EAD-ASEB-Ssolidissima-OA

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1.1 Welcome

This is the Atlantic surfclam project.

1.2 Quarto information

This project uses quarto. Quarto is similar to Rmarkdown and enables you to weave together content and executable code into a finished document. To learn more about Quarto see https://quarto.org.

When you click the **Render** button a document will be generated that includes both content and the output of embedded code.

2 Biodeposition

2.1 Quarto

Quarto enables you to weave together content and executable code into a finished document. To learn more about Quarto see https://quarto.org.

2.2 Running Code

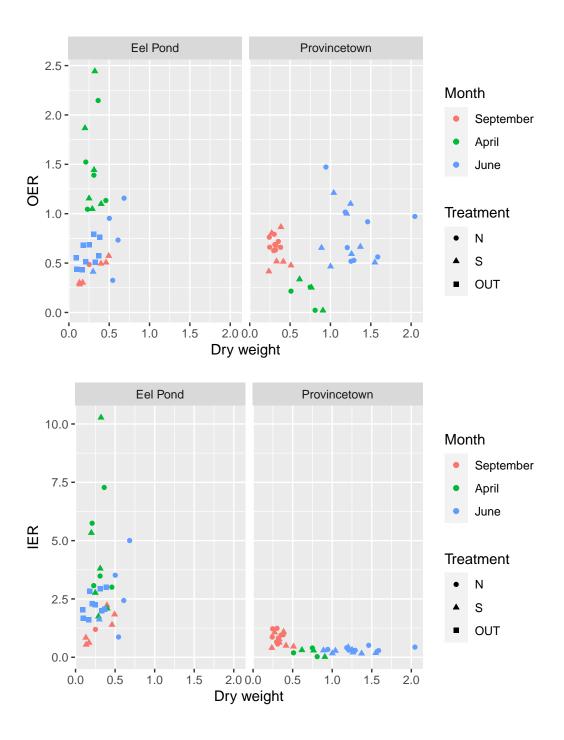
When you click the **Render** button a document will be generated that includes both content and the output of embedded code. You can embed code like this:

```
# bd <- bd[!is.na(bd$Month),]
# bd <- bd[!is.na(bd$Site),]
#print(rownames(bd))

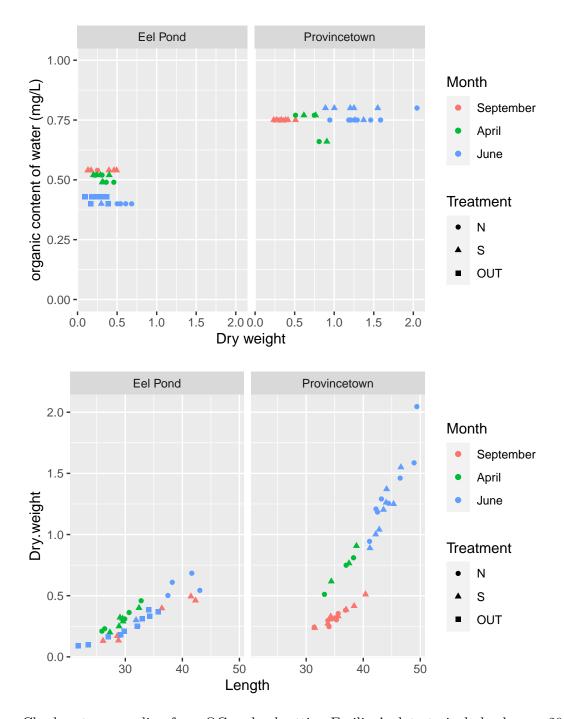
#bd <- bd[complete.cases(bd), ]
bd<- bd[bd$Month =="September"|bd$Month =="April"|bd$Month =="June",]
bd$Site <- as.factor(bd$Site)
bd$Month <- as.factor(bd$Month)
bd$Month <- factor(bd$Month, levels=c("September", "April", "June"))
bd$Treatment <- as.factor(bd$Treatment)
bd$Treatment <- factor(bd$Treatment, levels=c("N", "S", "OUT"))

#print(rownames(bd))
#bd$Treatment[bd$Site=="Eel Pond"] <- "N"
bd <- bd[bd$TPM>=.9,]
#bd$OER
#str(bd)
```

OER.

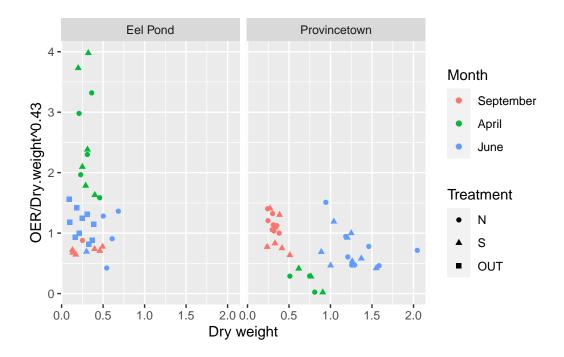


Warning: Removed 1 rows containing missing values (`geom_point()`).



Check out new scaling from QC and subsetting Emilien's data to include clams >20mm and June and July datasets only

```
gg1 <- ggplot(data = bd, aes(x=Dry.weight, y=OER/Dry.weight^.43, color = Month))+
  geom_point(aes(shape=Treatment))+
  xlab("Dry weight") +
  #ylab("OER new scaling")+
  #ylim(0,9)+
  #scale_color_discrete(name="Site")+
  #ggtitle ('OER as a function of DW')+
  facet_wrap(~ Site)</pre>
```



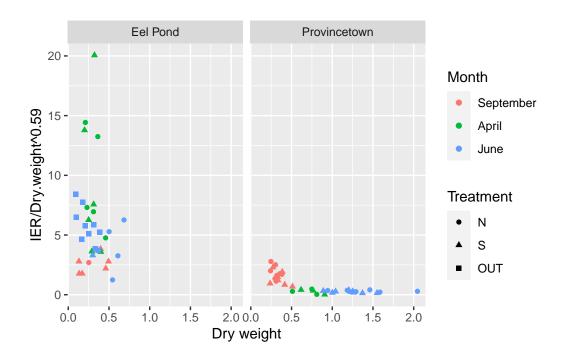
```
gg1 <- ggplot(data = bd, aes(x=Dry.weight, y=IER/Dry.weight^.59, color = Month))+
    geom_point(aes(shape=Treatment))+
    xlab("Dry weight") +

# ylab("IER")+

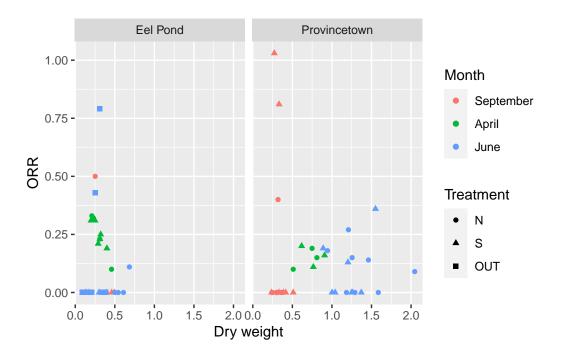
#ylim(0,9)+

#scale_color_discrete(name="Site")+

#ggtitle ('OER as a function of DW')+
    facet_wrap(~ Site)</pre>
gg1
```

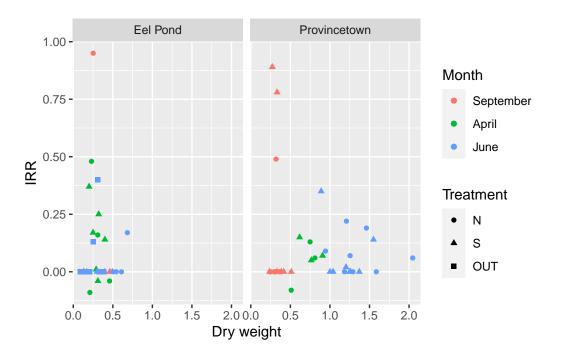


```
gg1 <- ggplot(data = bd, aes(x=Dry.weight, y=ORR, color = Month))+
  geom_point(aes(shape=Treatment))+
  xlab("Dry weight") +
  #ylab("ORR")+
  #ylim(0,9)+
  #scale_color_discrete(name="Site")+
  #ggtitle ('OER as a function of DW')+
  facet_wrap(~ Site)
gg1</pre>
```



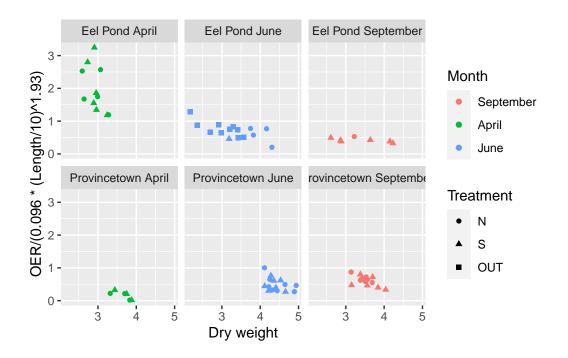
```
gg1 <- ggplot(data = bd, aes(x=Dry.weight, y=IRR, color = Month))+
  geom_point(aes(shape=Treatment))+
  xlab("Dry weight") +

# ylab("IRR")+
  #ylim(0,9)+
  #scale_color_discrete(name="Site")+
  #ggtitle ('OER as a function of DW')+
  facet_wrap(~ Site)</pre>
gg1
```

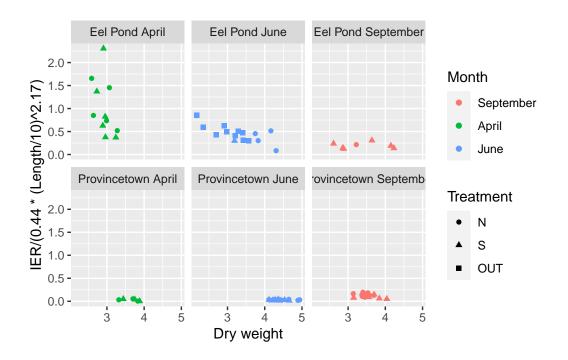


Now using length scaling

```
gg1 <- ggplot(data = bd, aes(x=Length/10, y=0ER/(0.096*(Length/10)^1.93), color = Month))+
    geom_point(aes(shape=Treatment))+
    xlab("Dry weight") +
    #ylab("0ER new scaling")+
    #ylim(0,9)+
    #scale_color_discrete(name="Site")+-
    #ggtitle ('OER as a function of DW')+
    facet_wrap(~ paste(Site,Month))</pre>
```



```
gg1 <- ggplot(data = bd, aes(x=Length/10, y=IER/(.44*(Length/10)^2.17), color = Month))+
    geom_point(aes(shape=Treatment))+
    xlab("Dry weight") +
    #ylab("OER new scaling")+
    #ylim(0,9)+
    #scale_color_discrete(name="Site")+-
    #ggtitle ('OER as a function of DW')+
    facet_wrap(~ paste(Site,Month))
gg1</pre>
```



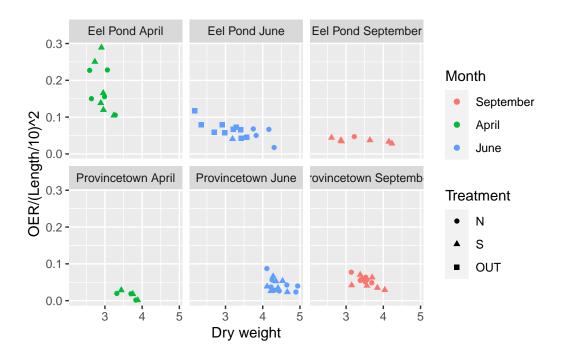
```
gg1 <- ggplot(data = bd, aes(x=Length/10, y=OER/(Length/10)^2, color = Month))+
  geom_point(aes(shape=Treatment))+
  xlab("Dry weight") +

# ylab("IER")+

#ylim(0,9)+

#scale_color_discrete(name="Site")+

#ggtitle ('OER as a function of DW')+
  facet_wrap(~ paste(Site,Month))</pre>
```



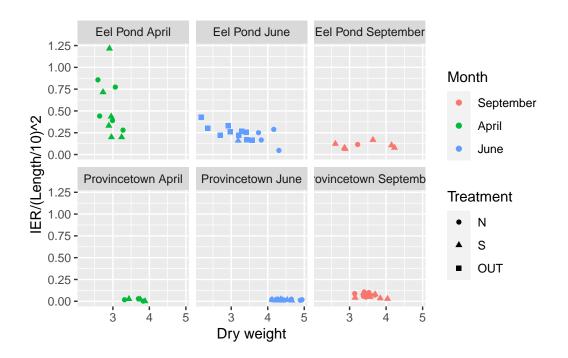
```
gg1 <- ggplot(data = bd, aes(x=Length/10, y=IER/(Length/10)^2, color = Month))+
  geom_point(aes(shape=Treatment))+
  xlab("Dry weight") +

# ylab("IER")+

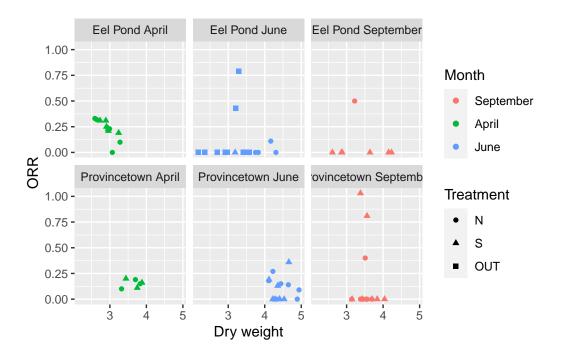
#ylim(0,9)+

#scale_color_discrete(name="Site")+

#ggtitle ('OER as a function of DW')+
  facet_wrap(~ paste(Site,Month))</pre>
```



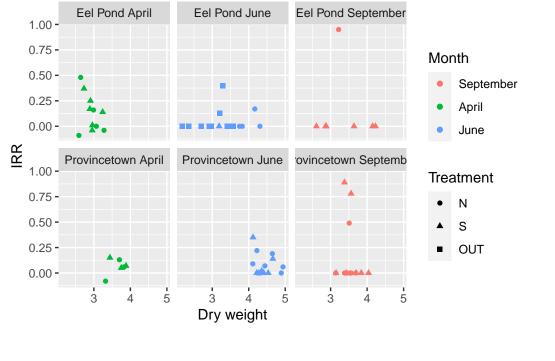
```
gg1 <- ggplot(data = bd, aes(x=Length/10, y=ORR, color = Month))+
  geom_point(aes(shape=Treatment))+
  xlab("Dry weight") +
  #ylab("ORR")+
  #ylim(0,9)+
  #scale_color_discrete(name="Site")+
  #ggtitle ('OER as a function of DW')+
  facet_wrap(~ paste(Site,Month))
gg1</pre>
```



```
gg1 <- ggplot(data = bd, aes(x=Length/10, y=IRR, color = Month))+
  geom_point(aes(shape=Treatment))+
  xlab("Dry weight") +

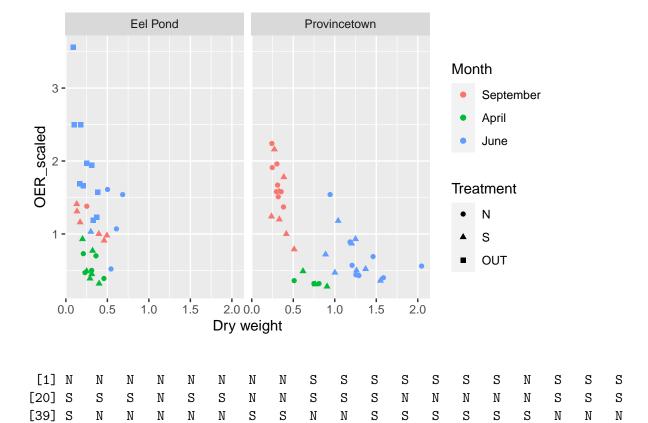
# ylab("IRR")+
  #ylim(0,9)+

#scale_color_discrete(name="Site")+
  #ggtitle ('OER as a function of DW')+
  facet_wrap(~ paste(Site,Month))</pre>
```



[1] N N N N N N N S S S S S S S N [20] S S S S S S S N N S S N S N N [39] S N N N N S S N S S S S S N N N [58] N OUT OUT OUT OUT OUT OUT OUT OUT N S

Levels: N S OUT



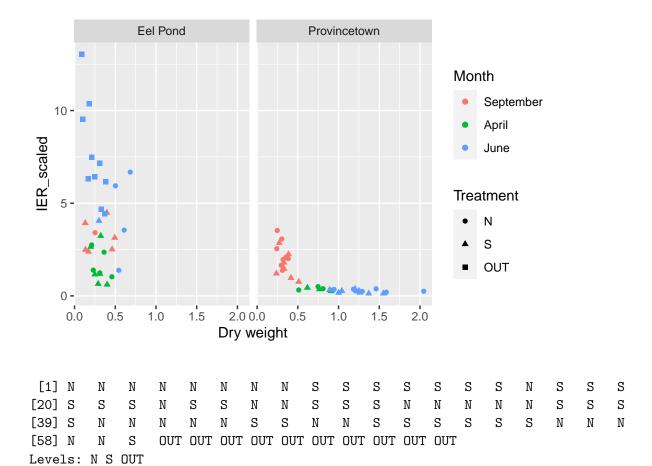
OUT OUT OUT OUT OUT OUT OUT OUT

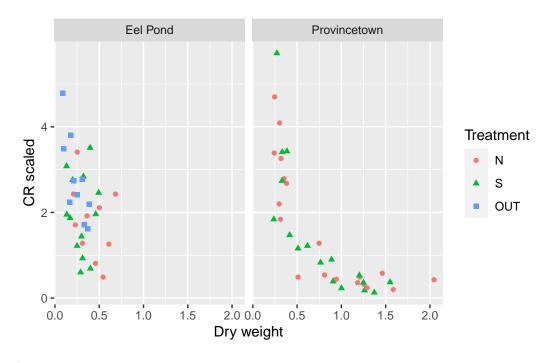
Levels: N S OUT

N

S

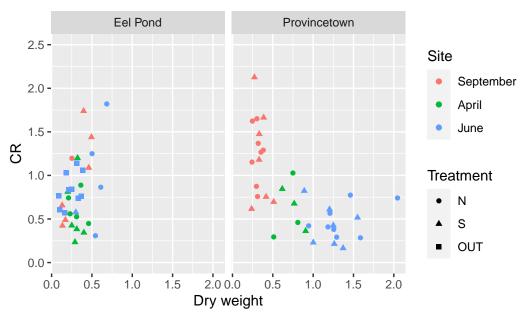
[58] N





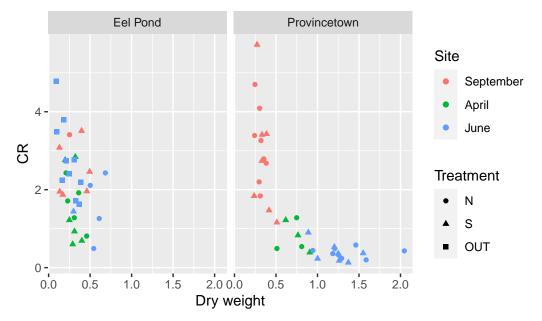
```
gg1 <- ggplot(data = bd, aes(x=Dry.weight, y=CR*Dry.weight^.76, color = Month ))+
    geom_point(aes(shape=Treatment))+
    xlab("Dry weight") +
    ylab("CR")+
    ylim(0,2.5)+
    scale_color_discrete(name="Site")+
    ggtitle ('Unscaled CR as a function of DW')+
    facet_wrap(~ Site)
gg1</pre>
```

Unscaled CR as a function of DW



```
gg2 <- ggplot(data = bd, aes(x=Dry.weight, y=CR, color = Month))+
  geom_point(aes(shape=Treatment))+
  xlab("Dry weight") +
  ylab("CR")+
  #ylim(0,9)+
  scale_color_discrete(name="Site")+
  ggtitle ('Scaled CR as a function of DW')+
  facet_wrap(~ Site)</pre>
```

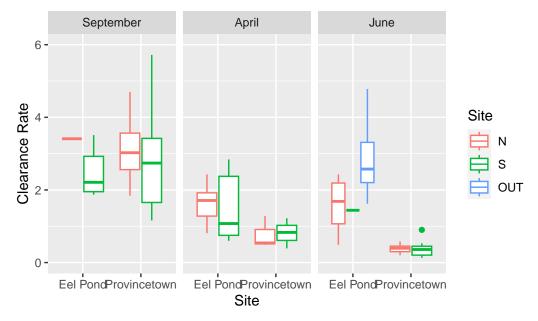
Scaled CR as a function of DW



```
gg3 <- ggplot(data = bd, aes(x=Site, y=CR, col = Treatment))+
    geom_boxplot()+
    #xlab("Dry weight") +
    ylab("Clearance Rate")+
    ylim(0,6)+
    scale_color_discrete(name="Site")+
    ggtitle ('Scaled CR')+
    facet_wrap(~ Month)
gg3</pre>
```

Warning: Removed 1 rows containing non-finite values (`stat_boxplot()`).

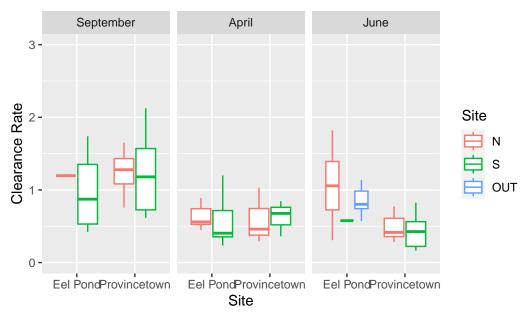
Scaled CR



```
gg4 <- ggplot(data = bd, aes(x=Site, y=CR*Dry.weight^.76, col = Treatment))+
    geom_boxplot()+
    #xlab("Dry weight") +
    ylab("Clearance Rate")+
    ylim(0,3)+
    scale_color_discrete(name="Site")+
    ggtitle ('Unscaled CR')+
    facet_wrap(~ Month)
    gg4</pre>
```

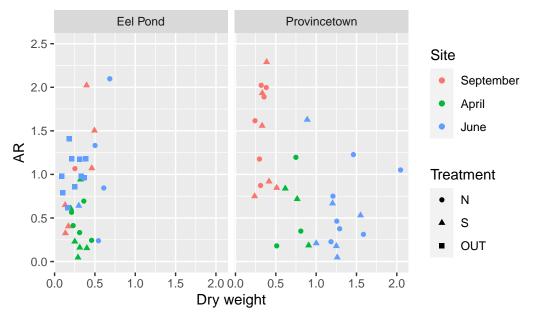
Warning: Removed 1 rows containing non-finite values (`stat_boxplot()`).

Unscaled CR



```
gg1 <- ggplot(data = bd, aes(x=Dry.weight, y=AR*Dry.weight^.76, color = Month ))+
    geom_point(aes(shape=Treatment))+
    xlab("Dry weight") +
    ylab("AR")+
    ylim(0,2.5)+
    scale_color_discrete(name="Site")+
    ggtitle ('Unscaled AR as a function of DW')+
    facet_wrap(~ Site)
gg1</pre>
```

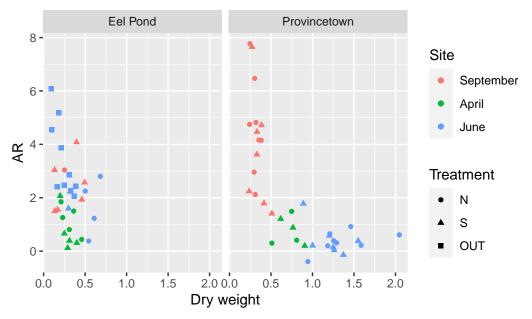
Unscaled AR as a function of DW



```
gg2 <- ggplot(data = bd, aes(x=Dry.weight, y=AR, color = Month))+
  geom_point(aes(shape=Treatment))+
  xlab("Dry weight") +
  ylab("AR")+
  #ylim(0,9)+
  scale_color_discrete(name="Site")+
  ggtitle ('Scaled AR as a function of DW')+
  facet_wrap(~ Site)

gg2</pre>
```

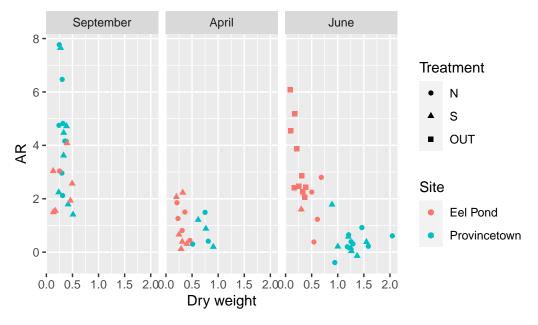
Scaled AR as a function of DW



```
gg2 <- ggplot(data = bd, aes(x=Dry.weight, y=AR, color = Site ))+
  geom_point(aes(shape=Treatment))+
  xlab("Dry weight") +
  ylab("AR")+
  #ylim(0,9)+
  scale_color_discrete(name="Site")+
  ggtitle ('Scaled AR as a function of DW')+
  facet_wrap(~ Month)

gg2</pre>
```

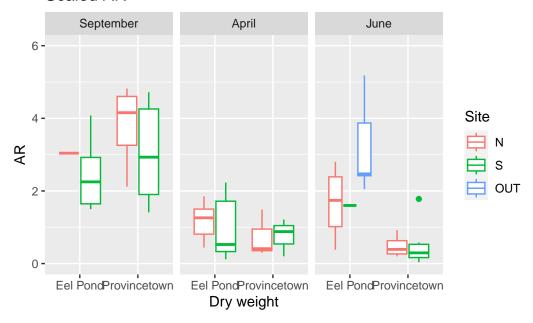
Scaled AR as a function of DW



```
gg3 <- ggplot(data = bd, aes(x=Site, y=AR, col = Treatment))+
  geom_boxplot()+
  xlab("Dry weight") +
  ylab("AR")+
  ylim(0,6)+
  scale_color_discrete(name="Site")+
  ggtitle ('Scaled AR')+
  facet_wrap(~ Month)
  gg3</pre>
```

Warning: Removed 7 rows containing non-finite values (`stat_boxplot()`).

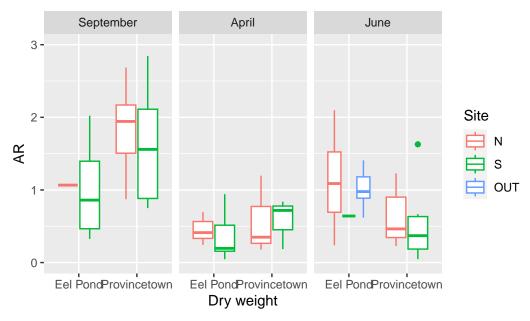
Scaled AR



```
gg4 <- ggplot(data = bd, aes(x=Site, y=AR*Dry.weight^.76, col = Treatment))+
    geom_boxplot()+
    xlab("Dry weight") +
    ylab("AR")+
    ylim(0,3)+
    scale_color_discrete(name="Site")+
    ggtitle ('Unscaled AR')+
    facet_wrap(~ Month)
    gg4</pre>
```

Warning: Removed 3 rows containing non-finite values (`stat_boxplot()`).

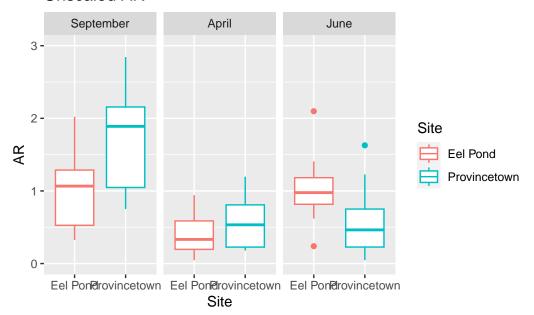
Unscaled AR



```
gg4 <- ggplot(data = bd, aes(x=Site, y=AR*Dry.weight^.76, col = Site))+
    geom_boxplot()+
    #xlab("Dry weight") +
    ylab("AR")+
    ylim(0,3)+
    scale_color_discrete(name="Site")+
    ggtitle ('Unscaled AR')+
    facet_wrap(~ Month)
    gg4</pre>
```

Warning: Removed 3 rows containing non-finite values (`stat_boxplot()`).

Unscaled AR



Call:

```
lm(formula = AR * Dry.weight^0.76 ~ Site * Dry.weight * Month +
    Treatment, data = bd)
```

Residuals:

Min 1Q Median 3Q Max -1.18407 -0.26698 -0.07919 0.26462 1.32257

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	0.17523	0.45048	0.389	0.69879	
SiteProvincetown	2.47106	0.78259	3.158	0.00258	**
Dry.weight	3.09740	1.33860	2.314	0.02444	*

```
MonthApril
                                       0.51011
                                                  0.76689
                                                            0.665
                                                                   0.50872
MonthJune
                                       0.13123
                                                  0.79744 0.165
                                                                   0.86989
TreatmentS
                                      -0.08378
                                                  0.14717 -0.569
                                                                   0.57149
TreatmentOUT
                                                           0.822
                                                                   0.41444
                                       0.37085
                                                  0.45098
SiteProvincetown:Dry.weight
                                      -5.74908
                                                  2.32024 -2.478 0.01632 *
SiteProvincetown: MonthApril
                                      -2.50614
                                                  1.55666 -1.610
                                                                   0.11314
SiteProvincetown:MonthJune
                                      -2.65692
                                                  1.21551 -2.186
                                                                   0.03310 *
Dry.weight:MonthApril
                                      -3.88836
                                                  2.42129 -1.606
                                                                   0.11402
Dry.weight:MonthJune
                                                  1.80279 -0.939 0.35177
                                      -1.69307
SiteProvincetown:Dry.weight:MonthApril 6.49811
                                                  3.46105 1.877
                                                                   0.06576 .
SiteProvincetown:Dry.weight:MonthJune
                                       4.64709
                                                  2.64008 1.760 0.08393 .
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.5139 on 55 degrees of freedom
  (1 observation deleted due to missingness)
Multiple R-squared: 0.573, Adjusted R-squared: 0.4721
F-statistic: 5.678 on 13 and 55 DF, p-value: 1.958e-06
  summary(gm2)
Call:
lm(formula = AR * Dry.weight^0.76 ~ Site + Dry.weight, data = bd)
Residuals:
               1Q
                   Median
                                3Q
                                        Max
-1.26383 -0.45350 -0.07471 0.38898 1.60996
Coefficients:
                Estimate Std. Error t value Pr(>|t|)
                             0.1281
                                      8.473 3.78e-12 ***
(Intercept)
                   1.0858
SiteProvincetown
                  0.6307
                             0.1827
                                      3.452 0.000975 ***
                             0.2089 -4.187 8.54e-05 ***
Dry.weight
                 -0.8748
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.6312 on 66 degrees of freedom
  (1 observation deleted due to missingness)
Multiple R-squared: 0.227, Adjusted R-squared: 0.2036
```

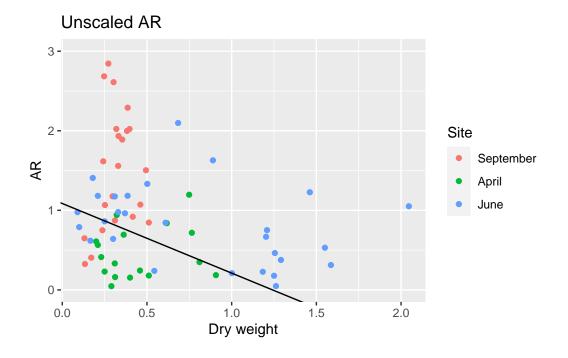
F-statistic: 9.692 on 2 and 66 DF, p-value: 0.000204

summary(gm3)

```
Call:
lm(formula = AR ~ Treatment, data = bd[bd$Site == "Provincetown",
    ])
Residuals:
             1Q Median
    Min
                            3Q
                                   Max
-2.6168 -1.7122 -0.8458 1.8198 5.8153
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
                        0.5265 4.229 0.000167 ***
(Intercept) 2.2268
TreatmentS -0.3921
                         0.7662 -0.512 0.612107
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 2.295 on 34 degrees of freedom
  (1 observation deleted due to missingness)
Multiple R-squared: 0.007645, Adjusted R-squared: -0.02154
F-statistic: 0.2619 on 1 and 34 DF, p-value: 0.6121
  gm2 <- lm(AR*Dry.weight<sup>.76</sup> ~ Site*Dry.weight,
                data = bd
  summary(gm2)
Call:
lm(formula = AR * Dry.weight^0.76 ~ Site * Dry.weight, data = bd)
Residuals:
    Min
               1Q
                   Median
                                 3Q
                                        Max
-1.24011 -0.47053 0.01922 0.37110 1.32959
Coefficients:
                            Estimate Std. Error t value Pr(>|t|)
(Intercept)
                              0.4496
                                        0.2502 1.797 0.07705 .
SiteProvincetown
                              1.3986
                                        0.3158 4.429 3.71e-05 ***
                              1.1409
                                        0.7210 1.583 0.11838
Dry.weight
```

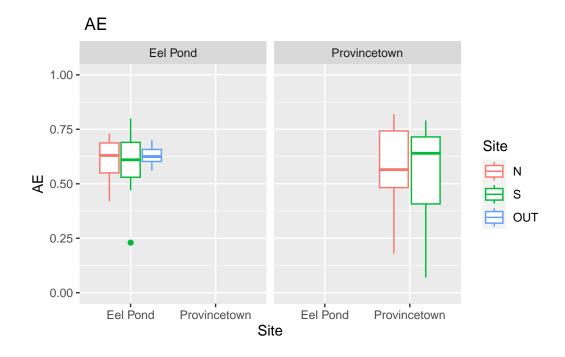
```
SiteProvincetown:Dry.weight -2.1803 0.7498 -2.908 0.00497 **
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.5983 on 65 degrees of freedom
  (1 observation deleted due to missingness)
Multiple R-squared: 0.316, Adjusted R-squared: 0.2844
F-statistic: 10.01 on 3 and 65 DF, p-value: 1.644e-05
  gm2 <- lm(AR*Dry.weight^.76 ~ Site+Dry.weight,
                data = bd
  summary(gm2)
Call:
lm(formula = AR * Dry.weight^0.76 ~ Site + Dry.weight, data = bd)
Residuals:
    Min
               1Q
                   Median
                                30
                                        Max
-1.26383 -0.45350 -0.07471 0.38898 1.60996
Coefficients:
                Estimate Std. Error t value Pr(>|t|)
                             0.1281 8.473 3.78e-12 ***
(Intercept)
                  1.0858
SiteProvincetown 0.6307
                             0.1827
                                      3.452 0.000975 ***
Dry.weight
                 -0.8748
                             0.2089 -4.187 8.54e-05 ***
___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.6312 on 66 degrees of freedom
  (1 observation deleted due to missingness)
Multiple R-squared: 0.227, Adjusted R-squared: 0.2036
F-statistic: 9.692 on 2 and 66 DF, p-value: 0.000204
  #effects(gm2)
  gg4 <- ggplot(data = bd, aes(x=Dry.weight, y=AR*Dry.weight^.76, col = Month))+
    geom_point()+
    geom_abline(slope = coef(gm2)[["Dry.weight"]],
                intercept = coef(gm2)[["(Intercept)"]])+
```

```
xlab("Dry weight") +
ylab("AR")+
ylim(0,3)+
scale_color_discrete(name="Site")+
ggtitle ('Unscaled AR')
#+
#facet_wrap(~ Site)
gg4
```

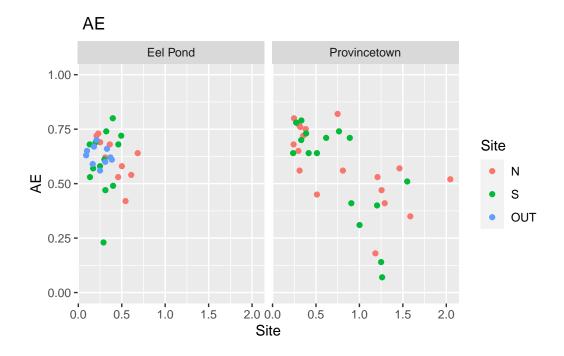


```
gg1 <- ggplot(data = bd, aes(x=Site, y=AE, col = Treatment))+
    geom_boxplot()+
    xlab("Site") +
    ylab("AE")+
    ylim(0,1)+
    scale_color_discrete(name="Site")+
    ggtitle (' AE')+
        facet_wrap(~ Site)
    gg1</pre>
```

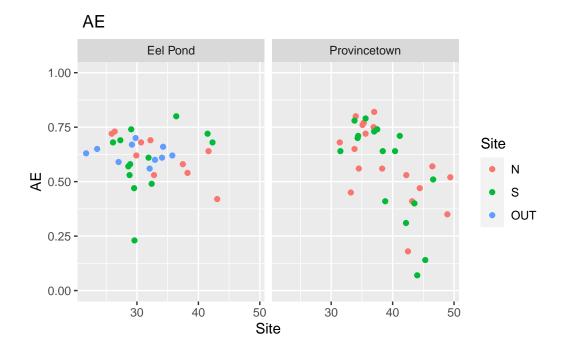
Warning: Removed 3 rows containing non-finite values (`stat_boxplot()`).



```
gg1 <- ggplot(data = bd, aes(x=Dry.weight, y=AE, col = Treatment))+
    geom_point()+
    xlab("Site") +
    ylab("AE")+
    ylim(0,1)+
    scale_color_discrete(name="Site")+
    ggtitle (' AE')+
    facet_wrap(~ Site)
gg1</pre>
```



```
gg1 <- ggplot(data = bd, aes(x=Length, y=AE, col = Treatment))+
    geom_point()+
    xlab("Site") +
    ylab("AE")+
    ylim(0,1)+
    scale_color_discrete(name="Site")+
    ggtitle (' AE')+
    facet_wrap(~ Site)
gg1</pre>
```



Generalized nonlinear least squares fit
 Model: OER ~ A * (Length/10)^B
 Data: bd
 AIC BIC logLik
 25.77933 43.7673 -4.889667

Coefficients:

		Value	Std.Error	t-value	p-value
A.paste(Site,	Month)Eel Pond April	1.1912439	0.5413981	2.200310	0.0315
A.paste(Site,	Month)Eel Pond June	0.5022929	0.2561003	1.961313	0.0543
A.paste(Site,	Month)Eel Pond September	0.3326396	0.1875521	1.773585	0.0810
A.paste(Site,	Month)Provincetown April	0.1407754	0.1150654	1.223438	0.2257
A.paste(Site,	Month)Provincetown June	0.5938241	0.3739325	1.588052	0.1173
A.paste(Site,	Month)Provincetown September	0.5072131	0.2731387	1.856980	0.0680
В		0.2022421	0.4196051	0.481982	0.6315

```
Correlation:
```

```
A. (MPA A. (MPJ A. (MPS A. (SMA A. (SMJ
A.paste(Site, Month)Eel Pond June
                                             0.969
A.paste(Site, Month)Eel Pond September
                                            0.895 0.880
A.paste(Site, Month)Provincetown April
                                             0.660 0.649 0.600
A.paste(Site, Month)Provincetown June
                                             0.983 0.967 0.894 0.659
A.paste(Site, Month)Provincetown September 0.972 0.956 0.884 0.651 0.971
                                           -0.992 -0.976 -0.902 -0.665 -0.991
                                           A.(SMS
A.paste(Site, Month)Eel Pond June
A.paste(Site, Month)Eel Pond September
A.paste(Site, Month)Provincetown April
A.paste(Site, Month)Provincetown June
A.paste(Site, Month)Provincetown September
                                           -0.980
Standardized residuals:
        Min
                     Q1
                                Med
                                              QЗ
                                                         Max
-1.59259716 -0.59069467 -0.03900329 0.44377293 3.52749181
Residual standard error: 0.2735131
Degrees of freedom: 70 total; 63 residual
  # fit2 <- nlme::gnls(OER~ A*(Length/10)^B,</pre>
               params = list(A~paste(Site,Month)-1, B~paste(Site,Month)-1),
               data=bd,
               na.action = na.exclude,
               start = list(A=c(1,1,1,1,1,1), B = c(1,1,1,1,1,1)))
  # summary(fit2)
  fit3 <- nlme::gnls(OER~ A*(Length/10)^B,
             params = list(A \sim 1, B \sim 1),
             data=bd,
             na.action = na.exclude,
             start = list(A=1, B = .3)
  summary(fit1)
Generalized nonlinear least squares fit
  Model: OER ~ A * (Length/10)^B
  Data: bd
```

AIC BIC logLik 25.77933 43.7673 -4.889667

Coefficients:

		Value	Std.Error	t-value	p-value
A.paste(Site,	Month)Eel Pond April	1.1912439	0.5413981	2.200310	0.0315
A.paste(Site,	Month)Eel Pond June	0.5022929	0.2561003	1.961313	0.0543
A.paste(Site,	Month)Eel Pond September	0.3326396	0.1875521	1.773585	0.0810
A.paste(Site,	Month)Provincetown April	0.1407754	0.1150654	1.223438	0.2257
A.paste(Site,	Month)Provincetown June	0.5938241	0.3739325	1.588052	0.1173
A.paste(Site,	Month)Provincetown September	0.5072131	0.2731387	1.856980	0.0680
В		0.2022421	0.4196051	0.481982	0.6315

Correlation:

		A.(MPA	A.(MPJ	A.(MPS	A.(SMA	A.(SMJ
A.paste(Site,	Month)Eel Pond June	0.969	9			
A.paste(Site,	Month) Eel Pond Septemb	ber 0.895	0.880			
A.paste(Site,	Month)Provincetown Apr	ril 0.660	0.649	0.600		
A.paste(Site,	Month)Provincetown Jun	ne 0.983	0.967	0.894	0.659	
A.paste(Site,	Month)Provincetown Sep	ptember 0.972	0.956	0.884	0.651	0.971
В		-0.992	2 -0.976	-0.902	-0.665	-0.991
		A.(SMS	3			

A.paste(Site, Month)Eel Pond June
A.paste(Site, Month)Eel Pond September
A.paste(Site, Month)Provincetown April
A.paste(Site, Month)Provincetown June
A.paste(Site, Month)Provincetown September
B -0.980

Standardized residuals:

Min Q1 Med Q3 Max -1.59259716 -0.59069467 -0.03900329 0.44377293 3.52749181

Residual standard error: 0.2735131

Degrees of freedom: 70 total; 63 residual

summary(fit3)

Generalized nonlinear least squares fit

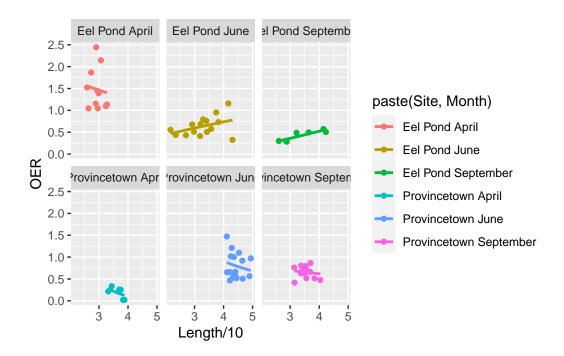
Model: OER ~ A * (Length/10)^B

Data: bd

```
AIC
               BIC
                       logLik
  88.47039 95.21587 -41.23519
Coefficients:
       Value Std.Error t-value p-value
A 1.4461728 0.6820059 2.120470 0.0376
B -0.5246761 0.3808781 -1.377544 0.1729
 Correlation:
 Α
B -0.989
Standardized residuals:
       Min
                   Q1
                             Med
                                         Q3
                                                   Max
-1.5659052 -0.5604893 -0.1701492 0.5193628 3.6540902
Residual standard error: 0.4424753
Degrees of freedom: 70 total; 68 residual
  AIC(fit1,fit3)
             AIC
     df
fit1 8 25.77933
fit3 3 88.47039
  BIC(fit1,fit3)
             BIC
     df
fit1 8 43.76730
fit3 3 95.21587
  gg1 <- ggplot(data = bd, aes(x=Length/10, y=0ER, color = paste(Site,Month)))+
    geom_point()+
    #xlab("Dry weight") +
    #ylab("ORR")+
    #ylim(0,9)+
    #scale_color_discrete(name="Site")+
    #ggtitle ('ORR as a function of DW')+
    stat_smooth(
```

```
method = 'nls',
  formula = y ~ A * x^B,
  method.args = list(start = c(A = 2.3, B = 2)),
  se = FALSE
) +
  facet_wrap(~ paste(Site,Month))

gg1
```



CR is scaled in excel spreadsheet, so to unscale it multiply it... and then here I'm looking at the relationship between length and unscaled CR... I'm not getting a value of 2...

```
Generalized nonlinear least squares fit
  Model: CR * Dry.weight^0.76 ~ A * (Length/10)^B
```

```
Data: bd
      AIC
               BIC
                      logLik
  56.88835 74.7612 -20.44417
Coefficients:
                                               Value Std.Error t-value p-value
A.paste(Site, Month)Eel Pond April
                                           0.2560434 0.1213874 2.109308 0.0390
A.paste(Site, Month)Eel Pond June
                                           0.3488175 0.1772280 1.968185 0.0535
A.paste(Site, Month)Eel Pond September
                                           0.4024601 0.2107537 1.909623 0.0608
A.paste(Site, Month)Provincetown April
                                           0.2230504 0.1287165 1.732881 0.0881
A.paste(Site, Month)Provincetown June
                                           0.1438257 0.0918314 1.566192 0.1224
A.paste(Site, Month)Provincetown September 0.4625192 0.2402667 1.925024 0.0588
                                           0.7765549 0.4075953 1.905211 0.0614
 Correlation:
                                           A. (MPA A. (MPJ A. (MPS A. (SMA A. (SMJ
A.paste(Site, Month)Eel Pond June
                                            0.911
A.paste(Site, Month)Eel Pond September
                                            0.903 0.952
A.paste(Site, Month)Provincetown April
                                                   0.899 0.891
                                            0.853
A.paste(Site, Month)Provincetown June
                                            0.885
                                                   0.934 0.925 0.874
                                            0.920 0.971 0.962 0.909 0.944
A.paste(Site, Month)Provincetown September
                                           -0.929 -0.980 -0.971 -0.917 -0.953
В
                                           A.(SMS
A.paste(Site, Month)Eel Pond June
A.paste(Site, Month)Eel Pond September
A.paste(Site, Month)Provincetown April
A.paste(Site, Month)Provincetown June
A.paste(Site, Month)Provincetown September
В
                                           -0.990
Standardized residuals:
                                Med
                                             QЗ
                                                        Max
        Min
-2.25934022 -0.61005308 -0.06371923 0.67295723 2.72548315
Residual standard error: 0.3432967
Degrees of freedom: 69 total; 62 residual
  fit2 <- nlme::gnls(CR*Dry.weight^.76~ A*(Length/10)^B,
```

data=bd,

na.action = na.exclude,

params = list(A~paste(Site,Month)-1, B~paste(Site,Month)-1),

```
start = list(A=c(1,1,1,1,1,1), B = c(1,1,1,1,1,1)))
  summary(fit2)
Generalized nonlinear least squares fit
 Model: CR * Dry.weight^0.76 ~ A * (Length/10)^B
 Data: bd
       AIC
                BIC
                       logLik
  61.76457 90.80796 -17.88229
Coefficients:
                                               Value Std.Error
                                                                   t-value
A.paste(Site, Month)Eel Pond April
                                            4.335165 10.998719 0.3941518
A.paste(Site, Month)Eel Pond June
                                            0.401470
                                                      0.286231 1.4026097
A.paste(Site, Month)Eel Pond September
                                            0.128397
                                                      0.128146 1.0019640
A.paste(Site, Month)Provincetown April
                                                      2.915631 0.1901769
                                            0.554486
A.paste(Site, Month)Provincetown June
                                            0.092114
                                                      0.472748 0.1948480
A.paste(Site, Month)Provincetown September
                                                      2.716251 0.7079471
                                            1.922962
B.paste(Site, Month)Eel Pond April
                                                      2.402488 -0.7727109
                                           -1.856429
B.paste(Site, Month)Eel Pond June
                                                      0.580814 1.1386753
                                            0.661359
B.paste(Site, Month)Eel Pond September
                                            1.683562
                                                      0.761724 2.2101980
B.paste(Site, Month)Provincetown April
                                            0.075060
                                                      4.058026 0.0184968
B.paste(Site, Month)Provincetown June
                                            1.074964
                                                      3.431508 0.3132627
B.paste(Site, Month)Provincetown September -0.354019
                                                      1.126131 -0.3143675
                                           p-value
A.paste(Site, Month)Eel Pond April
                                            0.6949
A.paste(Site, Month)Eel Pond June
                                            0.1662
A.paste(Site, Month)Eel Pond September
                                            0.3206
A.paste(Site, Month)Provincetown April
                                            0.8498
A.paste(Site, Month)Provincetown June
                                            0.8462
A.paste(Site, Month)Provincetown September
                                            0.4819
B.paste(Site, Month)Eel Pond April
                                            0.4429
B.paste(Site, Month)Eel Pond June
                                            0.2596
B.paste(Site, Month)Eel Pond September
                                            0.0311
B.paste(Site, Month)Provincetown April
                                            0.9853
B.paste(Site, Month)Provincetown June
                                            0.7552
B.paste(Site, Month)Provincetown September
                                            0.7544
Correlation:
                                           A. (MPA A. (MPJ A. (MPS A. (SMA A. (SMJ
A.paste(Site, Month)Eel Pond June
                                            0.000
```

0.000 0.000

0.000 0.000 0.000

A.paste(Site, Month)Eel Pond September

A.paste(Site, Month)Provincetown April

```
A.paste(Site, Month)Provincetown June
                                           0.000
                                                  0.000
                                                         0.000
                                                                0.000
                                                  0.000 0.000
                                                                0.000 0.000
A.paste(Site, Month)Provincetown September
                                           0.000
B.paste(Site, Month)Eel Pond April
                                           -0.998
                                                  0.000 0.000
                                                                0.000 0.000
B.paste(Site, Month)Eel Pond June
                                           0.000 -0.990 0.000
                                                                0.000 0.000
                                                  0.000 - 0.992
                                                                0.000 0.000
B.paste(Site, Month)Eel Pond September
                                           0.000
B.paste(Site, Month)Provincetown April
                                           0.000
                                                  0.000 0.000 -0.999 0.000
B.paste(Site, Month)Provincetown June
                                           0.000
                                                  0.000
                                                         0.000
                                                                0.000 - 0.999
B.paste(Site, Month)Provincetown September
                                           0.000 0.000 0.000 0.000 0.000
                                          A. (SMS B. (MPA B. (MPJ B. (MPS B. (SMA
A.paste(Site, Month)Eel Pond June
A.paste(Site, Month)Eel Pond September
A.paste(Site, Month)Provincetown April
A.paste(Site, Month)Provincetown June
A.paste(Site, Month)Provincetown September
B.paste(Site, Month)Eel Pond April
                                           0.000
B.paste(Site, Month)Eel Pond June
                                           0.000
                                                  0.000
B.paste(Site, Month)Eel Pond September
                                           0.000
                                                  0.000
                                                         0.000
B.paste(Site, Month)Provincetown April
                                           0.000
                                                  0.000
                                                         0.000
                                                                0.000
B.paste(Site, Month)Provincetown June
                                           0.000
                                                  0.000
                                                         0.000
                                                                0.000 0.000
B.paste(Site, Month)Provincetown September -0.999
                                                  0.000
                                                         0.000
                                                                0.000 0.000
                                          B.(SMJ
A.paste(Site, Month)Eel Pond June
A.paste(Site, Month)Eel Pond September
A.paste(Site, Month)Provincetown April
A.paste(Site, Month)Provincetown June
A.paste(Site, Month)Provincetown September
B.paste(Site, Month)Eel Pond April
B.paste(Site, Month)Eel Pond June
B.paste(Site, Month)Eel Pond September
B.paste(Site, Month)Provincetown April
B.paste(Site, Month)Provincetown June
B.paste(Site, Month)Provincetown September 0.000
Standardized residuals:
                               Med
                                            QЗ
       Min
                                                       Max
-2.16273012 -0.58659285 -0.07930908 0.62931579 2.54226464
```

Residual standard error: 0.3449874

Degrees of freedom: 69 total; 57 residual

```
fit3 <- nlme::gnls(CR*Dry.weight^.76~ A*(Length/10)^B,
             params = list(A~1, B~1),
             data=bd,
             na.action = na.exclude,
             start = list(A=1, B = .3)
  summary(fit1)
Generalized nonlinear least squares fit
  Model: CR * Dry.weight^0.76 ~ A * (Length/10)^B
  Data: bd
       AIC
               BIC
                      logLik
  56.88835 74.7612 -20.44417
Coefficients:
                                               Value Std.Error t-value p-value
A.paste(Site, Month)Eel Pond April
                                           0.2560434 \ 0.1213874 \ 2.109308 \ 0.0390
A.paste(Site, Month)Eel Pond June
                                           0.3488175 0.1772280 1.968185 0.0535
A.paste(Site, Month)Eel Pond September
                                           0.4024601 0.2107537 1.909623 0.0608
A.paste(Site, Month)Provincetown April
                                           0.2230504 0.1287165 1.732881 0.0881
A.paste(Site, Month)Provincetown June
                                           0.1438257 0.0918314 1.566192 0.1224
A.paste(Site, Month)Provincetown September 0.4625192 0.2402667 1.925024 0.0588
                                           0.7765549 0.4075953 1.905211 0.0614
 Correlation:
                                           A. (MPA A. (MPJ A. (MPS A. (SMA A. (SMJ
A.paste(Site, Month)Eel Pond June
                                            0.911
A.paste(Site, Month)Eel Pond September
                                            0.903
                                                   0.952
A.paste(Site, Month)Provincetown April
                                            0.853
                                                   0.899
                                                          0.891
A.paste(Site, Month)Provincetown June
                                            0.885
                                                   0.934 0.925 0.874
                                            0.920
A.paste(Site, Month)Provincetown September
                                                   0.971
                                                          0.962 0.909 0.944
                                           -0.929 -0.980 -0.971 -0.917 -0.953
                                           A. (SMS
A.paste(Site, Month)Eel Pond June
A.paste(Site, Month)Eel Pond September
A.paste(Site, Month)Provincetown April
A.paste(Site, Month)Provincetown June
A.paste(Site, Month)Provincetown September
В
                                           -0.990
Standardized residuals:
        Min
                     Q1
                                Med
                                             Q3
                                                        Max
-2.25934022 -0.61005308 -0.06371923 0.67295723 2.72548315
```

Residual standard error: 0.3432967

Degrees of freedom: 69 total; 62 residual

summary(fit3)

Generalized nonlinear least squares fit

Model: CR * Dry.weight^0.76 ~ A * (Length/10)^B

Data: bd

AIC BIC logLik 88.54044 95.24276 -41.27022

Coefficients:

Value Std.Error t-value p-value
A 0.9936092 0.4586190 2.1665241 0.0338
B -0.1644273 0.3666569 -0.4484499 0.6553

Correlation:

Α

B -0.99

Standardized residuals:

Min Q1 Med Q3 Max -1.3732219 -0.7802869 -0.1683142 0.6812916 2.9405469

Residual standard error: 0.4465908

Degrees of freedom: 69 total; 67 residual

AIC(fit1,fit3)

df AIC fit1 8 56.88835 fit3 3 88.54044

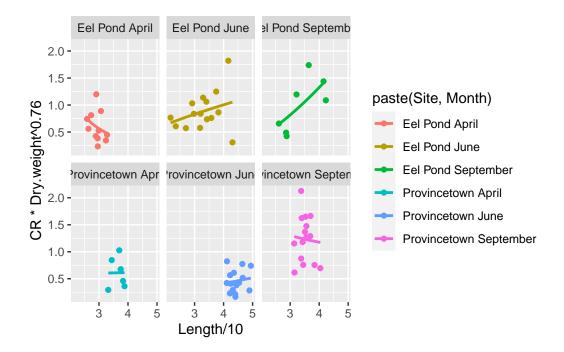
BIC(fit1,fit3)

df BIC fit1 8 74.76120 fit3 3 95.24276

```
gg1 <- ggplot(data = bd, aes(x=Length/10, y=CR*Dry.weight^.76, color = paste(Site,Month)))
geom_point()+
#xlab("Dry weight") +
#ylab("ORR")+
#ylim(0,9)+
#scale_color_discrete(name="Site")+
#ggtitle ('ORR as a function of DW')+
stat_smooth(
   method = 'nls',
   formula = y ~ A * x^B,
   method.args = list(start = c(A = 2.3, B = 2)),
   se = FALSE
) +
facet_wrap(~ paste(Site,Month))</pre>
```

Warning: Removed 1 rows containing non-finite values (`stat_smooth()`).

Warning: Removed 1 rows containing missing values (`geom_point()`).



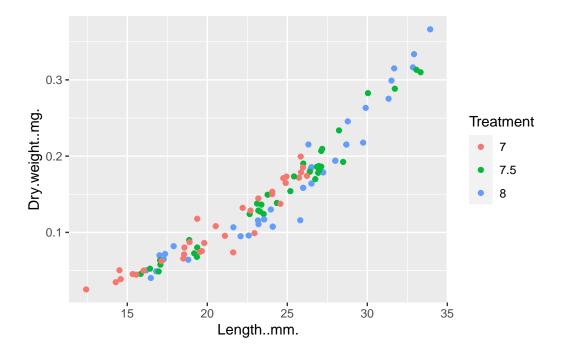
3 Biodeposition_size_scaling

3.1 Quarto

```
library(ggplot2)
  library(ggpubr)
  library(nlme)
Warning: package 'nlme' was built under R version 4.3.2
  library(data.table)
Warning: package 'data.table' was built under R version 4.3.2
  setwd("~/GitHub/EAD-ASEB-Ssolidissima-OA/projects/Biodeposition/data")
  bd_Pousse <- read.csv("Pousse_Meseck_Raw Data Surfclam Biodeposition.csv", stringsAsFactor
  bd_Pousse$Dry.weight..mg. <- bd_Pousse$DW_corr</pre>
  str(bd_Pousse)
'data.frame': 112 obs. of 32 variables:
                : chr "5/15/2019" "5/15/2019" "5/15/2019" "5/15/2019" ...
$ Date
$ pH
                 : num 8 8 8 8 8 8 8 7.5 7.5 7.5 ...
$ Length..mm.
                : num 16.5 16.2 17.3 17.9 16.8 ...
$ Dry.weight..mg.: num 0.0402 0.0503 0.0648 0.0819 0.049 0.0717 0.07 0.0802 0.0634 0.09 ..
$ Length20mm : num NA ...
$ DW_Pred
                 : num NA NA NA NA NA NA NA NA NA ...
$ DW_diff
                : num NA NA NA NA NA NA NA NA NA ...
                : num 0.0402 0.0503 0.0648 0.0819 0.049 0.0717 0.07 0.0802 0.0634 0.09 ..
$ DW_corr
$ time.h.
                 : num 1.5 1.5 1.5 1.5 1.5 1.5 1.5 1.5 1.5 ...
```

```
: num 0.0176 NA NA NA NA ...
$ slope
$ int
                : num
                       -0.269 NA NA NA NA ...
                : num 15.9 NA NA NA NA ...
$ sum
                        2 3 4 5 6 7 8 2 3 4 ...
$ feces
                : int
                : num 1.86 3.79 2.65 1.88 3.46 ...
$ TPM
$ PIM
                        1.56 3.31 2.25 1.54 3.03 ...
                : num
                        0.303\ 0.48\ 0.395\ 0.335\ 0.431\ 0.245\ 0.128\ 0.367\ 0.31\ 0.376\ \dots
$ POM
                : num
$ X..organic
                : num 16.3 12.7 14.9 17.8 12.4 ...
$ ER..mg.h.
                : num 1.24 2.53 1.77 1.25 2.31 ...
                : num 0.202 0.32 0.263 0.223 0.287 ...
$ OER
$ IER
                : num 1.04 2.21 1.5 1.03 2.02 ...
$ X
                : logi NA NA NA NA NA NA ...
$ pseudofeces
                : int 2345678234...
                        2.91 2.5 2.58 3.3 2.05 ...
$ TPM.1
                : num
$ PIM.1
                : num
                        1.64 2.18 2.2 2.87 1.78 ...
$ POM.1
                : num 1.269 0.313 0.389 0.429 0.272 ...
$ X..organ
                        43.6 12.5 15 13 13.3 ...
                : num
$ RR..mg.h.
                : num
                        1.94 1.66 1.72 2.2 1.37 ...
$ ORR
                : num 0.846 0.209 0.259 0.286 0.181 ...
$ IRR
                        1.09 1.46 1.46 1.92 1.19 ...
                : num
$ TSM
                : num
                        15.8 15.8 15.8 15.8 15.8 ...
$ PIM.2
                : num 13.4 13.4 13.4 13.4 13.4 ...
$ POM.2
                : num 2.38 2.38 2.38 2.38 2.38 ...
 bd_Pousse$Treatment <- as.factor(bd_Pousse$pH)</pre>
 bd_Pousse$OER[bd_Pousse$OER<=-.5]<-NA
 bd_Pousse$IER[bd_Pousse$IER<=-.5]<-NA
 bd_Pousse$Dry.weight..mg.[bd_Pousse$Dry.weight..mg.>=.5]<-NA
 ambient <- bd_Pousse[bd_Pousse$Treatment==8,]</pre>
 gg1 <- ggplot(data = bd_Pousse, aes(x=Length..mm., y=Dry.weight..mg., color = Treatment))+</pre>
   gg1
```

Warning: Removed 1 rows containing missing values (`geom_point()`).



```
Formula: Dry.weight..mg. ~ A * Length..mm.^B
```

Parameters:

```
Estimate Std. Error t value Pr(>|t|)
A 3.035e-05 6.047e-06 5.018 2.04e-06 ***
B 2.652e+00 6.003e-02 44.180 < 2e-16 ***
---
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

Residual standard error: 0.0155 on 109 degrees of freedom

```
Number of iterations to convergence: 6
Achieved convergence tolerance: 1.432e-07
(1 observation deleted due to missingness)
```

```
2.652*.76
```

[1] 2.01552

#Note the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included, so am getting the relationship between the summary data does not have length included and so am getting the relationship between the summary data does not have length included and so am getting the relationship between the summary data does not have a summary data does not have a

You can add options to executable code like this

```
gg1 <- ggplot(data = bd_Pousse, aes(x=Dry.weight..mg., y=OER, color = Treatment))+
    geom_point()+
    xlab("Dry weight") +
    ylab("OER")+
    #ylim(0,9)+
    #scale_color_discrete(name="Site")+
    #ggtitle ('OER as a function of DW')+
    stat_smooth(
      method = 'nls',
      formula = y \sim A * x^B,
      method.args = list(start = c(A = 2.3, B = 0.72)),
      se = FALSE
    )#stat_smooth(color = 1, method = 'nls', formula = 'y~a*exp(b*x)',
                   method.args = list(start=c(A = 2.3, B=.72)), se=FALSE) +
    facet_wrap(~ Treatment)
<ggproto object: Class FacetWrap, Facet, gg>
    compute_layout: function
    draw_back: function
    draw_front: function
    draw_labels: function
    draw_panels: function
    finish_data: function
```

init_scales: function
map_data: function

params: list

setup_data: function
setup_params: function

shrink: TRUE

train_scales: function

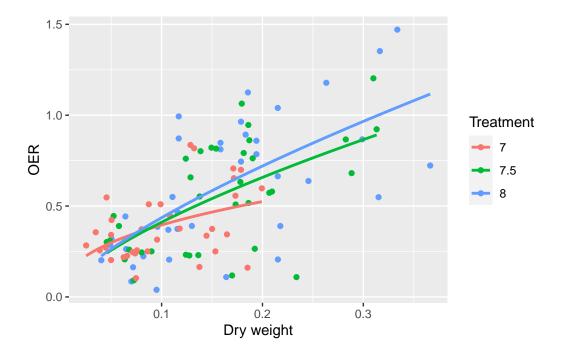
vars: function

super: <ggproto object: Class FacetWrap, Facet, gg>

gg1

Warning: Removed 5 rows containing non-finite values (`stat_smooth()`).

Warning: Removed 5 rows containing missing values (`geom_point()`).



```
nls_7.5 \leftarrow nls(OER \sim A * Dry.weight..mg.^B,
                   start = list(A = 2.3, B = 0.72),
                   data = bd_Pousse[bd_Pousse$Treatment=="7.5",],
                   na.action=na.exclude)
  nls_7 <- nls(OER ~ A * Dry.weight..mg.^B,</pre>
                   start = list(A = 2.3, B = 0.72),
                   data = bd Pousse[bd Pousse$Treatment=="7",],
                   na.action=na.exclude)
  summary(nls 8)
Formula: OER ~ A * Dry.weight..mg.^B
Parameters:
  Estimate Std. Error t value Pr(>|t|)
  2.3096
             0.5933 3.893 0.000412 ***
             0.1567 4.622 4.73e-05 ***
B 0.7242
___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.2824 on 36 degrees of freedom
Number of iterations to convergence: 3
Achieved convergence tolerance: 2.313e-06
  (4 observations deleted due to missingness)
  summary(nls_7.5)
Formula: OER ~ A * Dry.weight..mg.^B
Parameters:
  Estimate Std. Error t value Pr(>|t|)
A 1.9579
             0.6121 3.199 0.003044 **
              0.1806 3.754 0.000672 ***
B 0.6779
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.2477 on 33 degrees of freedom
```

```
Number of iterations to convergence: 3
Achieved convergence tolerance: 9.866e-07
  summary(nls_7)
Formula: OER ~ A * Dry.weight..mg.^B
Parameters:
 Estimate Std. Error t value Pr(>|t|)
  0.1580 2.574 0.01488 *
В
  0.4067
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.1761 on 32 degrees of freedom
Number of iterations to convergence: 5
Achieved convergence tolerance: 7.596e-06
  (1 observation deleted due to missingness)
  nls_all <- nls(OER ~ A * Dry.weight..mg.^B,</pre>
                  start = list(A = 2.3, B = 0.72),
                  data = bd_Pousse,
                  na.action=na.exclude)
  summary(nls_all)
Formula: OER ~ A * Dry.weight..mg.^B
Parameters:
 Estimate Std. Error t value Pr(>|t|)
  2.1245
            0.3439 6.178 1.25e-08 ***
            0.0913 7.800 4.78e-12 ***
B 0.7121
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.2414 on 105 degrees of freedom
Number of iterations to convergence: 3
```

```
Achieved convergence tolerance: 8.352e-07
  (5 observations deleted due to missingness)
  gg1 <- ggplot(data = bd_Pousse, aes(x=Dry.weight..mg., y=IER, color = Treatment))+
    geom_point()+
    xlab("Dry weight") +
    ylab("IER")+
    #ylim(0,9)+
    #scale_color_discrete(name="Site")+
    #ggtitle ('IER as a function of DW')+
    stat_smooth(
      method = 'nls',
      formula = y \sim A * x^B,
      method.args = list(start = c(A = 2.3, B = 0.72)),
      se = FALSE
    )#stat smooth(color = 1, method = 'nls', formula = 'y~a*exp(b*x)',
                   method.args = list(start=c(A = 2.3, B=.72)), se=FALSE) +
    facet_wrap(~ Treatment)
<ggproto object: Class FacetWrap, Facet, gg>
    compute_layout: function
    draw_back: function
    draw_front: function
    draw labels: function
    draw panels: function
    finish data: function
    init_scales: function
    map_data: function
    params: list
    setup_data: function
    setup_params: function
```

```
gg1
```

shrink: TRUE

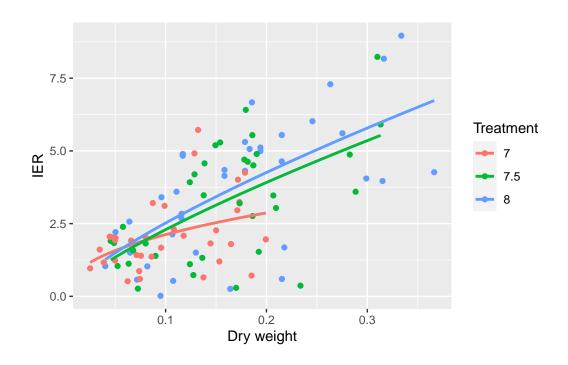
vars: function

train_scales: function

Warning: Removed 5 rows containing non-finite values (`stat_smooth()`).

super: <ggproto object: Class FacetWrap, Facet, gg>

Warning: Removed 5 rows containing missing values (`geom_point()`).



Formula: IER ~ A * Dry.weight..mg.^B

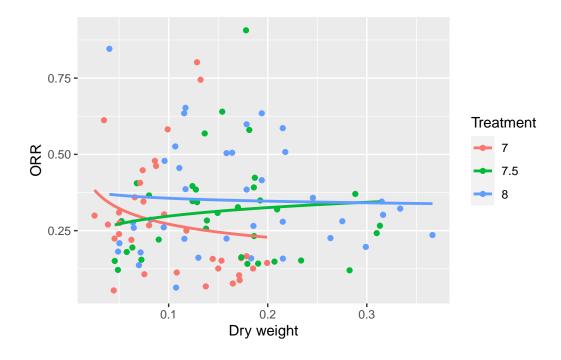
Parameters:

Estimate Std. Error t value Pr(>|t|)

```
3.8196 3.769 0.000588 ***
A 14.3955
              0.1657 4.569 5.56e-05 ***
B 0.7569
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 1.725 on 36 degrees of freedom
Number of iterations to convergence: 5
Achieved convergence tolerance: 4.534e-06
  (4 observations deleted due to missingness)
  summary(nls_7.5)
Formula: IER ~ A * Dry.weight..mg.^B
Parameters:
 Estimate Std. Error t value Pr(>|t|)
A 13.5589 4.8227 2.812 0.008235 **
            0.2094 3.686 0.000812 ***
B 0.7718
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 1.638 on 33 degrees of freedom
Number of iterations to convergence: 5
Achieved convergence tolerance: 3.898e-06
  summary(nls_7)
Formula: IER ~ A * Dry.weight..mg.^B
Parameters:
 Estimate Std. Error t value Pr(>|t|)
A 5.7753 2.4843 2.325
                              0.0266 *
B 0.4346
            0.1934 2.247 0.0317 *
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

```
Residual standard error: 1.147 on 32 degrees of freedom
Number of iterations to convergence: 5
Achieved convergence tolerance: 3.588e-06
  (1 observation deleted due to missingness)
  nls_all <- nls(IER ~ A * Dry.weight..mg.^B,</pre>
                   start = list(A = 2.3, B = 0.72),
                   data = bd_Pousse,
                   na.action=na.exclude)
  summary(nls_all)
Formula: IER ~ A * Dry.weight..mg.^B
Parameters:
 Estimate Std. Error t value Pr(>|t|)
             2.5090 5.647 1.40e-07 ***
A 14.1691
B 0.7912
             0.1026 7.709 7.55e-12 ***
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Residual standard error: 1.54 on 105 degrees of freedom
Number of iterations to convergence: 6
Achieved convergence tolerance: 2.999e-06
  (5 observations deleted due to missingness)
  gg1 <- ggplot(data = bd_Pousse, aes(x=Dry.weight..mg., y=ORR, color = Treatment))+
    geom_point()+
    xlab("Dry weight") +
    ylab("ORR")+
    #ylim(0,9)+
    #scale_color_discrete(name="Site")+
    #ggtitle ('ORR as a function of DW')+
    stat_smooth(
      method = 'nls',
      formula = y \sim A * x^B,
      method.args = list(start = c(A = 2.3, B = 0.72)),
      se = FALSE
```

```
)#stat_smooth(color = 1, method = 'nls', formula = 'y~a*exp(b*x)',
                   method.args = list(start=c(A = 2.3, B=.72)), se=FALSE) +
    facet_wrap(~ Treatment)
<ggproto object: Class FacetWrap, Facet, gg>
    compute_layout: function
    draw_back: function
    draw_front: function
    draw_labels: function
    draw_panels: function
    finish_data: function
    init_scales: function
    map_data: function
    params: list
    setup_data: function
    setup_params: function
    shrink: TRUE
    train_scales: function
    vars: function
    super: <ggproto object: Class FacetWrap, Facet, gg>
  gg1
Warning: Removed 4 rows containing non-finite values (`stat_smooth()`).
Warning: Removed 4 rows containing missing values (`geom_point()`).
```



Formula: ORR ~ A * Dry.weight..mg.^B

Parameters:

```
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.1798 on 37 degrees of freedom
Number of iterations to convergence: 6
Achieved convergence tolerance: 7.096e-06
  (3 observations deleted due to missingness)
  summary(nls_7.5)
Formula: ORR ~ A * Dry.weight..mg.^B
Parameters:
  Estimate Std. Error t value Pr(>|t|)
  0.4011 0.1433 2.798 0.00852 **
              0.1806 0.718 0.47785
  0.1297
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.1692 on 33 degrees of freedom
Number of iterations to convergence: 10
Achieved convergence tolerance: 3.312e-06
  summary(nls_7)
Formula: ORR ~ A * Dry.weight..mg.^B
Parameters:
 Estimate Std. Error t value Pr(>|t|)
A 0.15264 0.08087
                       1.888
                               0.0682 .
B -0.24997
            0.20202 - 1.237
                               0.2250
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.19 on 32 degrees of freedom
Number of iterations to convergence: 11
```

```
Achieved convergence tolerance: 2.023e-06
  (1 observation deleted due to missingness)
  nls_all <- nls(ORR ~ A * Dry.weight..mg.^B,</pre>
                   start = list(A = 2.3, B = 0.72),
                   data = bd_Pousse,
                   na.action=na.exclude)
  summary(nls_all)
Formula: ORR ~ A * Dry.weight..mg.^B
Parameters:
   Estimate Std. Error t value Pr(>|t|)
A 0.315202 0.064442 4.891 3.59e-06 ***
B -0.001062 0.093649 -0.011
                                 0.991
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.181 on 106 degrees of freedom
Number of iterations to convergence: 7
Achieved convergence tolerance: 2.754e-06
  (4 observations deleted due to missingness)
  gg1 <- ggplot(data = bd_Pousse, aes(x=Dry.weight..mg., y=IRR, color = Treatment))+
    geom_point()+
    xlab("Dry weight") +
    ylab("IRR")+
    #ylim(0,9)+
    #scale_color_discrete(name="Site")+
    #ggtitle ('IRR as a function of DW')+
    stat_smooth(
      method = 'nls',
      formula = y \sim A * x^B,
      method.args = list(start = c(A = 2.3, B = 0.72)),
      se = FALSE
    )#stat_smooth(color = 1, method = 'nls', formula = 'y~a*exp(b*x)',
                   method.args = list(start=c(A = 2.3, B=.72)), se=FALSE) +
    facet_wrap(~ Treatment)
```

<ggproto object: Class FacetWrap, Facet, gg>

compute_layout: function

draw_back: function
draw_front: function
draw_labels: function
draw_panels: function
finish_data: function
init_scales: function
map_data: function

params: list

setup_data: function
setup_params: function

shrink: TRUE

train_scales: function

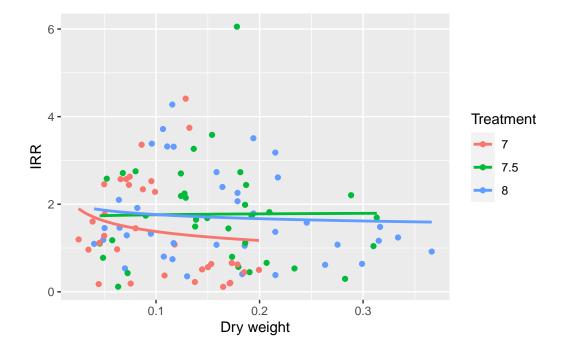
vars: function

super: <ggproto object: Class FacetWrap, Facet, gg>

gg1

Warning: Removed 4 rows containing non-finite values (`stat_smooth()`).

Warning: Removed 4 rows containing missing values (`geom_point()`).



```
nls_8 <- nls(IRR ~ A * Dry.weight..mg.^B,</pre>
                   start = list(A = 2.3, B = 0.72),
                   data = bd_Pousse[bd_Pousse$Treatment=="8",],
                   na.action=na.exclude)
  nls_7.5 <- nls(IRR ~ A * Dry.weight..mg.^B,</pre>
                   start = list(A = 2.3, B = 0.72),
                   data = bd Pousse[bd Pousse$Treatment=="7.5",],
                   na.action=na.exclude)
  nls_7 <- nls(IRR ~ A * Dry.weight..mg.^B,</pre>
                   start = list(A = 2.3, B = 0.72),
                   data = bd_Pousse[bd_Pousse$Treatment=="7",],
                   na.action=na.exclude)
  summary(nls_8)
Formula: IRR ~ A * Dry.weight..mg.^B
Parameters:
 Estimate Std. Error t value Pr(>|t|)
             0.5196
                      2.830 0.00747 **
A 1.4707
B -0.0789
              0.1698 -0.465 0.64489
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Residual standard error: 1.058 on 37 degrees of freedom
Number of iterations to convergence: 9
Achieved convergence tolerance: 3.173e-06
  (3 observations deleted due to missingness)
  summary(nls_7.5)
Formula: IRR ~ A * Dry.weight..mg.^B
Parameters:
  Estimate Std. Error t value Pr(>|t|)
A 1.8275 0.7976 2.291 0.0285 *
  0.0165 0.2137 0.077
                               0.9389
В
```

```
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 1.18 on 33 degrees of freedom
Number of iterations to convergence: 8
Achieved convergence tolerance: 5e-06
  summary(nls_7)
Formula: IRR ~ A * Dry.weight..mg.^B
Parameters:
  Estimate Std. Error t value Pr(>|t|)
A 0.8056
             0.5072
                      1.588
                                0.122
B -0.2328
                                0.342
              0.2411 -0.965
Residual standard error: 1.144 on 32 degrees of freedom
Number of iterations to convergence: 12
Achieved convergence tolerance: 5.953e-06
  (1 observation deleted due to missingness)
  nls_all <- nls(IRR ~ A * Dry.weight..mg.^B,</pre>
                   start = list(A = 2.3, B = 0.72),
                   data = bd_Pousse,
                   na.action=na.exclude)
  summary(nls_all)
Formula: IRR ~ A * Dry.weight..mg.^B
Parameters:
 Estimate Std. Error t value Pr(>|t|)
A 1.4948 0.3667 4.076 8.88e-05 ***
B -0.0437
             0.1109 -0.394
                                0.694
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 1.122 on 106 degrees of freedom
```

```
Number of iterations to convergence: 8
Achieved convergence tolerance: 3.669e-06
  (4 observations deleted due to missingness)
  fit1 <- nlme::gnls(OER~ A*Dry.weight..mg.^B,</pre>
             params = list(A~Treatment-1, B~1),
             data=bd_Pousse,
             na.action = na.exclude,
             start = list(A=c(1,1,1), B = .7)
  summary(fit1)
Generalized nonlinear least squares fit
  Model: OER ~ A * Dry.weight..mg.^B
  Data: bd_Pousse
       AIC
               BIC
                      logLik
  4.853311 18.21745 2.573345
Coefficients:
                   Value Std.Error t-value p-value
A.Treatment7 1.7335946 0.3907440 4.436651
A.Treatment7.5 1.9105770 0.3384029 5.645864
                                                  0
A.Treatment8 2.0961135 0.3464099 6.050963
                                                  0
               0.6633785 0.0960590 6.905949
                                                  0
 Correlation:
               A.Trt7 A.T7.5 A.Trt8
A.Treatment7.5 0.811
A.Treatment8
             0.825 0.852
В
               0.886 0.915 0.931
Standardized residuals:
        Min
                                Med
                                                        Max
                     Q1
                                             Q3
-2.57264637 -0.54686202 0.01601255 0.69985020 2.02837182
Residual standard error: 0.2407639
Degrees of freedom: 107 total; 103 residual
  fit2 <- nlme::gnls(OER~ A*Dry.weight..mg.^B,</pre>
             params = list(A~Treatment-1, B~Treatment-1),
```

```
data=bd Pousse,
             na.action = na.exclude,
             start = list(A=c(1,1,1), B = c(1,1,1))
  summary(fit2)
Generalized nonlinear least squares fit
  Model: OER ~ A * Dry.weight..mg.^B
  Data: bd_Pousse
      AIC
               BIC
                     logLik
  7.379111 26.08891 3.310444
Coefficients:
                   Value Std.Error t-value p-value
A.Treatment7 1.0097516 0.4893917 2.063279 0.0417
A.Treatment7.5 1.9579171 0.5966757 3.281376 0.0014
A.Treatment8 2.3096866 0.5072903 4.552988 0.0000
B.Treatment7
              0.4066719 0.2166225 1.877330 0.0634
B.Treatment7.5 0.6778915 0.1760267 3.851071 0.0002
B.Treatment8 0.7241743 0.1339397 5.406720 0.0000
 Correlation:
               A.Trt7 A.T7.5 A.Trt8 B.Trt7 B.T7.5
A.Treatment7.5 0.000
A.Treatment8 0.000 0.000
B.Treatment7 0.977 0.000 0.000
B.Treatment7.5 0.000 0.972 0.000 0.000
B.Treatment8 0.000 0.000 0.961 0.000 0.000
Standardized residuals:
        Min
                     Q1
                               Med
                                            QЗ
                                                       Max
-2.57536384 -0.50673800 -0.05415554 0.68404928 2.09119265
Residual standard error: 0.2414668
Degrees of freedom: 107 total; 101 residual
  fit3 <- nlme::gnls(OER~ A*Dry.weight..mg.^B,</pre>
             params = list(A \sim 1, B \sim 1),
             data=bd_Pousse,
             na.action = na.exclude,
             start = list(A=1, B = .7)
  summary(fit3)
```

```
Generalized nonlinear least squares fit
  Model: OER ~ A * Dry.weight..mg.^B
  Data: bd_Pousse
       AIC
                BIC
                      logLik
  3.487331 11.50582 1.256334
Coefficients:
      Value Std.Error t-value p-value
A 2.1244875 0.3438641 6.178276
B 0.7121494 0.0912972 7.800344
                                     0
Correlation:
B 0.965
Standardized residuals:
        Min
                     Q1
                                Med
                                              QЗ
                                                         Max
-2.67390205 -0.53146579 -0.02817086 0.67123866 2.20523656
Residual standard error: 0.2414131
Degrees of freedom: 107 total; 105 residual
  AIC(fit1,fit2,fit3)
     df
             AIC
fit1 5 4.853311
fit2 7 7.379111
fit3 3 3.487331
  BIC(fit1,fit2,fit3)
             BIC
     df
fit1 5 18.21745
fit2 7 26.08891
fit3 3 11.50582
  fit_OER_treat <- nlme(OER~ A*Dry.weight..mg.^B,</pre>
             fixed = A + B \sim 1,
             random = A ~ 1 | Treatment,
```

```
data=bd Pousse,
              na.action = na.exclude,
              start = c(A=1, B = .7)
  fit_OER_month <- nlme(OER~ A*Dry.weight..mg.^B,</pre>
             fixed = A + B \sim 1,
              random = A ~ 1 | Month,
              data=bd_Pousse,
             na.action = na.exclude,
              start = c(A=1, B = .7)
  fit_OER_crossed <- nlme(OER~ A*Dry.weight..mg.^B,</pre>
             fixed = A + B \sim 1,
              random = list(Treatment=(A+B~1), Month=(A~1)),
              data=bd_Pousse,
             na.action = na.exclude,
              start = c(A=1, B = .7)
  summary(fit_OER_treat)
Nonlinear mixed-effects model fit by maximum likelihood
  Model: OER ~ A * Dry.weight..mg.^B
  Data: bd_Pousse
       ATC
                BIC
                      logLik
  5.487331 16.17865 1.256334
Random effects:
 Formula: A ~ 1 | Treatment
                   A Residual
StdDev: 2.939036e-05 0.2391463
Fixed effects: A + B ~ 1
      Value Std.Error DF t-value p-value
A 2.1244875 0.3438642 103 6.178276
B 0.7121494 0.0912972 103 7.800344
Correlation:
B 0.965
Standardized Within-Group Residuals:
       Min
                   Q1
                              Med
                                          QЗ
                                                    Max
-2.6992477 -0.5365035 -0.0284379 0.6776013 2.2261397
```

```
Number of Observations: 107
Number of Groups: 3
  summary(fit_OER_month)
Nonlinear mixed-effects model fit by maximum likelihood
  Model: OER ~ A * Dry.weight..mg.^B
  Data: bd_Pousse
        AIC
                 BIC logLik
  -10.21828 0.4730363 9.10914
Random effects:
 Formula: A ~ 1 | Month
                A Residual
StdDev: 0.4298785 0.2139649
Fixed effects: A + B ~ 1
      Value Std.Error DF t-value p-value
A 2.3583317 0.5052313 101 4.667826
B 0.7689894 0.1024072 101 7.509137
 Correlation:
 Α
B 0.899
Standardized Within-Group Residuals:
                    Q1
                                Med
                                             QЗ
                                                        Max
-3.33824715 -0.54182672 0.07714807 0.65787227 2.03575835
Number of Observations: 107
Number of Groups: 5
  summary(fit_OER_crossed)
Nonlinear mixed-effects model fit by maximum likelihood
  Model: OER ~ A * Dry.weight..mg.^B
 Data: bd_Pousse
      AIC
               BIC
                      logLik
  1.333443 20.04324 6.333279
```

```
Random effects:
 Formula: list(A ~ 1, B ~ 1)
 Level: Treatment
 Structure: General positive-definite, Log-Cholesky parametrization
  StdDev
A 1.904067e-05 A
B 1.916928e-08 0
Formula: A ~ 1 | Month %in% Treatment
                A Residual
StdDev: 0.3330295 0.2102274
Fixed effects: A + B ~ 1
      Value Std.Error DF t-value p-value
A 1.4383438 0.3197207 91 4.498750
B 0.5142906 0.1077050 91 4.774993
                                         0
 Correlation:
  Α
B 0.943
Standardized Within-Group Residuals:
        Min
                     Q1
                                 Med
                                              QЗ
                                                          Max
-2.85392669 -0.54127778 0.01701554 0.59453380 2.20672688
Number of Observations: 107
Number of Groups:
           Treatment Month %in% Treatment
                   3
                                        15
  fit_OER_crossed <- nlme(OER~ A*Dry.weight..mg.^B,</pre>
             fixed = A + B \sim 1,
             random = list(Treatment=(A+B~1), Month=(A~1)),
              data=bd_Pousse,
             na.action = na.exclude,
              start = c(A=1, B = .7)
  fit_IER_crossed <- nlme(IER~ A*Dry.weight..mg.^B,</pre>
              fixed = A + B \sim 1,
              random = list(Treatment=(A+B~1), Month=(A~1)),
              data=bd_Pousse,
             na.action = na.exclude,
              start = c(A=1, B = .7)
  fit_ORR_crossed <- nlme(ORR~ A*Dry.weight..mg.^B,</pre>
```

```
fixed = A + B \sim 1,
             random = list(Treatment=(A~1)), #Removed random effect of month bc would not co
             data=bd_Pousse,
             na.action = na.exclude,
             start = c(A=1, B = .7)
  fit_IRR_crossed <- nlme(IRR~ A*Dry.weight..mg.^B,</pre>
             fixed = A + B \sim 1,
             random = list(Treatment=(A+B~1), Month=(A~1)),
             data=bd_Pousse,
             na.action = na.exclude,
             start = c(A=1, B = .7)
  summary(fit_OER_crossed)
Nonlinear mixed-effects model fit by maximum likelihood
  Model: OER ~ A * Dry.weight..mg.^B
  Data: bd_Pousse
       AIC
                BIC
                      logLik
  1.333443 20.04324 6.333279
Random effects:
 Formula: list(A ~ 1, B ~ 1)
 Level: Treatment
 Structure: General positive-definite, Log-Cholesky parametrization
  StdDev
               Corr
A 1.904067e-05 A
B 1.916928e-08 0
 Formula: A ~ 1 | Month %in% Treatment
                A Residual
StdDev: 0.3330295 0.2102274
Fixed effects: A + B ~ 1
      Value Std.Error DF t-value p-value
A 1.4383438 0.3197207 91 4.498750
B 0.5142906 0.1077050 91 4.774993
 Correlation:
  Α
B 0.943
Standardized Within-Group Residuals:
                     Q1
                                Med
                                              QЗ
        Min
                                                         Max
```

-2.85392669 -0.54127778 0.01701554 0.59453380 2.20672688

Number of Observations: 107

Number of Groups:

Treatment Month %in% Treatment 3 15

summary(fit_IER_crossed)

Nonlinear mixed-effects model fit by maximum likelihood

Model: IER ~ A * Dry.weight..mg.^B

Data: bd Pousse

AIC BIC logLik 399.8229 418.5327 -192.9114

Random effects:

Formula: list(A ~ 1, B ~ 1)

Level: Treatment

Structure: General positive-definite, Log-Cholesky parametrization

StdDev Corr A 2.486702e-04 A B 2.244356e-07 0

Formula: A ~ 1 | Month %in% Treatment

A Residual

StdDev: 2.252674 1.375624

Fixed effects: A + B \sim 1

Value Std.Error DF t-value p-value A 10.459513 2.3585562 91 4.434710 0 B 0.631461 0.1125008 91 5.612946 0

Correlation:

A 0.4

B 0.944

Standardized Within-Group Residuals:

Min Q1 Med Q3 Max -2.836393322 -0.472751163 -0.004826624 0.539504100 2.275859320

Number of Observations: 107

Number of Groups:

Treatment Month %in% Treatment 3 15

summary(fit_ORR_crossed)

Nonlinear mixed-effects model fit by maximum likelihood

Model: ORR ~ A * Dry.weight..mg.^B

Data: bd_Pousse

AIC BIC logLik

-56.755 -46.02648 32.3775

Random effects:

Formula: A ~ 1 | Treatment

A Residual

StdDev: 0.000644228 0.1792925

Fixed effects: A + B ~ 1

Value Std.Error DF t-value p-value

A 0.31522344 0.06445193 104 4.890830 0.0000

B -0.00102379 0.09365732 104 -0.010931 0.9913

Correlation:

Α

B 0.963

Standardized Within-Group Residuals:

Min Q1 Med Q3 Max -1.4624874 -0.8394831 -0.2115732 0.4595231 3.2956680

Number of Observations: 108

Number of Groups: 3

summary(fit_IRR_crossed)

Nonlinear mixed-effects model fit by maximum likelihood

Model: IRR ~ A * Dry.weight..mg.^B

Data: bd_Pousse

AIC BIC logLik

343.3118 362.0868 -164.6559

Random effects:

```
Formula: list(A ~ 1, B ~ 1)
 Level: Treatment
 Structure: General positive-definite, Log-Cholesky parametrization
               Corr
A 3.134309e-04 A
B 3.966123e-05 0.001
Formula: A ~ 1 | Month %in% Treatment
                   A Residual
StdDev: 8.081668e-05 1.111436
Fixed effects: A + B ~ 1
      Value Std.Error DF t-value p-value
A 1.496221 0.3670782 92 4.076028 0.0001
B -0.043238 0.1108656 92 -0.390004 0.6974
Correlation:
  Α
B 0.963
Standardized Within-Group Residuals:
                   Q1
                             Med
                                         QЗ
                                                   Max
-1.4117544 -0.8373900 -0.2023488 0.6797753 3.9965273
Number of Observations: 108
Number of Groups:
           Treatment Month %in% Treatment
                                       15
  fit1 <- nlme::gnls(OER~ A*(Length..mm./10)^B,
             params = list(A~Treatment-1, B~1),
             data=bd_Pousse,
             na.action = na.exclude,
             start = list(A=c(1,1,1), B = .3))
  summary(fit1)
Generalized nonlinear least squares fit
  Model: OER ~ A * (Length..mm./10)^B
  Data: bd_Pousse
       AIC
                BIC
                      logLik
  5.582898 18.94704 2.208551
```

Coefficients:

Value Std.Error t-value p-value
A.Treatment7 0.0973070 0.02380870 4.087037 1e-04
A.Treatment7.5 0.1029169 0.02855707 3.603904 5e-04
A.Treatment8 0.1111913 0.03149990 3.529893 6e-04
B 1.8344623 0.27164763 6.753095 0e+00

Correlation:

A.Trt7 A.T7.5 A.Trt8

A.Treatment7.5 0.871

A.Treatment8 0.881 0.944

B -0.901 -0.966 -0.977

Standardized residuals:

Min Q1 Med Q3 Max -2.40806503 -0.53830392 -0.00303497 0.70306688 1.99271682

Residual standard error: 0.2415861

Degrees of freedom: 107 total; 103 residual

Generalized nonlinear least squares fit

Model: OER ~ A * (Length..mm./10)^B

Data: bd_Pousse

AIC BIC logLik 6.687948 25.39775 3.656026

Coefficients:

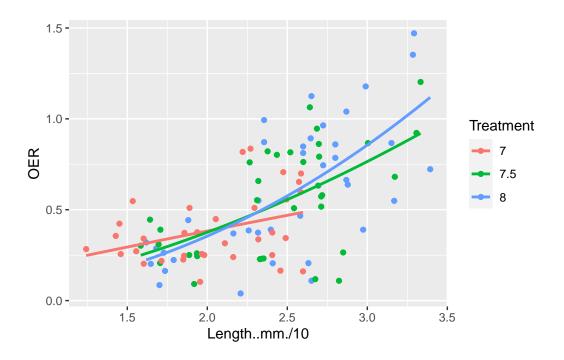
ValueStd.Errort-valuep-valueA.Treatment70.20368310.09228502.2071100.0296A.Treatment7.50.11149250.05005202.2275350.0281A.Treatment80.07870010.03345692.3522810.0206B.Treatment70.91075500.57938661.5719300.1191B.Treatment7.51.75327330.45097703.8877220.0002B.Treatment82.17234590.40680035.3400790.0000

```
Correlation:
              A.Trt7 A.T7.5 A.Trt8 B.Trt7 B.T7.5
A.Treatment7.5 0.000
A.Treatment8 0.000 0.000
B.Treatment7 -0.973 0.000 0.000
B.Treatment7.5 0.000 -0.987 0.000 0.000
B.Treatment8
               0.000 0.000 -0.990 0.000 0.000
Standardized residuals:
        Min
                               Med
                                            QЗ
                                                       Max
                    Q1
-2.40495512 -0.51428075 0.01691518 0.70238271 2.02514620
Residual standard error: 0.2406882
Degrees of freedom: 107 total; 101 residual
  fit3 <- nlme::gnls(OER~ A*(Length..mm./10)^B,
             params = list(A-1, B-1),
             data=bd_Pousse,
             na.action = na.exclude,
             start = list(A=1, B = .3)
  summary(fit3)
Generalized nonlinear least squares fit
  Model: OER ~ A * (Length..mm./10)^B
  Data: bd_Pousse
       AIC
               BIC
                     logLik
  2.943235 10.96172 1.528382
Coefficients:
      Value Std.Error t-value p-value
A 0.0956827 0.02339266 4.090289
                                 1e-04
B 1.9355590 0.24718721 7.830336
                                 0e+00
 Correlation:
  Α
B -0.985
Standardized residuals:
        Min
                               Med
                                                       Max
```

-2.50929987 -0.50686280 -0.01971863 0.66995680 2.11967734

```
Residual standard error: 0.2408001
Degrees of freedom: 107 total; 105 residual
  AIC(fit1,fit2,fit3)
     df
             AIC
fit1 5 5.582898
fit2 7 6.687948
fit3 3 2.943235
  BIC(fit1,fit2,fit3)
             BIC
     df
fit1 5 18.94704
fit2 7 25.39775
fit3 3 10.96172
  gg1 <- ggplot(data = bd_Pousse, aes(x=Length..mm./10, y=OER, color = Treatment))+
    geom_point()+
    #xlab("Dry weight") +
    #ylab("ORR")+
    #ylim(0,9)+
    #scale_color_discrete(name="Site")+
    #ggtitle ('ORR as a function of DW')+
    stat_smooth(
      method = 'nls',
      formula = y \sim A * x^B,
      method.args = list(start = c(A = 2.3, B = 2)),
      se = FALSE
    )
  #+
    #facet_wrap(~ Treatment)
  gg1
Warning: Removed 5 rows containing non-finite values (`stat_smooth()`).
```

Warning: Removed 5 rows containing missing values (`geom_point()`).



Generalized nonlinear least squares fit
 Model: IER ~ A * (Length..mm./10)^B
 Data: bd_Pousse
 AIC BIC logLik
 401.1693 414.5334 -195.5846

Coefficients:

Value Std.Error t-value p-value
A.Treatment7 0.4572221 0.1279428 3.573646 0.0005
A.Treatment7.5 0.5181582 0.1623990 3.190649 0.0019
A.Treatment8 0.5547207 0.1788415 3.101745 0.0025
B 1.9990503 0.3054852 6.543852 0.0000

Correlation:

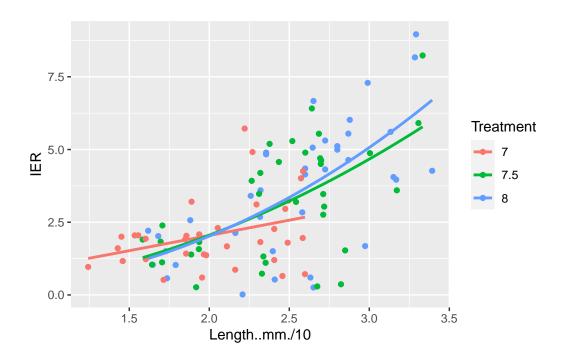
A.Trt7 A.T7.5 A.Trt8

```
A.Treatment7.5 0.867
               0.877 0.951
A.Treatment8
              -0.894 -0.970 -0.981
Standardized residuals:
      Min
                            Med
                                        QЗ
                                                  Max
-2.4498758 -0.5468184 0.1328319 0.6152599 2.2597997
Residual standard error: 1.534194
Degrees of freedom: 107 total; 103 residual
  fit2 <- nlme::gnls(IER~ A*(Length..mm./10)^B,
             params = list(A~Treatment-1, B~Treatment-1),
             data=bd Pousse,
             na.action = na.exclude,
             start = list(A=c(1,1,1), B=c(.3,.3,.3)))
  summary(fit2)
Generalized nonlinear least squares fit
  Model: IER ~ A * (Length..mm./10)^B
  Data: bd_Pousse
      AIC
              BIC
                    logLik
  403.072 421.7818 -194.536
Coefficients:
                  Value Std.Error t-value p-value
A.Treatment7
              1.0039269 0.5537211 1.813055 0.0728
A.Treatment7.5 0.5054045 0.2563311 1.971686 0.0514
              0.4206189 0.2003098 2.099842 0.0382
A.Treatment8
B.Treatment7
              1.0260338 0.6996277 1.466542 0.1456
B.Treatment7.5 2.0240970 0.5034159 4.020725 0.0001
B.Treatment8
             2.2657591 0.4513386 5.020087 0.0000
 Correlation:
              A.Trt7 A.T7.5 A.Trt8 B.Trt7 B.T7.5
A.Treatment7.5 0.000
               0.000 0.000
A.Treatment8
B.Treatment7 -0.975 0.000 0.000
B.Treatment7.5 0.000 -0.988 0.000 0.000
B.Treatment8 0.000 0.000 -0.991 0.000 0.000
```

```
Standardized residuals:
        Min
                    Q1
                                Med
                                             QЗ
                                                        Max
-2.45275038 -0.49878689 0.06754884 0.57154993 2.24491008
Residual standard error: 1.5342
Degrees of freedom: 107 total; 101 residual
  fit3 <- nlme::gnls(IER~ A*(Length..mm./10)^B,
             params = list(A~1, B~1),
             data=bd_Pousse,
             na.action = na.exclude,
             start = list(A=1, B = .3)
  summary(fit3)
Generalized nonlinear least squares fit
  Model: IER ~ A * (Length..mm./10)^B
  Data: bd_Pousse
       AIC
               BIC
                       logLik
  399.0157 407.0342 -196.5079
Coefficients:
     Value Std.Error t-value p-value
A 0.443513 0.1241451 3.572538
                                5e-04
B 2.169206 0.2784169 7.791214
                                0e+00
 Correlation:
 Α
B -0.987
Standardized residuals:
                             Med
       Min
                   Q1
                                         QЗ
                                                   Max
-2.5095101 -0.5885213 0.1244992 0.5942316 2.0986938
Residual standard error: 1.53268
Degrees of freedom: 107 total; 105 residual
  AIC(fit1,fit2,fit3)
             AIC
     df
fit1 5 401.1693
```

```
fit2 7 403.0720
fit3 3 399.0157
  BIC(fit1,fit2,fit3)
             BIC
     df
fit1 5 414.5334
fit2 7 421.7818
fit3 3 407.0342
  gg1 <- ggplot(data = bd_Pousse, aes(x=Length..mm./10, y=IER, color = Treatment))+
    geom_point()+
    #xlab("Dry weight") +
    #ylab("ORR")+
    #ylim(0,9)+
    #scale_color_discrete(name="Site")+
    #ggtitle ('ORR as a function of DW')+
    stat_smooth(
      method = 'nls',
      formula = y \sim A * x^B,
      method.args = list(start = c(A = 2.3, B = 2)),
      se = FALSE
    )
  #+
    #facet_wrap(~ Treatment)
  gg1
Warning: Removed 5 rows containing non-finite values (`stat_smooth()`).
```

Warning: Removed 5 rows containing missing values (`geom_point()`).



Generalized nonlinear least squares fit
 Model: ORR ~ A * (Length..mm./10)^B
 Data: bd_Pousse
 AIC BIC logLik
 -57.68407 -44.27341 33.84203

Coefficients:

Value Std.Error t-value p-value
A.Treatment7 0.2752024 0.05910500 4.656161 0.0000
A.Treatment7.5 0.3038002 0.07622111 3.985776 0.0001
A.Treatment8 0.3431002 0.08671211 3.956774 0.0001
B 0.0265632 0.26502546 0.100229 0.9204

Correlation:

A.Trt7 A.T7.5 A.Trt8

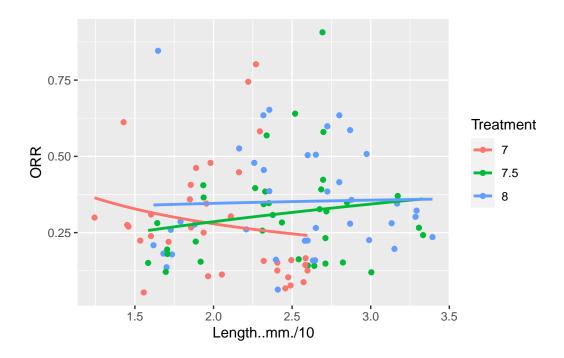
```
A.Treatment7.5 0.790
               0.812 0.871
A.Treatment8
              -0.858 -0.921 -0.946
Standardized residuals:
      Min
                            Med
                                        QЗ
                                                  Max
-1.5971008 -0.7673370 -0.1998771 0.4506859 3.2997067
Residual standard error: 0.1802482
Degrees of freedom: 108 total; 104 residual
  fit2 <- nlme::gnls(ORR~ A*(Length..mm./10)^B,
             params = list(A~Treatment-1, B~Treatment-1),
             data=bd Pousse,
             na.action = na.exclude,
             start = list(A=c(1,1,1), B=c(.3,.3,.3)))
  summary(fit2)
Generalized nonlinear least squares fit
  Model: ORR ~ A * (Length..mm./10)^B
  Data: bd_Pousse
       AIC
                 BIC
                       logLik
  -56.13407 -37.35915 35.06704
Coefficients:
                   Value Std.Error
                                      t-value p-value
A.Treatment7
               0.4116627 0.1433324 2.8720841 0.0050
A.Treatment7.5 0.2086449 0.0988228 2.1113028 0.0372
A.Treatment8
               0.3288236 0.1188672 2.7663114 0.0067
B.Treatment7 -0.5612090 0.5131991 -1.0935501 0.2767
B.Treatment7.5 0.4543274 0.5127255 0.8861026 0.3776
B.Treatment8
               0.0736379 0.3885703 0.1895098 0.8501
 Correlation:
              A.Trt7 A.T7.5 A.Trt8 B.Trt7 B.T7.5
A.Treatment7.5 0.000
A.Treatment8
               0.000 0.000
B.Treatment7 -0.950 0.000 0.000
B.Treatment7.5 0.000 -0.979 0.000 0.000
B.Treatment8 0.000 0.000 -0.974 0.000 0.000
```

```
Min
                  Q1
                             Med
                                         QЗ
                                                   Max
-1.5975417 -0.6982954 -0.3194695 0.4782606 3.2198221
Residual standard error: 0.179954
Degrees of freedom: 108 total; 102 residual
  fit3 <- nlme::gnls(ORR~ A*(Length..mm./10)^B,
             params = list(A~1, B~1),
             data=bd_Pousse,
             na.action = na.exclude,
             start = list(A=1, B = .3)
  summary(fit3)
Generalized nonlinear least squares fit
  Model: ORR ~ A * (Length..mm./10)^B
  Data: bd_Pousse
        AIC
                  BIC
                        logLik
  -59.13845 -51.09205 32.56922
Coefficients:
      Value Std.Error t-value p-value
A 0.2795575 0.06026682 4.638664 0.0000
B 0.1475814 0.24830764 0.594349 0.5535
 Correlation:
 Α
B -0.967
Standardized residuals:
                             Med
       Min
                   Q1
                                         QЗ
                                                   Max
-1.4112834 -0.8284690 -0.1847145 0.4589861 3.2276697
Residual standard error: 0.1806562
Degrees of freedom: 108 total; 106 residual
  AIC(fit1,fit2,fit3)
              AIC
     df
fit1 5 -57.68407
```

Standardized residuals:

```
fit2 7 -56.13407
fit3 3 -59.13845
  BIC(fit1,fit2,fit3)
              BIC
     df
fit1 5 -44.27341
fit2 7 -37.35915
fit3 3 -51.09205
  gg1 <- ggplot(data = bd_Pousse, aes(x=Length..mm./10, y=ORR, color = Treatment))+
    geom_point()+
    #xlab("Dry weight") +
    #ylab("ORR")+
    #ylim(0,9)+
    #scale_color_discrete(name="Site")+
    #ggtitle ('ORR as a function of DW')+
    stat_smooth(
      method = 'nls',
      formula = y \sim A * x^B,
      method.args = list(start = c(A = 2.3, B = 2)),
      se = FALSE
    )
  #+
    #facet_wrap(~ Treatment)
  gg1
Warning: Removed 4 rows containing non-finite values (`stat_smooth()`).
```

Warning: Removed 4 rows containing missing values (`geom_point()`).



Generalized nonlinear least squares fit
 Model: IRR ~ A * (Length..mm./10)^B
 Data: bd_Pousse
 AIC BIC logLik
 337.2855 350.6962 -163.6428

Coefficients:

Value Std.Error t-value p-value
A.Treatment7 1.5411751 0.3910295 3.941327 0.0001
A.Treatment7.5 1.9556583 0.5682653 3.441453 0.0008
A.Treatment8 1.9066430 0.5682767 3.355131 0.0011
B -0.1175519 0.3140424 -0.374319 0.7089

Correlation:

A.Trt7 A.T7.5 A.Trt8

```
A.Treatment7.5 0.786
               0.792 0.870
A.Treatment8
              -0.846 -0.929 -0.936
Standardized residuals:
      Min
                            Med
                                        QЗ
                                                  Max
-1.5330178 -0.7479955 -0.2680821 0.6905940 3.8442038
Residual standard error: 1.122033
Degrees of freedom: 108 total; 104 residual
  fit2 <- nlme::gnls(IRR~ A*(Length..mm./10)^B,
             params = list(A~Treatment-1, B~Treatment-1),
             data=bd Pousse,
             na.action = na.exclude,
             start = list(A=c(1,1,1), B=c(.3,.3,.3)))
  summary(fit2)
Generalized nonlinear least squares fit
  Model: IRR ~ A * (Length..mm./10)^B
  Data: bd_Pousse
      AIC
               BIC
                      logLik
  340.4051 359.1801 -163.2026
Coefficients:
                   Value Std.Error
                                      t-value p-value
A.Treatment7
               2.0351779 0.8846866 2.3004506 0.0235
A.Treatment7.5 1.5443301 0.7633024 2.0232218 0.0457
A.Treatment8
               1.8561279 0.8323219 2.2300600 0.0279
B.Treatment7 -0.5280379 0.6381486 -0.8274529 0.4099
B.Treatment7.5 0.1556260 0.5471325 0.2844394 0.7767
B.Treatment8
              -0.0873537 0.4889758 -0.1786463 0.8586
 Correlation:
              A.Trt7 A.T7.5 A.Trt8 B.Trt7 B.T7.5
A.Treatment7.5 0.000
A.Treatment8
               0.000 0.000
B.Treatment7 -0.950 0.000 0.000
B.Treatment7.5 0.000 -0.976 0.000 0.000
B.Treatment8 0.000 0.000 -0.972 0.000 0.000
```

```
Min
                   Q1
                             Med
                                         QЗ
                                                   Max
-1.3837406 -0.6710881 -0.2796251 0.7739305 3.7684779
Residual standard error: 1.128372
Degrees of freedom: 108 total; 102 residual
  fit3 <- nlme::gnls(IRR~ A*(Length..mm./10)^B,
             params = list(A~1, B~1),
             data=bd_Pousse,
             na.action = na.exclude,
             start = list(A=1, B = .3)
  summary(fit3)
Generalized nonlinear least squares fit
  Model: IRR ~ A * (Length..mm./10)^B
  Data: bd_Pousse
       AIC
               BIC
                       logLik
  335.4969 343.5433 -164.7484
Coefficients:
      Value Std.Error t-value p-value
A 1.6248086 0.4080025 3.982350 0.0001
B 0.0105338 0.2934046 0.035902 0.9714
 Correlation:
 Α
B -0.965
Standardized residuals:
       Min
                   Q1
                             Med
                                         QЗ
                                                   Max
-1.3592906 -0.8172054 -0.1691198 0.6802826 3.9294734
Residual standard error: 1.122834
Degrees of freedom: 108 total; 106 residual
  AIC(fit1,fit2,fit3)
             AIC
     df
fit1 5 337.2855
```

Standardized residuals:

```
fit2 7 340.4051
fit3 3 335.4969
  BIC(fit1,fit2,fit3)
             BIC
     df
fit1 5 350.6962
fit2 7 359.1801
fit3 3 343.5433
  gg1 <- ggplot(data = bd_Pousse, aes(x=Length..mm./10, y=IRR, color = Treatment))+
    geom_point()+
    #xlab("Dry weight") +
    #ylab("ORR")+
    #ylim(0,9)+
    #scale_color_discrete(name="Site")+
    #ggtitle ('ORR as a function of DW')+
    stat_smooth(
      method = 'nls',
      formula = y \sim A * x^B,
      method.args = list(start = c(A = 2.3, B = 2)),
      se = FALSE
    )
  #+
    #facet_wrap(~ Treatment)
  gg1
Warning: Removed 4 rows containing non-finite values (`stat_smooth()`).
```

Warning: Removed 4 rows containing missing values (`geom_point()`).

