	peso.tot	1243	4.00	189.30	16.85	23.50	31.50
	F	peso.addome	1242	1.09	28.00	4.80	6.30	8.30
	F	peso.epato	1242	0.10	11.60	1.10	1.50	2.20
	F	Hiw	1243	0.00	26.36	5.38	7.01	8.33
	F	TwB	1243	0.00	102.56	24.91	27.16	30.02
	F	peso.gonadi	409	0.10	4.20	0.60	1.10	2.00
	M	lt.carapace	2494	4.09	49039.00	43.63	47.41	51.45
	M	peso.tot	2493	4.00	74.10	16.80	21.90	28.70
	M	peso.addome	2494	1.20	64.00	4.30	5.30	6.56
	M	peso.epato	2491	0.10	24.00	0.90	1.20	1.60
	M	Hiw	2493	0.00	59.42	4.65	5.75	6.74
	M	TwB	2493	6.25	271.94	21.65	24.10	26.67

```

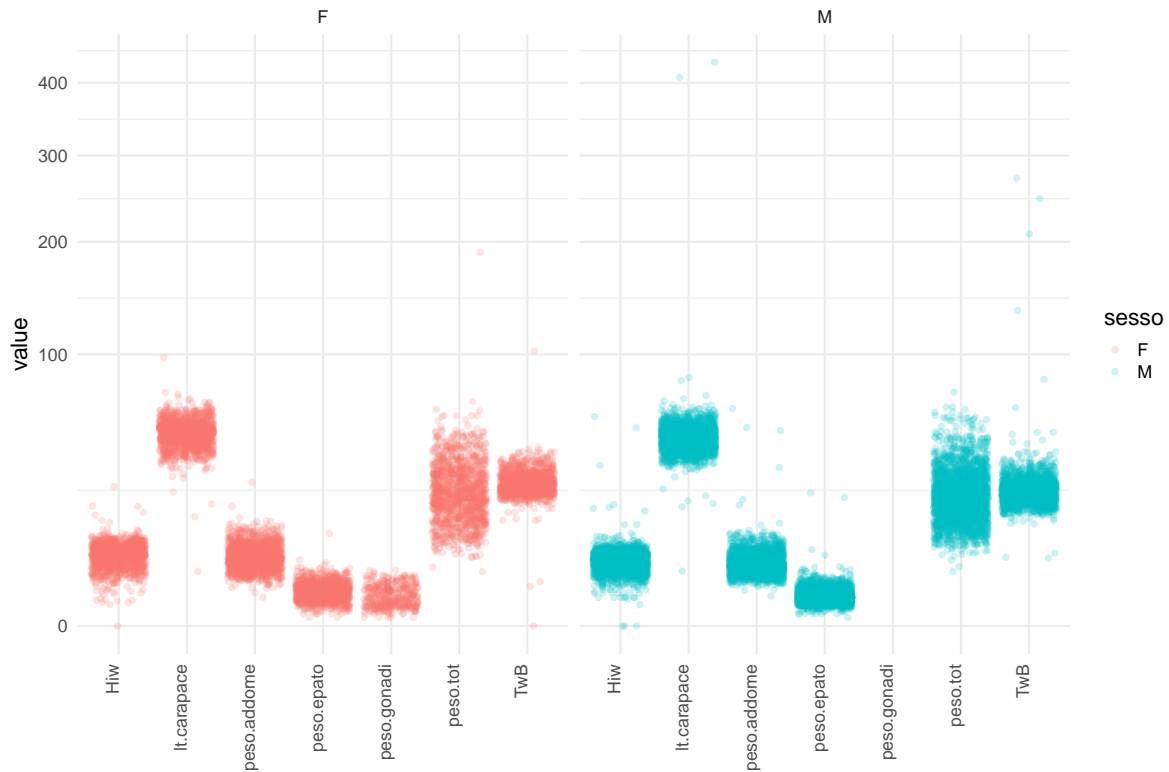
pc %>%
  select(where(is.numeric), sesso) %>%
  pivot_longer(lt.carapace:peso.gonadi,
               names_to = 'var', values_to = 'value') %>%
  ggplot(aes(var, value)) +
  geom_jitter(aes(col=sezzo), alpha=0.5, size=2) +
  scale_y_sqrt() +

```

```
facet_grid(~sesso)+
theme(axis.text.x = element_text(angle = 90, vjust = 0, hjust=1))+
labs(x='')
```



```
pc %>%
  select(where(is.numeric),sesso) %>%
  pivot_longer(lt.carapace:peso.gonadi,
               names_to = 'var', values_to = 'value') %>%
  filter(value<800) %>%
  ggplot(aes(var,value))+
  geom_jitter(aes(col=sesso),alpha=0.2, size=2)+
  scale_y_sqrt()+
  facet_grid(~sesso)+
  theme(axis.text.x = element_text(angle = 90, vjust = 0, hjust=1))+
  labs(x='')
```



```
# add month and year
pc <- pc %>%
  mutate(month=month(data),
         year=year(data),
         date=floor_date(data, unit = 'month'))
```

```
clark.sex <- pc %>%
  group_by(date, sesso) %>%
  summarise(n=n()) %>%
  mutate(tot=cumsum(n))
```

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

```
clark.sexratioF <- clark.sex %>%
  filter(sesso=='F')
clark.sexratioM <- clark.sex %>%
  filter(sesso=='M')
```

```
sexratio <- cbind(data.frame(ratio=round(clark.sexratioM$n/clark.sexratioF$n,1)),
                  distinct(pc,date))
```

```
pc %>%
  count(sezzo,date) %>%
  group_by(date) %>%
  mutate(perc=n/sum(n)) %>%
  ggplot(aes(date,perc,group = sesso))+
  geom_bar(stat = 'identity', alpha=0.9, aes(fill=sesso))+
  scale_y_continuous(labels = scales::percent_format())+
  labs(x='',y='')+
  geom_label(data=sexratio,
            aes(x=date,y=1.1,
                label=paste0(ratio,':1')),
            inherit.aes = FALSE,
            angle=90)+
  theme(legend.position = 'none')
```



```

mute <- pc %>%
  filter(muta=='SI') %>%
  count( Sesso, date) %>%
  mutate(perc=n/sum(n))

mute %>%
  mutate(perc=round(perc*100,2)) %>%
  pivot_wider(id_cols=-n, names_from = Sesso, values_from = perc) %>%
  kable()

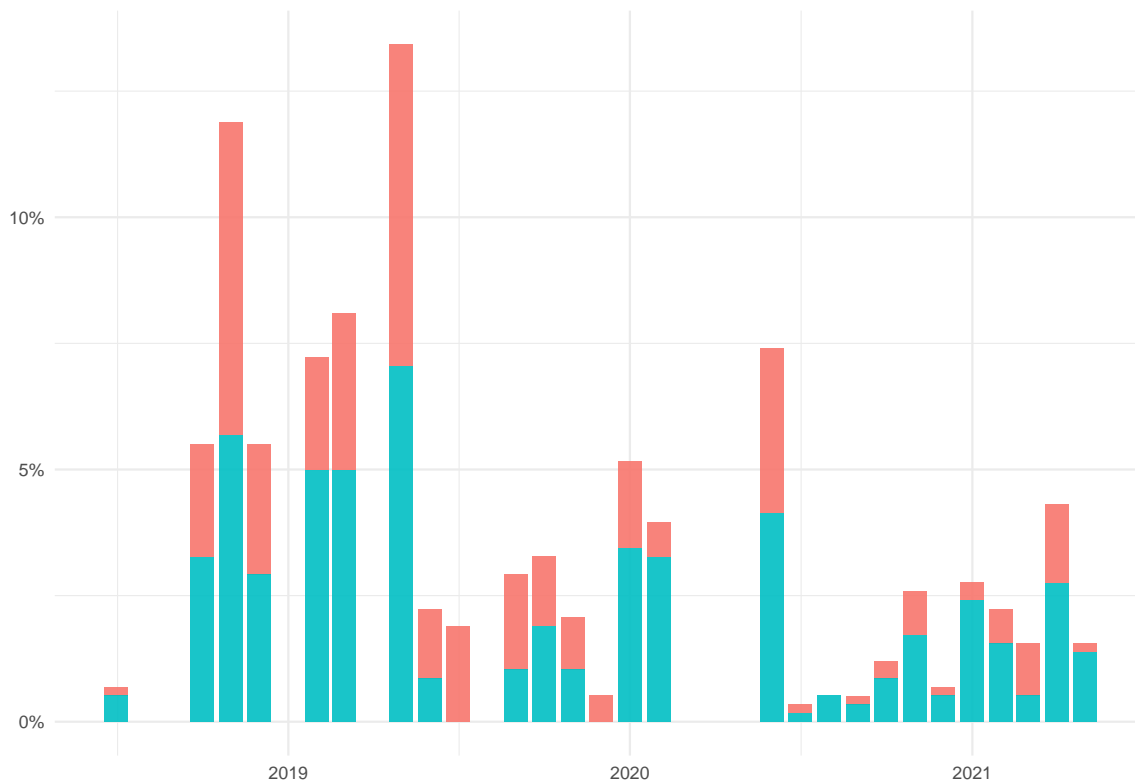
```

date	F	M
2018-07-01	0.17	0.52
2018-10-01	2.24	3.27
2018-11-01	6.20	5.68
2018-12-01	2.58	2.93
2019-02-01	2.24	4.99
2019-03-01	3.10	4.99
2019-05-01	6.37	7.06
2019-06-01	1.38	0.86
2019-07-01	1.89	NA
2019-09-01	1.89	1.03
2019-10-01	1.38	1.89
2019-11-01	1.03	1.03
2019-12-01	0.52	NA
2020-01-01	1.72	3.44
2020-02-01	0.69	3.27
2020-06-01	3.27	4.13
2020-07-01	0.17	0.17
2020-09-01	0.17	0.34
2020-10-01	0.34	0.86
2020-11-01	0.86	1.72
2020-12-01	0.17	0.52
2021-01-01	0.34	2.41
2021-02-01	0.69	1.55
2021-03-01	1.03	0.52
2021-04-01	1.55	2.75
2021-05-01	0.17	1.38
2020-08-01	NA	0.52

```

mute %>% ggplot(aes(date,perc,group = sesso))+
  geom_bar(stat = 'identity', alpha=0.9, aes(fill=sesso))+
  scale_y_continuous(labels = scales::percent_format())+
  labs(x='',y='')+
  theme(legend.position = 'none')

```



```

unique(pc$maturità.gonadi)

```

```

[1] NA "MATURE " "IN VIA DI MATURAZIONE" "INIZIO MATURAZIONE"
[4] "MATURE " "CON LARVE" "UOVA PLEOPODALI" "QUASI MATURE"
[7] "IN VIA DI MATURAZIONE" "UOVA PLEOPODALI" "DEGENERATE"
[10] "INIZIO MATURAZIONE" "QUASI MATURE"

```

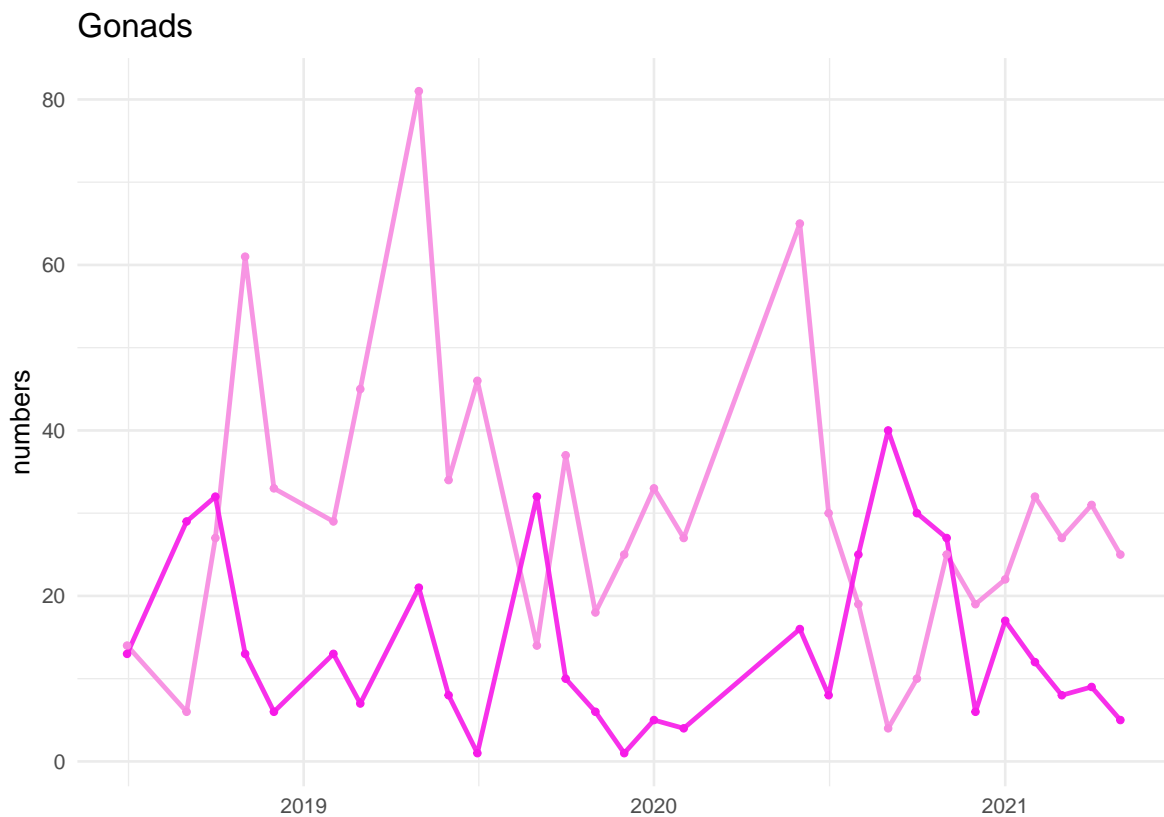
```

# corregge i nomi
pc$maturità.gonadi <- as.factor(pc$maturità.gonadi)
levels(pc$maturità.gonadi)[8] <- 'MATURE'

```

```
# crea una nuova variabile con stadio gonadi: 0, 1
pc.f <- pc %>%
  filter(sesso=='F') %>%
  mutate(gonadBin=ifelse(maturità.gonadi=='MATURE','1','0'))

pc.f %>%
  count(date,gonadBin) %>%
  ggplot(aes(x=date,y=n, group=gonadBin,col=gonadBin)) +
  geom_line(alpha=0.9, lwd=1.5) +
  geom_point(size=2) +
  ggtitle('Gonads') + ylab('numbers')+ xlab('')+
  scale_color_manual(values = c('#f78ae0','#f71ae8'))+
  theme(legend.position = 'none')
```



```
pc.f %>%
  count(date,gonadBin) %>%
  group_by(date) %>%
```



```
mutate(perc=round(n/sum(n)*100,2)) %>%
pivot_wider(id_cols = date,
             values_from = perc,
             names_from = gonadBin) %>%
kable()
```

date	0	1
2018-07-01	51.85	48.15
2018-09-01	17.14	82.86
2018-10-01	45.76	54.24
2018-11-01	82.43	17.57
2018-12-01	84.62	15.38
2019-02-01	69.05	30.95
2019-03-01	86.54	13.46
2019-05-01	79.41	20.59
2019-06-01	80.95	19.05
2019-07-01	97.87	2.13
2019-09-01	30.43	69.57
2019-10-01	78.72	21.28
2019-11-01	75.00	25.00
2019-12-01	96.15	3.85
2020-01-01	86.84	13.16
2020-02-01	87.10	12.90
2020-06-01	80.25	19.75
2020-07-01	78.95	21.05
2020-08-01	43.18	56.82
2020-09-01	9.09	90.91
2020-10-01	25.00	75.00
2020-11-01	48.08	51.92
2020-12-01	76.00	24.00
2021-01-01	56.41	43.59
2021-02-01	72.73	27.27
2021-03-01	77.14	22.86
2021-04-01	77.50	22.50
2021-05-01	83.33	16.67

```
pc.f %>%
count(date,gonadBin) %>%
group_by(date) %>%
mutate(perc=n/sum(n)) %>%
```

```
ggplot(aes(date,perc,group = gonadBin, fill=gonadBin))+
  geom_bar(stat = 'identity', alpha=0.9)+
  scale_y_continuous(labels = scales::percent_format())+
  labs(x='',y='')+
  scale_fill_manual(values = c('#f78ae0','#f71ae8'))+
  theme(legend.position = 'none')
```

