

Inorganic carbon and nitrogen assimilation in cellular compartments of a benthic kleptoplastic foraminifer

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Supplementary material 1: Results of the different post-hoc Tukey multiple comparisons tests following the linear mixed-effects model analyses.

Note: “4, 8, 12, 20 and dark” in the columns “Time points” correspond to the time points 4, 8, 12, and 20 h of the Experiment 1, and to the time point 8 h of the Experiment 2, respectively.

- $\delta^{13}\text{C}$ cytoplasm

Simultaneous Tests for General Linear Hypotheses

Multiple Comparisons of Means: Tukey Contrasts

Fit: lme.formula(fixed = Y ~ time, data = mydata, random = ~1 | Plotb)

Linear Hypotheses:

Time points	Estimate	Std. Error	z value	Pr(> z)	
20 - 12 == 0	45.6	24.0	1.90	0.3166	
4 - 12 == 0	-32.7	24.3	-1.35	0.6619	
8 - 12 == 0	31.3	24.0	1.31	0.6876	
dark - 12 == 0	-71.9	25.3	-2.85	0.0357	*
4 - 20 == 0	-78.3	24.0	-3.27	0.0095	**
8 - 20 == 0	-14.3	23.6	-0.60	0.9746	
dark - 20 == 0	-117.5	24.9	-4.71	<0.001	***
8 - 4 == 0	64.0	24.0	2.67	0.0585	.
dark - 4 == 0	-39.2	25.3	-1.55	0.5289	
dark - 8 == 0	-103.2	24.9	-4.14	<0.001	***

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

- $\delta^{15}\text{N}$ cytoplasm

Simultaneous Tests for General Linear Hypotheses

Multiple Comparisons of Means: Tukey Contrasts

Fit: lme.formula(fixed = Z ~ time, data = mydata, random = ~1 | Plotb)

Linear Hypotheses:

Time points	Estimate	Std. Error	z value	Pr(> z)	
20 - 12 == 0	137.4	102.8	1.34	0.6677	
4 - 12 == 0	-201.5	104.2	-1.93	0.2991	
8 - 12 == 0	91.6	102.8	0.89	0.9002	
dark - 12 == 0	638.7	108.5	5.88	<0.001	***
4 - 20 == 0	-339.0	102.8	-3.30	0.0085	**
8 - 20 == 0	-45.8	101.3	-0.45	0.9914	
dark - 20 == 0	501.2	107.1	4.68	<0.001	***
8 - 4 == 0	293.1	102.8	2.85	0.0350	*
dark - 4 == 0	840.2	108.5	7.74	<0.001	***
dark - 8 == 0	547.1	107.1	5.11	<0.001	***

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

- $\delta^{13}\text{C}$ electron-opaque bodies

Simultaneous Tests for General Linear Hypotheses

Multiple Comparisons of Means: Tukey Contrasts

Fit: lme.formula(fixed = C ~ time, data = mydata, random = ~1 | Plotb)

Linear Hypotheses:

Time points	Estimate	Std. Error	z value	Pr(> z)	
20 - 12 == 0	132.9	51.8	2.56	0.0768	.
4 - 12 == 0	-49.5	50.0	-0.99	0.8598	
8 - 12 == 0	92.1	49.6	1.86	0.3406	
dark - 12 == 0	-75.3	49.8	-1.51	0.5547	
4 - 20 == 0	-182.4	53.7	-3.40	0.0061	**
8 - 20 == 0	-40.8	53.3	-0.77	0.9404	
dark - 20 == 0	-208.2	53.5	-3.89	<0.001	***
8 - 4 == 0	141.6	51.5	2.75	0.0475	*
dark - 4 == 0	-25.8	51.8	-0.50	0.9875	
dark - 8 == 0	-167.4	51.4	-3.26	0.0098	**

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

- $\delta^{15}\text{N}$ electron-opaque bodies

Simultaneous Tests for General Linear Hypotheses

Multiple Comparisons of Means: Tukey Contrasts

Fit: lme.formula(fixed = N ~ time, data = mydata, random = ~1 | Plotb)

Linear Hypotheses:

Time points	Estimate	Std. Error	z value	Pr(> z)	
20 - 12 == 0	-792	689	-1.15	0.780	
4 - 12 == 0	1133	685	1.65	0.462	
8 - 12 == 0	48	683	0.07	1.000	
dark - 12 == 0	882	684	1.29	0.698	
4 - 20 == 0	1925	694	2.78	0.044	*
8 - 20 == 0	840	692	1.21	0.743	
dark - 20 == 0	1674	692	2.42	0.111	
8 - 4 == 0	-1085	688	-1.58	0.512	
dark - 4 == 0	-251	688	-0.36	0.996	
dark - 8 == 0	834	687	1.21	0.743	

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

- $\delta^{13}\text{C}$ fibrillar vesicles

Simultaneous Tests for General Linear Hypotheses

Multiple Comparisons of Means: Tukey Contrasts

Fit: lme.formula(fixed = C ~ time, data = mydata, random = ~1 | Plotb)

Linear Hypotheses:

Time points	Estimate	Std. Error	z value	Pr(> z)	
20 - 12 == 0	64.82	63.91	1.01	0.8460	
4 - 12 == 0	71.09	86.30	0.82	0.9217	
8 - 12 == 0	-9.74	62.25	-0.16	0.9999	
dark - 12 == 0	-208.71	63.86	-3.27	0.0094	**
4 - 20 == 0	6.27	88.19	0.07	1.0000	
8 - 20 == 0	-74.57	64.84	-1.15	0.7759	
dark - 20 == 0	-273.53	66.39	-4.12	<0.001	***
8 - 4 == 0	-80.83	86.99	-0.93	0.8833	
dark - 4 == 0	-279.80	88.16	-3.17	0.0125	*
dark - 8 == 0	-198.97	64.80	-3.07	0.0177	*

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

- $\delta^{15}\text{N}$ fibrillar vesicles

Simultaneous Tests for General Linear Hypotheses

Multiple Comparisons of Means: Tukey Contrasts

Fit: lme.formula(fixed = N ~ time, data = mydata, random = ~1 | Plotb)

Linear Hypotheses:

Time points	Estimate	Std. Error	z value	Pr(> z)	
20 - 12 == 0	324	197	1.64	0.460	
4 - 12 == 0	-519	293	-1.77	0.383	
8 - 12 == 0	-218	190	-1.15	0.774	
dark - 12 == 0	1469	197	7.45	<0.001	***
4 - 20 == 0	-843	301	-2.80	0.039	*
8 - 20 == 0	-542	202	-2.69	0.053	.
dark - 20 == 0	1145	208	5.49	<0.001	***
8 - 4 == 0	301	296	1.01	0.844	
dark - 4 == 0	1988	301	6.60	<0.001	***
dark - 8 == 0	1688	202	8.36	<0.001	***

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

- $\delta^{13}\text{C}$ lipid droplets

Simultaneous Tests for General Linear Hypotheses

Multiple Comparisons of Means: Tukey Contrasts

Fit: lme.formula(fixed = C ~ time, data = mydata, random = ~1 | Plotb)

Linear Hypotheses:

Time points	Estimate	Std. Error	z value	Pr(> z)
20 - 12 == 0	148.4	64.6	2.30	0.1452
4 - 12 == 0	-20.4	57.7	-0.35	0.9966
8 - 12 == 0	76.1	57.5	1.32	0.6756
dark - 12 == 0	-87.7	65.3	-1.34	0.6632
4 - 20 == 0	-168.8	64.9	-2.60	0.0699 .
8 - 20 == 0	-72.3	64.8	-1.11	0.7979
dark - 20 == 0	-236.1	71.9	-3.29	0.0089 **
8 - 4 == 0	96.5	57.9	1.67	0.4522
dark - 4 == 0	-67.3	65.6	-1.03	0.8429
dark - 8 == 0	-163.8	65.5	-2.50	0.0899 .

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