Supplementary Material

¹H NMR Spectroscopy and MVA to Evaluate the Effects of Caulerpin-Based Diet on *Diplodus Sargus* Lipid Profiles

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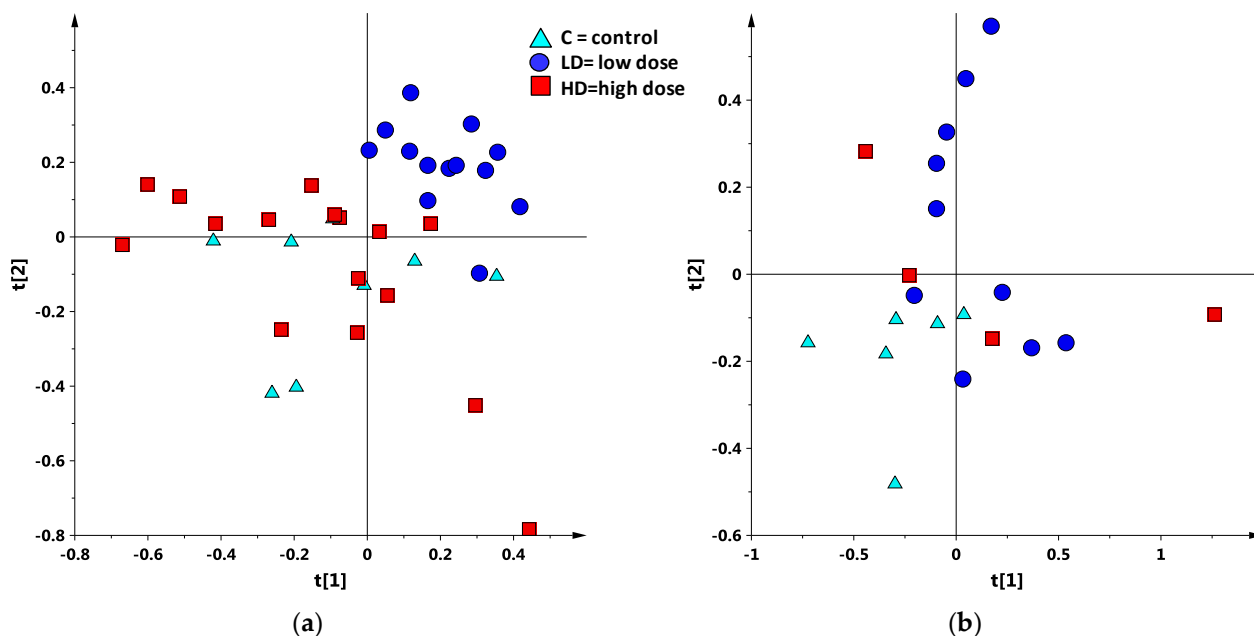
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Figure S1. (**a**) PCA (4 components, R²X=0.80, Q²=0.70) score plot obtained from ¹H NMR lipid extracts for experiment 1 (3 days treatment); (**b**) PCA (4 components, R²X=0.89, Q²=0.74) score plot for experiment 2 (10 days treatment).

FOLD CHANGE EXP-1 (37 samples)

Table S1. Fatty acid percentage calculated by integration of unbiased signals in the ^{1}H NMR spectra of lipid extracts for EXP-1 samples.

	%PUFA	%EPA	%DHA	%DUFA	%MUFA	%UFA	%SFA
C	23.23	3.75	12.25	6.32	22.37	48.08	51.92
HD	22.31	3.57	11.29	7.47	19.20	51.02	48.98
LD	24.13	3.46	10.11	8.01	14.00	53.86	46.14

Table S2. Ratio calculated from percentage of FA values.

	%PUFA	%EPA	%DHA	%DUFA	%MUFA	%UFA	%SFA
C/LD	0.963	1.083	1.212	0.789	1.598	0.893	1.125
C/HD	1.041	1.050	1.085	0.845	1.165	0.942	1.060

Table S3. Fold change (FC) values.

FC	%PUFA	%EPA	%DHA	%DUFA	%MUFA	%UFA	%SFA
C/LD	0.054	-0.115	-0.277	0.342	-0.676	0.164	-0.170
C/HD	-0.058	-0.070	-0.117	0.243	-0.221	0.085	-0.084

FOLD CHANGE EXP-2 (20 samples)

Table S4. Fatty acid percentage calculated by integration of unbiased signals in the ^{1}H NMR spectra of lipid extracts for EXP-2 samples.

	%PUFA	%EPA	%DHA	%DUFA	%MUFA	%UFA	%SFA
C	16.88	1.77	9.22	12.44	14.70	55.99	44.01
HD	16.52	2.48	8.89	11.78	19.88	51.82	48.18
LD	22.28	3.37	10.08	9.07	17.31	51.33	48.67

Table S5. Ratio calculated from percentage of FA values.

	%PUFA	%EPA	%DHA	%DUFA	%MUFA	%UFA	%SFA
C/LD	1.320	1.902	1.094	0.729	1.178	0.917	1.106
C/HD	1.349	1.359	1.134	0.770	0.871	0.991	1.010

Table S6. Fold change (FC) values.

FC	%PUFA	%EPA	%DHA	%DUFA	%MUFA	%UFA	%SFA
C/LD	-0.401	-0.9278	-0.130	0.455	-0.236	0.125	-0.145
C/HD	-0.432	-0.442	-0.181	0.377	0.199	0.014	-0.014

Table S7. One-way Anova results for EXP-1 samples.

> AnovaModel.2 <- aov(X.PUFA ~ trattamento, data=dsE7_TOT)

> summary(AnovaModel.2)
          Df Sum Sq Mean Sq F value Pr(>F)
trattamento 2 23.69 11.847   2.127 0.135
Residuals   34 189.33   5.568

```
> with(dsE7_TOT, numSummary(X.PUFA, groups=trattamento, statistics=c("mean",
+   "sd")))
      mean       sd data:n
C  23.23309 2.519209     8
HD 22.31423 2.485671    16
LD 24.12908 2.086125    13


Fit: aov(formula = X.PUFA ~ trattamento, data = dsE7_TOT)


Linear Hypotheses:
          Estimate Std. Error t value Pr(>|t|)
HD - C == 0   -0.9189    1.0218  -0.899   0.643
LD - C == 0    0.8960    1.0604   0.845   0.677
LD - HD == 0   1.8148    0.8811   2.060   0.113
(Adjusted p values reported -- single-step method)


> AnovaModel.5 <- aov(X.EPA ~ trattamento, data=dsE7_TOT)


> summary(AnovaModel.5)
            Df Sum Sq Mean Sq F value Pr(>F)
trattamento  2  0.410  0.2051   0.967   0.39
Residuals   34  7.211  0.2121


> with(dsE7_TOT, numSummary(X.EPA, groups=trattamento, statistics=c("mean", "sd")))
      mean        sd data:n
C  3.751333 0.3737482     8
HD 3.572989 0.4962436    16
LD 3.463528 0.4600527    13


Fit: aov(formula = X.EPA ~ trattamento, data = dsE7_TOT)


Linear Hypotheses:
          Estimate Std. Error t value Pr(>|t|)
HD - C == 0   -0.1783    0.1994  -0.894   0.646
LD - C == 0   -0.2878    0.2069  -1.391   0.355
LD - HD == 0  -0.1095    0.1720  -0.637   0.800
(Adjusted p values reported -- single-step method)




> AnovaModel.6 <- aov(X.DHA ~ trattamento, data=dsE7_TOT)


> summary(AnovaModel.6)
            Df Sum Sq Mean Sq F value Pr(>F)
trattamento  2  23.92  11.961   5.034 0.0122 *
Residuals   34  80.78   2.376
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
> with(dsE7_TOT, numSummary(X.DHA, groups=trattamento, statistics=c("mean", "sd")))
     mean      sd data:n
C  12.24687 1.093859     8
HD 11.28827 1.842136    16
LD 10.10589 1.338741    13
```

Multiple Comparisons of Means: Tukey Contrasts

Fit: aov(formula = X.DHA ~ trattamento, data = dsE7_TOT)

Linear Hypotheses:
```
           Estimate Std. Error t value Pr(>|t|)
HD - C == 0  -0.9586     0.6675  -1.436   0.3326
LD - C == 0  -2.1410     0.6927  -3.091   0.0107 *
LD - HD == 0 -1.1824     0.5756  -2.054   0.1141
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Adjusted p values reported -- single-step method)
```

```
> with(dsE7_TOT, numSummary(X.DUFA, groups=trattamento, statistics=c("mean",
+  "sd")))
     mean      sd data:n
C  6.315064 1.908038     8
HD 7.473351 2.257739    16
LD 8.005193 1.440296    13
```

        Simultaneous Tests for General Linear Hypotheses

Multiple Comparisons of Means: Tukey Contrasts

Fit: aov(formula = X.DUFA ~ trattamento, data = dsE7_TOT)

Linear Hypotheses:
```
           Estimate Std. Error t value Pr(>|t|)
HD - C == 0   1.1583     0.8363   1.385    0.358
LD - C == 0   1.6901     0.8679   1.947    0.140
LD - HD == 0  0.5318     0.7212   0.737    0.742
```

```
> AnovaModel.8 <- aov(X.MUFA ~ trattamento, data=dsE7_TOT)

> summary(AnovaModel.8)
            Df Sum Sq Mean Sq F value Pr(>F)
trattamento  2  383.5  191.75   3.892 0.0301 *
```

Residuals   34 1675.2   49.27

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

> with(dsE7_TOT, numSummary(X.MUFA, groups=trattamento, statistics=c("mean",
+   "sd")))
     mean   sd data:n
C 22.37170 8.244414   8
HD 19.19655 8.009060   16
LD 14.00173 4.446400   13

Multiple Comparisons of Means: Tukey Contrasts

Fit: aov(formula = X.MUFA ~ trattamento, data = dsE7_TOT)

Linear Hypotheses:
          Estimate Std. Error t value Pr(>|t|)
HD - C == 0   -3.175   3.039 -1.045   0.5529
LD - C == 0   -8.370   3.154 -2.654   0.0312 *
LD - HD == 0 -5.195   2.621 -1.982   0.1312

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Adjusted p values reported -- single-step method)

> AnovaModel.9 <- aov(X.UFA ~ trattamento, data=dsE7_TOT)

> summary(AnovaModel.9)
            Df Sum Sq Mean Sq F value Pr(>F)
trattamento 2 169.4   84.72   5.318 0.00978 **
Residuals   34 541.6   15.93

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

> with(dsE7_TOT, numSummary(X.UFA, groups=trattamento, statistics=c("mean", "sd")))
     mean   sd data:n
C 48.08015 5.023999   8
HD 51.01588 4.292175   16
LD 53.86400 2.717480   13

Multiple Comparisons of Means: Tukey Contrasts

Fit: aov(formula = X.UFA ~ trattamento, data = dsE7_TOT)

Linear Hypotheses:

```
        Estimate Std. Error t value Pr(>|t|)
HD - C == 0    2.936    1.728  1.699  0.21910
LD - C == 0    5.784    1.794  3.225  0.00756 **
LD - HD == 0   2.848    1.490  1.911  0.14999
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Adjusted p values reported -- single-step method)


> AnovaModel.10 <- aov(X.SFA ~ trattamento, data=dsE7_TOT)

> summary(AnovaModel.10)
            Df Sum Sq Mean Sq F value  Pr(>F)
trattamento  2  169.4   84.72   5.318 0.00978 **
Residuals   34  541.6   15.93
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1


> with(dsE7_TOT, numSummary(X.SFA, groups=trattamento, statistics=c("mean", "sd")))
      mean       sd data:n
C  51.91985 5.023999      8
HD 48.98412 4.292175     16
LD 46.13600 2.717480     13
```

          Simultaneous Tests for General Linear Hypotheses

Multiple Comparisons of Means: Tukey Contrasts

Fit: aov(formula = X.SFA ~ trattamento, data = dsE7_TOT)

Linear Hypotheses:

```
          Estimate Std. Error t value Pr(>|t|)
HD - C == 0   -2.936    1.728  -1.699  0.21908
LD - C == 0   -5.784    1.794  -3.225  0.00761 **
LD - HD == 0  -2.848    1.490  -1.911  0.15007
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Adjusted p values reported -- single-step method)
```


Table S8. One-way Anova results for EXP-2 samples.

          Simultaneous Tests for General Linear Hypotheses

Multiple Comparisons of Means: Tukey Contrasts

> with(DSL_M25_simca, numSummary(PUFA, groups=trattamento, statistics=c("mean", "sd")))

```
        mean       sd data:n
C 22.28495 1.526364      6
HD 16.51727 3.000546     4
LD 16.87872 2.283117    10
```

Fit: aov(formula = X.**PUFA** ~ trattamento, data = DSL_M25_simca)

Linear Hypotheses:

```
           Estimate Std. Error t value Pr(>|t|)
HD - C == 0  -5.7677    1.4482  -3.983  0.00265 **
LD - C == 0  -5.4062    1.1586  -4.666  < 0.001 ***
LD - HD == 0  0.3615    1.3273   0.272  0.95973
---
```

> with(DSL_M25_simca, numSummary(**EPA**, groups=trattamento, statistics=c("mean", "sd")))

```
       mean        sd data:n
C 3.366663 0.5306937      6
HD 2.478220 0.5833759     4
LD 1.769966 0.2715293    10
```

Fit: aov(formula = **EPA** ~ trattamento, data = DSL_M25_simca)

Linear Hypotheses:

```
           Estimate Std. Error t value Pr(>|t|)
HD - C == 0  -0.8884    0.2753  -3.227  0.0128 *
LD - C == 0  -1.5967    0.2203  -7.249  <0.001 ***
LD - HD == 0 -0.7083    0.2523  -2.807  0.0307 *
```

> AnovaModel.4 <- aov(**DHA** ~ trattamento, data=DSL_M25_simca)

```
            Df Sum Sq Mean Sq F value Pr(>F)
trattamento  2   4.17   2.083   0.449  0.646
Residuals   17  78.87   4.639
```

> with(DSL_M25_simca, numSummary(**DHA**, groups=trattamento,
+   statistics=c("mean", "sd")))

```
        mean       sd data:n
C 10.081935 1.886349      6
HD  8.891368 1.876738     4
LD  9.215016 2.369036    10
```

Multiple Comparisons of Means: Tukey Contrasts

Fit: aov(formula = X.**DHA** ~ trattamento, data = DSL_M25_simca)

Linear Hypotheses:

```
           Estimate Std. Error t value Pr(>|t|)
HD - C == 0  -1.1906    1.3903  -0.856   0.672
```

```
LD - C == 0  -0.8669    1.1123  -0.779   0.719
LD - HD == 0   0.3236    1.2743   0.254   0.965
(Adjusted p values reported -- single-step method)
```

```
> AnovaModel.5 <- aov(DUFA ~ trattamento, data=DSL_M25_simca)

> summary(AnovaModel.5)
        Df Sum Sq Mean Sq F value Pr(>F)
trattamento  2  43.69  21.844   2.827 0.0871 .
Residuals   17 131.34   7.726
---

> with(DSL_M25_simca, numSummary(DUFA, groups=trattamento,
+   statistics=c("mean", "sd")))
      mean       sd data:n
C  9.069537 1.927444      6
HD 11.781788 5.177614      4
LD 12.436267 1.895603     10

Fit: aov(formula = DUFA ~ trattamento, data = DSL_M25_simca)

Linear Hypotheses:
        Estimate Std. Error t value Pr(>|t|)
HD - C == 0   2.7123     1.7942   1.512   0.3084
LD - C == 0   3.3667     1.4353   2.346   0.0757 .
LD - HD == 0   0.6545     1.6444   0.398   0.9162
---


> AnovaModel.7 <- aov(MUFA ~ trattamento, data=DSL_M25_simca)

> summary(AnovaModel.7)
        Df Sum Sq Mean Sq F value Pr(>F)
trattamento  2   82.1   41.06   0.617  0.551
Residuals   17 1131.9   66.58

> with(DSL_M25_simca, numSummary(MUFA, groups=trattamento,
+   statistics=c("mean", "sd")))
    mean       sd data:n
C 17.31323  4.851955      6
HD 19.88096 14.671742      4
LD 14.69904  6.397993     10

Fit: aov(formula = MUFA ~ trattamento, data = DSL_M25_simca)

Linear Hypotheses:
        Estimate Std. Error t value Pr(>|t|)
HD - C == 0   2.568      5.267   0.488   0.877
```

LD - C == 0 -2.614   4.214 -0.620   0.810
LD - HD == 0 -5.182   4.827 -1.073   0.541

> AnovaModel.12 <- aov(X.**UFA** ~ trattamento, data=DSL_M25_simca)

> summary(AnovaModel.12)
        Df Sum Sq Mean Sq F value Pr(>F)
trattamento 2 100.0   49.98   2.107 0.152
Residuals   17 403.2   23.72

> with(DSL_M25_simca, numSummary(X.**UFA**, groups=trattamento,
+   statistics=c("mean", "sd")))
     mean   sd data:n
C 51.33229 2.776968   6
HD 51.81999 7.162467   4
LD 55.98598 4.839500   10

Fit: aov(formula = X.UFA ~ trattamento, data = DSL_M25_simca)

Linear Hypotheses:
        Estimate Std. Error t value Pr(>|t|)
HD - C == 0   0.4877   3.1438 0.155   0.987
LD - C == 0   4.6537   2.5150 1.850   0.182
LD - HD == 0   4.1660   2.8813 1.446   0.339

> AnovaModel.10 <- aov(X.**SFA** ~ trattamento, data=DSL_M25_simca)

> summary(AnovaModel.10)
        Df Sum Sq Mean Sq F value Pr(>F)
trattamento 2 100.0   49.98   2.107 0.152
Residuals   17 403.2   23.72

> with(DSL_M25_simca, numSummary(X.**SFA**, groups=trattamento,
+   statistics=c("mean", "sd")))
     mean   sd data:n
C 48.66771 2.776968   6
HD 48.18001 7.162467   4
LD 44.01402 4.839500   10

Fit: aov(formula = X.SFA ~ trattamento, data = DSL_M25_simca)

Linear Hypotheses:
        Estimate Std. Error t value Pr(>|t|)
HD - C == 0   -0.4877   3.1438 -0.155   0.987
LD - C == 0   -4.6537   2.5150 -1.850   0.182
LD - HD == 0   -4.1660   2.8813 -1.446   0.338