## D13C - Initial Analysis

### Species Differences

#### Boxplot

* Most negative d13C (highest MR):
  + E. Antarctica
  + G. braueri
  + K. andersoni
* Least negative d13C (lowest MR):
  + E. carlsbergi
  + G. nicholsi
  + P. bolini

#### Kruskal-Wallis Test

* Significant difference between species.
  + Chi-squared = 82.414
  + Df = 5
  + p-value = 2.61 x 10^-16

#### Dunn Test (from dunn.test package)

* Bonferroni correction

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
|  | E. antarctica | E. carlsbergi | G. braueri | G. nicholsi | P. bolini | K. anderssoni |
| E. Antarctica |  | <0.01 | N/S | <0.01 | <0.01 | N/S |
| E. carlsbergi | <0.01 |  | <0.01 | N/S | N/S | <0.01 |
| G. braueri | N/S | <0.01 |  | <0.01 | <0.01 | N/S |
| G. nicholsi | <0.01 | N/S | <0.01 |  | N/S | <0.01 |
| P. bolini | <0.01 | N/S | <0.01 | N/S |  | <0.05 |
| K. anderssoni | N/S | <0.01 | N/S | <0.01 | <0.05 |  |

* Supports three groups as seen in boxplot

### Log10\_Weight vs. d13C

#### Plot

* Decrease in d13C with weight with E. carlsbergi and G. nicholsi as outliers.
  + Decrease in mass specific metabolic rate with weight (expected).

#### Spearman’s Rank Test

* No significant correlation between d13C and log10\_Weight.
  + Rho = -0.0292
  + P = 0.7637

### D18O vs. d13C

#### Plot

* Decrease in d13C with increase in d18O. H
  + Higher metabolic rate at lower temperatures (counter-intuative).

#### Spearman’s Rank Test

* Significant negative correlation.
  + Rho = -0.6721
  + P = 1.695 x 10^-15

### D13C with Depth

* Nothing interesting on the metabolic side.
* E. carlsbergi and G. braueri mostly caught shallow (>250m).

## D13C – Sherwood & Rose Comparison

### Species Comparisons

* E. Antarctica – same value as *Chaetodon ulietensis* (Pacific double-saddle butterflyfish, Chaetodontidae).
* G. braueri – same value as *Oncorhynchus nerka* (sockeye salmon, Salmonidae).
* K. andersoni – same value as *Maena maena* (now *Spicara maena,* blotched picarel, Centracanthidae).
* E. carlsbergi – same values as *Clupea pallasii* (Pacific herring, Clupeidae).
* G. nicholsi – same values as *Beryx splendens* (splendid alfonsino, Berycidae) and *Osmerus mordax* (rainbow smelt, Osmeridae).
* P. bolini – between *Centrolophus niger* (rudderfish, Centrolophidae) and *Merluccius merluccius* (European hake, Gadidae).
* Possible action?: run data against known RMR and compare.

### K\_caud vs. d13C

#### Spearman’s Rank Analysis

* Significant negative correlation between K\_caud and d13C.
  + Rho = -0.6000
  + P = 2.04 x 10^-7

#### Bayesian Analysis (without n as a random factor)

* P. bolini (just), K. andersoni, G. braueri and E. Antarctica all plot just outside 95% HDPI.
  + Lower d13C than expected given K\_caud.
  + High metabolic rate for activity level.
  + C. maderensis is also below the line.

### D18O vs. d13C

#### Spearman’s Rank Analysis

* Significant positive relationship between d18O and d13C.
  + Increasing MR with increasing temperature.

#### Bayesian Analysis (without n as a random factor)

* Higher d18O than Sherwood & Rose fishes.
  + At lower temperatures.
* G. braueri, E. carlsbergi and P. bolini are all plotting within 95% HDPI.
* K. andersoni, E. Antarctica and G. braueri all plotting below the line.
  + Lower d13C than expected given d18O.
  + High metabolic rate for temperature.
  + C. maderensis just outside HDPI.
  + Not metabolic cold adaptation, or it would affect all of them?
* Possible action?: look into metabolic cold adaptation.
* Note: cannot do against weights as they are max. weights and missing for many species.

## M-Values – Initial Analysis

Action: do again with replicates

* 0.14 – 0.30
  + 95% HDI range of 10,000 replicates: 0.01-0.04

### Species Differences

#### Boxplot

* E. Antarctica, G. braueri and K. andersoni all have high M values.
* G. nicholsi and E. carlsbergi have low M values, with P. bolini straddling but on lower end.

#### Kruskal-Wallis Test

* Repeated 1000 times
* Significant difference between species.
  + Chi-squared = 49.82 (95% HDI 39.84 – 61.35)
  + Df = 5
  + P = 2.44 x10^-9 (95% HDI 6.43 x 10^-13 – 1.01 x 10^-07)

#### Dunn Test

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
|  | E. antarctica | E. carlsbergi | G. braueri | G. nicholsi | P. bolini | K. anderssoni |
| E. Antarctica |  | <0.01 | N/S | <0.01 | <0.01 | N/S |
| E. carlsbergi | <0.01 |  | <0.01 | N/S | N/S | <0.01 |
| G. braueri | N/S | <0.01 |  | <0.01 | <0.01 | N/S |
| G. nicholsi | <0.01 | N/S | <0.01 |  | N/S | <0.01 |
| P. bolini | <0.01 | N/S | <0.01 | N/S |  | <0.05 |
| K. anderssoni | N/S | <0.01 | N/S | <0.01 | <0.05 |  |

* Supports three groups as seen in boxplot

With multiple repeats

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
|  | E. antarctica | E. carlsbergi | G. braueri | G. nicholsi | P. bolini | K. anderssoni |
| E. Antarctica |  | <0.05 | N/S | <0.01 | <0.01 | N/S |
| E. carlsbergi | <0.05 |  | <0.1 | N/S | N/S | <0.01 |
| G. braueri | N/S | <0.01 |  | <0.01 | <0.02 | N/S |
| G. nicholsi | <0.01 | N/S | <0.01 |  | N/S | <0.01 |
| P. bolini | <0.01 | N/S | <0.02 | N/S |  | <0.01 |
| K. anderssoni | N/S | <0.01 | N/S | <0.01 | <0.01 |  |

### Log10\_Weight vs. M

* Action: repeat with multiples

#### Plot

* No discernible relationship.

#### Spearman’s Rank Test

* No significant correlation between M and log10\_Weight.
  + Rho = 0.0118276
  + P = 0.9033

## Temperature – Initial Analysis

* -3.43 – 3.88 C
  + 95% HDPI of 10,000 replicates: 1.17-1.34
* Seems low – may be due to parameters being derived from cod data.

### Species Differences

#### Boxplot

* E. carlsbergi & G. nicholsi experiencing highest temperatures.
* G. braueri experiencing low temperatures.
* E. Antarctica, K. andersoni and P. bolini straddling.

#### Repeated K-W Test

* Temperature significantly different with species.
  + Chi-squared = 50.81 (95% HDI 39.62-60.45)
  + P-value = 1.65x 10^-9 (95% HDI 2.54 x 10^-13 – 1.00 x 10^-7

### M vs. Temperature

#### Plot

* Negative relationship between M and temperature.

#### Repeated Regression

* Significant relationship between M and temperature.
  + R-squared = 0.21 (95% HDI 0.11-0.30).
  + P-value = 7.65 x 10^-6 (95% HDI 2.19 x 10^-11 – 1.68 x 10^-4)

Action: complete Spearman’s test

Possible action: look into Bayesian with data with error points