Principal Component Analysis

Dataset otu table

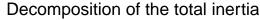
This dataset contains 151 individuals and 50 variables.						
## Warning: package 'FactoMineR' was built under R version 3.5.2						
1. Study of the outliers						
The analysis of the graphs does not detect any outlier.						

2. Inertia distribution

The inertia of the first dimensions shows if there are strong relationships between variables and suggests the number of dimensions that should be studied.

The first two dimensions of analyse express 15.88% of the total dataset inertia; that means that 15.88% of the individuals (or variables) cloud total variability is explained by the plane. This is a very small percentage and the first plane represents a small part of the data variability. This value is greater than the reference value that equals 9.47%, the variability explained by this plane is thus significant (the reference value is the 0.95-quantile of the inertia percentages distribution obtained by simulating 1726 data tables of equivalent size on the basis of a normal distribution).

From these observations, it is important to also interpret the dimensions greater or equal to the third one.



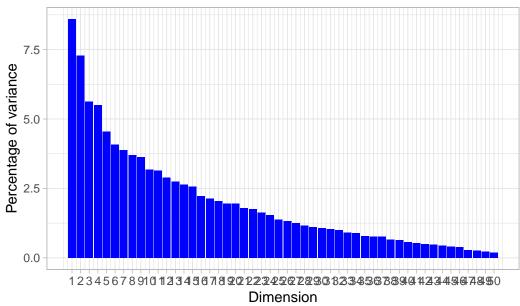


Figure 2 - Decomposition of the total inertia

An estimation of the right number of axis to interpret suggests to restrict the analysis to the description of the first 11 axis. These axis present an amount of inertia greater than those obtained by the 0.95-quantile of random distributions (53.14% against 41.8%). This observation suggests that only these axis are carrying a real information. As a consequence, the description will stand to these axis.

3. Description of the plane 1:2

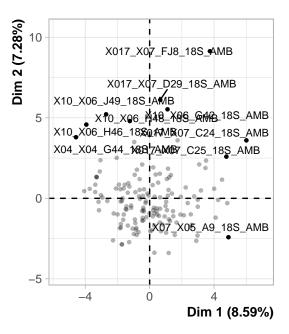


Figure 3.1 - Individuals factor map (PCA) The labeled individuals are those with the higher contribution to the plane construction.

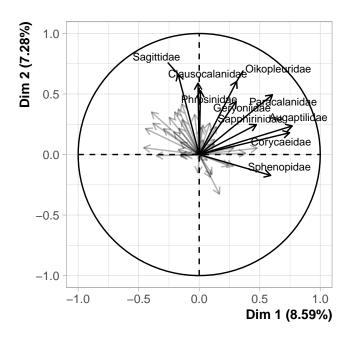


Figure 3.2 - Variables factor map (PCA) The labeled variables are those the best shown on the plane.

The dimension 1 opposes individuals such as $X017_X07_FJS_18S_AMB$, $X10_X06_G42_18S_AMB$, $X017_X07_C24_18S_AMB$, $X017_X07_D29_18S_AMB$, $X017_X07_C25_18S_AMB$ and $X07_X05_A9_18S_AMB$ (to the right of the graph, characterized by a strongly positive coordinate on the axis) to individuals such as $X10_X06_J49_18S_AMB$, $X10_X06_H46_18S_AMB$, $X04_X04_G44_18S_AMB$ and $X10_X06_H48_18S_AMB$ (to the left of the graph, characterized by a strongly negative coordinate on the axis).

The group in which the individual $X07_X05_A9_18S_AMB$ stands (characterized by a positive coordinate on the axis) is sharing:

- high values for variables like Sapphirinidae, Corycaeidae, Sphenopidae, Acartiidae, Temoridae, Doliolidae, Augaptilidae, Paracalanidae, Eupronoidae and Euphausiidae (variables are sorted from the strongest).
- low values for the variables Candaciidae, Subeucalanidae, Sagittidae, Sphaeronectidae, Hippolytidae, Squillidae, Phrosinidae, Clausocalanidae and Processidae (variables are sorted from the weakest).

The group in which the individuals $X017_X07_FJ8_18S_AMB$, $X10_X06_G42_18S_AMB$, $X017_X07_C24_18S_AMB$, $X017_X07_D29_18S_AMB$ and $X017_X07_C25_18S_AMB$ stand (characterized by a positive coordinate on the axis) is sharing:

• high values for variables like Geryoniidae, Oikopleuridae, Paracalanidae, Sagittidae, Augaptilidae, Clausocalanidae, Aetideidae, Phrosinidae, Halocyprididae and Corycaeidae (variables are sorted from the strongest).

The group in which the individuals $X10_X06_J49_18S_AMB$, $X10_X06_H46_18S_AMB$, $X04_X04_G44_18S_AMB$ and $X10_X06_H48_18S_AMB$ stand (characterized by a negative coordinate on the axis) is sharing:

- high values for variables like *Processidae*, *Callianassidae*, *Coenobitidae*, *Squillidae*, *Centropagidae*, *Candaciidae*, *Palaemonidae*, *Creseidae*, *Sagittidae* and *Subeucalanidae* (variables are sorted from the strongest).
- low values for the variables Augaptilidae, Corycaeidae, Doliolidae, Sphenopidae, Paracalanidae, Hormathiidae, Acartiidae and Euphausiidae (variables are sorted from the weakest).

The group 4 (characterized by a negative coordinate on the axis) is sharing:

• high values for the variable Calanidae.

• low values for variables like Paracalanidae, Sagittidae, Corycaeidae, Temoridae, Sapphirinidae, Clausocalanidae, Augaptilidae, Phrosinidae, Creseidae and Geryoniidae (variables are sorted from the weakest).

The dimension 2 opposes individuals such as $X017_X07_FJS_18S_AMB$, $X10_X06_G42_18S_AMB$, $X017_X07_C24_18S_AMB$, $X017_X07_D29_18S_AMB$, $X10_X06_J49_18S_AMB$, $X017_X07_C25_18S_AMB$, $X10_X06_H46_18S_AMB$, $X04_X04_G44_18S_AMB$ and $X10_X06_H48_18S_AMB$ (to the top of the graph, characterized by a strongly positive coordinate on the axis) to individuals such as ** and $X07_X05_A9_18S_AMB$ (to the bottom of the graph, characterized by a strongly negative coordinate on the axis).

The group in which the individuals $X10_X06_J49_18S_AMB$, $X10_X06_H46_18S_AMB$, $X04_X04_G44_18S_AMB$ and $X10_X06_H48_18S_AMB$ stand (characterized by a positive coordinate on the axis) is sharing :

- high values for variables like *Processidae*, *Callianassidae*, *Coenobitidae*, *Squillidae*, *Centropagidae*, *Candaciidae*, *Palaemonidae*, *Creseidae*, *Sagittidae* and *Subeucalanidae* (variables are sorted from the strongest).
- low values for the variables Augaptilidae, Corycaeidae, Doliolidae, Sphenopidae, Paracalanidae, Hormathiidae, Acartiidae and Euphausiidae (variables are sorted from the weakest).

The group in which the individuals $X017_X07_FJ8_18S_AMB$, $X10_X06_G42_18S_AMB$, $X017_X07_C24_18S_AMB$, $X017_X07_D29_18S_AMB$ and $X017_X07_C25_18S_AMB$ stand (characterized by a positive coordinate on the axis) is sharing:

• high values for variables like Geryoniidae, Oikopleuridae, Paracalanidae, Sagittidae, Augaptilidae, Clausocalanidae, Aetideidae, Phrosinidae, Halocyprididae and Corycaeidae (variables are sorted from the strongest).

The group 3 (characterized by a negative coordinate on the axis) is sharing:

- high values for the variable *Calanidae*.
- low values for variables like Paracalanidae, Sagittidae, Corycaeidae, Temoridae, Sapphirinidae, Clausocalanidae, Augaptilidae, Phrosinidae, Creseidae and Geryoniidae (variables are sorted from the weakest).

The group in which the individual $X07_X05_A9_18S_AMB$ stands (characterized by a negative coordinate on the axis) is sharing:

- high values for variables like Sapphirinidae, Corycaeidae, Sphenopidae, Acartiidae, Temoridae, Doliolidae, Augaptilidae, Paracalanidae, Eupronoidae and Euphausiidae (variables are sorted from the strongest).
- low values for the variables Candaciidae, Subeucalanidae, Sagittidae, Sphaeronectidae, Hippolytidae, Squillidae, Phrosinidae, Clausocalanidae and Processidae (variables are sorted from the weakest).

4. Description of the plane 3:4

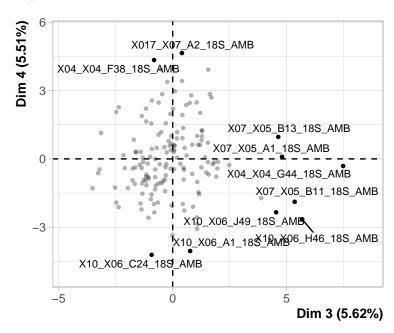


Figure 4.1 - Individuals factor map (PCA) The labeled individuals are those with the higher contribution to the plane construction.

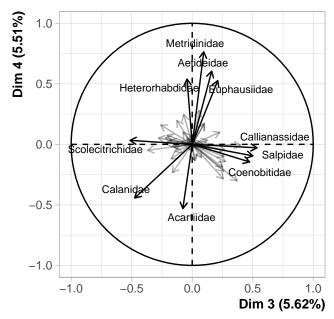


Figure 4.2 - Variables factor map (PCA) The labeled variables are those the best shown on the plane.

The dimension 3 opposes individuals such as $X04_X04_G44_18S_AMB$, $X10_X06_H46_18S_AMB$, $X10_X06_J49_18S_AMB$, $X07_X05_B13_18S_AMB$, $X07_X05_B11_18S_AMB$ and $X07_X05_A1_18S_AMB$ (to the right of the graph, characterized by a strongly positive coordinate on the axis) to individuals such as $X10_X06_C24_18S_AMB$ and $X10_X06_A1_18S_AMB$ (to the left of the graph, characterized by a strongly negative coordinate on the axis).

The group in which the individuals $X04_X04_G44_18S_AMB$, $X10_X06_H46_18S_AMB$, $X10_X06_J49_18S_AMB$,

X07_X05_B13_18S_AMB, X07_X05_B11_18S_AMB and X07_X05_A1_18S_AMB stand (characterized by a positive coordinate on the axis) is sharing:

- high values for variables like Coenobitidae, Callianassidae, Lycaeidae, Salpidae, Centropagidae, Temoridae, Creseidae, Pandalidae, Aequoreidae and Squillidae (variables are sorted from the strongest).
- low values for the variables Calanidae and Scolecitrichidae (variables are sorted from the weakest).

The group in which the individuals $X10_X06_C24_18S_AMB$ and $X10_X06_A1_18S_AMB$ stand (characterized by a negative coordinate on the axis) is sharing:

- high values for the variables Calanidae, Acartiidae, Hippolytidae, Sphaeronectidae and Palaemonidae (variables are sorted from the strongest).
- low values for the variables Metridinidae, Euphausiidae, Aetideidae, Heterorhabdidae, Coenobitidae, Pandalidae, Callianassidae and Typhloscolecidae (variables are sorted from the weakest).

The dimension 4 opposes individuals such as $X017_X07_A2_18S_AMB$ and $X04_X04_F38_18S_AMB$ (to the top of the graph, characterized by a strongly positive coordinate on the axis) to individuals such as $X10_X06_C24_18S_AMB$ and $X10_X06_A1_18S_AMB$ (to the bottom of the graph, characterized by a strongly negative coordinate on the axis).

The group in which the individuals $X017_X07_A2_18S_AMB$ and $X04_X04_F38_18S_AMB$ stand (characterized by a positive coordinate on the axis) is sharing:

- high values for the variables *Metridinidae*, *Aetideidae*, *Euphausiidae*, *Heterorhabdidae* and *Typhloscole-cidae* (variables are sorted from the strongest).
- low values for the variables *Calanidae*, *Acartiidae*, *Creseidae* and *Doliolidae* (variables are sorted from the weakest).

The group in which the individuals $X10_X06_C24_18S_AMB$ and $X10_X06_A1_18S_AMB$ stand (characterized by a negative coordinate on the axis) is sharing:

- high values for the variables Calanidae, Acartiidae, Hippolytidae, Sphaeronectidae and Palaemonidae (variables are sorted from the strongest).
- low values for the variables Metridinidae, Euphausiidae, Aetideidae, Heterorhabdidae, Coenobitidae, Pandalidae, Callianassidae and Typhloscolecidae (variables are sorted from the weakest).

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5. Description of the plane 5:6

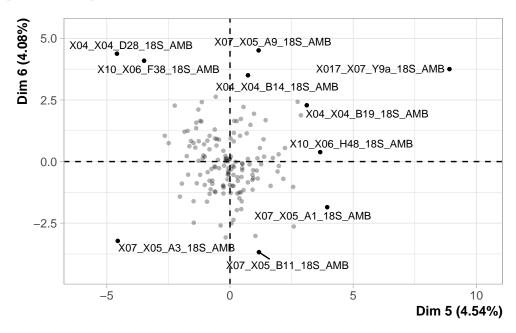


Figure 5.1 - Individuals factor map (PCA) The labeled individuals are those with the higher contribution to the plane construction.

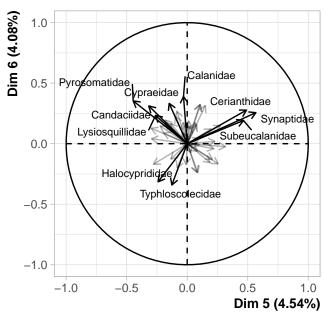


Figure 5.2 - Variables factor map (PCA) The labeled variables are those the best shown on the plane.

The dimension 5 opposes individuals such as $X017_X07_Y9a_18S_AMB$, $X04_X04_B19_18S_AMB$, $X04_X04_B14_18S_AMB$, $X07_X05_A9_18S_AMB$ and $X10_X06_H48_18S_AMB$ (to the right of the graph, characterized by a strongly positive coordinate on the axis) to individuals such as $X04_X04_D28_18S_AMB$ and $X10_X06_F38_18S_AMB$ (to the left of the graph, characterized by a strongly negative coordinate on the axis).

The group in which the individuals $X017_X07_Y9a_18S_AMB$, $X04_X04_B19_18S_AMB$,

X04_X04_B14_18S_AMB, X07_X05_A9_18S_AMB and X10_X06_H48_18S_AMB stand (characterized by a positive coordinate on the axis) is sharing:

• high values for variables like Synaptidae, Sphenopidae, Corycaeidae, Pontellidae, Hippolytidae, Cerianthidae, Hormathiidae, Scolecitrichidae, Subeucalanidae and Squillidae (variables are sorted from the strongest).

The group in which the individuals $X04_X04_D28_18S_AMB$ and $X10_X06_F38_18S_AMB$ stand (characterized by a negative coordinate on the axis) is sharing:

• high values for the variables Pyrosomatidae, Candaciidae, Metridinidae, Aetideidae, Acartiidae, Lysiosquillidae, Aequoreidae, Heterorhabdidae and Callianassidae (variables are sorted from the strongest).

The dimension 6 opposes individuals such as X017_X07_Y9a_18S_AMB, X04_X04_D28_18S_AMB, X04_X04_B19_18S_AMB, X10_X06_F38_18S_AMB, X04_X04_B14_18S_AMB, X07_X05_A9_18S_AMB and X10_X06_H48_18S_AMB (to the top of the graph, characterized by a strongly positive coordinate on the axis) to individuals such as X07_X05_A3_18S_AMB, X07_X05_A1_18S_AMB and X07_X05_B11_18S_AMB (to the bottom of the graph, characterized by a strongly negative coordinate on the axis).

The group in which the individuals $X04_X04_D28_18S_AMB$ and $X10_X06_F38_18S_AMB$ stand (characterized by a positive coordinate on the axis) is sharing:

• high values for the variables Pyrosomatidae, Candaciidae, Metridinidae, Aetideidae, Acartiidae, Lysiosquillidae, Aequoreidae, Heterorhabdidae and Callianassidae (variables are sorted from the strongest).

The group in which the individuals $X017_X07_Y9a_18S_AMB$, $X04_X04_B19_18S_AMB$, $X04_X04_B14_18S_AMB$, $X07_X05_A9_18S_AMB$ and $X10_X06_H48_18S_AMB$ stand (characterized by a positive coordinate on the axis) is sharing :

• high values for variables like Synaptidae, Sphenopidae, Corycaeidae, Pontellidae, Hippolytidae, Cerianthidae, Hormathiidae, Scolecitrichidae, Subeucalanidae and Squillidae (variables are sorted from the strongest).

The group in which the individuals $X07_X05_A3_18S_AMB$, $X07_X05_A1_18S_AMB$ and $X07_X05_B11_18S_AMB$ stand (characterized by a negative coordinate on the axis) is sharing :

- high values for the variables *Typhloscolecidae* and *Halocyprididae* (variables are sorted from the strongest).
- low values for variables like *Pyrosomatidae*, *Calanidae*, *Candaciidae*, *Scolecitrichidae*, *Temoridae*, *Callianassidae*, *Sphenopidae*, *Sapphirinidae*, *Processidae* and *Coenobitidae* (variables are sorted from the weakest).

8

6. Description of the plane 7:8

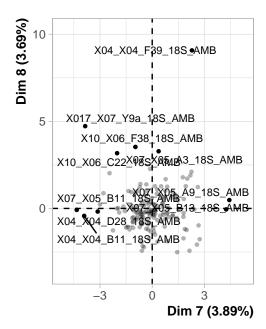


Figure 6.1 - Individuals factor map (PCA) The labeled individuals are those with the higher contribution to the plane construction.

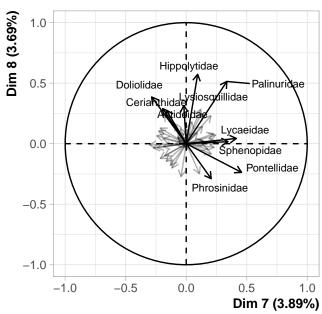


Figure 6.2 - Variables factor map (PCA) The labeled variables are those the best shown on the plane.

The dimension 7 opposes individuals such as $X07_X05_B13_18S_AMB$ and $X07_X05_A9_18S_AMB$ (to the right of the graph, characterized by a strongly positive coordinate on the axis) to individuals such as $X10_X06_C22_18S_AMB$, $X017_X07_Y9a_18S_AMB$, $X04_X04_D28_18S_AMB$, $X07_X05_B11_18S_AMB$, $X10_X06_F38_18S_AMB$, $X04_X04_B11_18S_AMB$ and $X07_X05_A3_18S_AMB$ (to the left of the graph, characterized by a strongly negative coordinate on the axis).

The group 1 (characterized by a positive coordinate on the axis) is sharing:

- high values for the variables *Phrosinidae*, *Pontellidae* and *Eupronoidae* (variables are sorted from the strongest).
- low values for the variables Aetideidae, Doliolidae, Sapphirinidae and Sphaeronectidae (variables are sorted from the weakest).

The group in which the individuals $X07_X05_B13_18S_AMB$ and $X07_X05_A9_18S_AMB$ stand (characterized by a positive coordinate on the axis) is sharing:

- high values for the variables Lycaeidae, Palinuridae, Palaemonidae and Sphenopidae (variables are sorted from the strongest).
- low values for the variables *Heterorhabdidae*, *Metridinidae* and *Geryoniidae* (variables are sorted from the weakest).

The group in which the individuals $X04_X04_D28_18S_AMB$, $X07_X05_B11_18S_AMB$ and $X04_X04_B11_18S_AMB$ stand (characterized by a negative coordinate on the axis) is sharing:

- high values for the variables Callianassidae, Aetideidae, Scombridae, Oikopleuridae, Centropagidae and Pyrosomatidae (variables are sorted from the strongest).
- low values for the variables *Pontellidae*, *Lycaeidae*, *Euphausiidae* and *Processidae* (variables are sorted from the weakest).

The group in which the individuals $X10_X06_C22_18S_AMB$, $X017_X07_Y9a_18S_AMB$, $X10_X06_F38_18S_AMB$ and $X07_X05_A3_18S_AMB$ stand (characterized by a negative coordinate on the axis) is sharing :

• high values for the variables *Doliolidae*, *Lysiosquillidae*, *Cerianthidae*, *Typhloscolecidae*, *Sapphirinidae*, *Metridinidae*, *Halocyprididae* and *Cypraeidae* (variables are sorted from the strongest).

The **dimension 8** opposes individuals such as $X04_X04_F39_18S_AMB$, $X07_X05_B13_18S_AMB$, $X10_X06_C22_18S_AMB$, $X017_X07_Y9a_18S_AMB$, $X07_X05_A9_18S_AMB$, $X10_X06_F38_18S_AMB$ and $X07_X05_A3_18S_AMB$ (to the top of the graph, characterized by a strongly positive coordinate on the axis) to individuals characterized by a strongly negative coordinate on the axis (to the bottom of the graph).

The group in which the individuals $X10_X06_C22_18S_AMB$, $X017_X07_Y9a_18S_AMB$, $X10_X06_F38_18S_AMB$ and $X07_X05_A3_18S_AMB$ stand (characterized by a positive coordinate on the axis) is sharing :

• high values for the variables *Doliolidae*, *Lysiosquillidae*, *Cerianthidae*, *Typhloscolecidae*, *Sapphirinidae*, *Metridinidae*, *Halocyprididae* and *Cypraeidae* (variables are sorted from the strongest).

The group in which the individuals $X07_X05_B13_18S_AMB$ and $X07_X05_A9_18S_AMB$ stand (characterized by a positive coordinate on the axis) is sharing:

- high values for the variables *Lycaeidae*, *Palinuridae*, *Palaemonidae* and *Sphenopidae* (variables are sorted from the strongest).
- low values for the variables *Heterorhabdidae*, *Metridinidae* and *Geryoniidae* (variables are sorted from the weakest).

The group in which the individual $X04_X04_F39_18S_AMB$ stands (characterized by a positive coordinate on the axis) is sharing:

• high values for the variables *Hippolytidae*, *Palinuridae* and *Clausocalanidae* (variables are sorted from the strongest).

The group 4 (characterized by a negative coordinate on the axis) is sharing:

• high values for the variables *Phrosinidae*, *Pontellidae* and *Eupronoidae* (variables are sorted from the strongest).

• low values for the variables Aetideidae, Doliolidae, Sapphirinidae and Sphaeronectidae (variables are sorted from the weakest).

7. Description of the plane 9:10

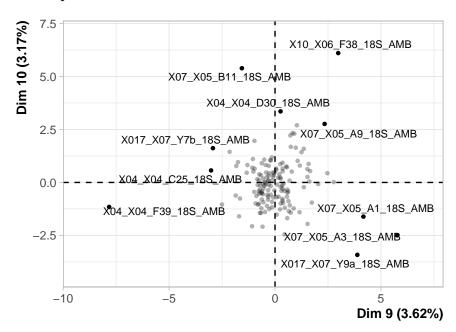


Figure 7.1 - Individuals factor map (PCA) The labeled individuals are those with the higher contribution to the plane construction.

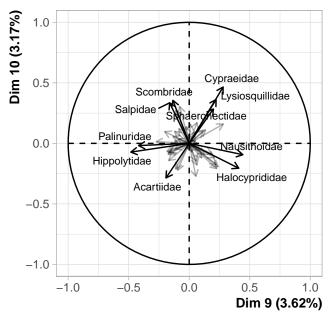


Figure 7.2 - Variables factor map (PCA) The labeled variables are those the best shown on the plane.

The dimension 9 opposes individuals such as $X07_X05_A3_18S_AMB$, $X017_X07_Y9a_18S_AMB$, $X07_X05_A1_18S_AMB$ and $X07_X05_A9_18S_AMB$ (to the right of the graph, characterized

by a strongly positive coordinate on the axis) to individuals such as $X04_X04_F39_18S_AMB$, $X017_X07_Y7b_18S_AMB$ and $X04_X04_C25_18S_AMB$ (to the left of the graph, characterized by a strongly negative coordinate on the axis).

The group in which the individual $X07_X05_A9_18S_AMB$ stands (characterized by a positive coordinate on the axis) is sharing :

- high values for the variables *Sphaeronectidae*, *Lysiosquillidae* and *Sphenopidae* (variables are sorted from the strongest).
- low values for the variables *Metridinidae*, *Aetideidae* and *Acartiidae* (variables are sorted from the weakest).

The group in which the individuals $X07_X05_A3_18S_AMB$, $X017_X07_Y9a_18S_AMB$ and $X07_X05_A1_18S_AMB$ stand (characterized by a positive coordinate on the axis) is sharing:

• high values for the variables Typhloscolecidae, Centropagidae, Cerianthidae, Subeucalanidae, Synaptidae, Doliolidae, Sapphirinidae, Nausithoidae, Hormathiidae and Creseidae (variables are sorted from the strongest).

The group in which the individual $X04_X04_F39_18S_AMB$ stands (characterized by a negative coordinate on the axis) is sharing:

• high values for the variables *Hippolytidae*, *Palinuridae* and *Clausocalanidae* (variables are sorted from the strongest).

The group in which the individuals $X017_X07_Y7b_18S_AMB$ and $X04_X04_C25_18S_AMB$ stand (characterized by a negative coordinate on the axis) is sharing:

• high values for the variables Mysidae, Squillidae, Coenobitidae, Palaemonidae, Oikopleuridae, Paracalanidae and Callianassidae (variables are sorted from the strongest).

The dimension 10 opposes individuals such as $X10_X06_F38_18S_AMB$, $X04_X04_D30_18S_AMB$, $X07_X05_B11_18S_AMB$ and $X07_X05_A9_18S_AMB$ (to the top of the graph, characterized by a strongly positive coordinate on the axis) to individuals such as $X07_X05_A3_18S_AMB$, $X017_X07_Y9a_18S_AMB$ and $X07_X05_A1_18S_AMB$ (to the bottom of the graph, characterized by a strongly negative coordinate on the axis).

The group in which the individuals $X10_X06_F38_18S_AMB$, $X04_X04_D30_18S_AMB$ and $X07_X05_B11_18S_AMB$ stand (characterized by a positive coordinate on the axis) is sharing :

• high values for the variables Pandalidae, Scombridae, Salpidae, Cypraeidae, Lysiosquillidae and Pyrosomatidae (variables are sorted from the strongest).

The group in which the individual $X07_X05_A9_18S_AMB$ stands (characterized by a positive coordinate on the axis) is sharing:

- high values for the variables *Sphaeronectidae*, *Lysiosquillidae* and *Sphenopidae* (variables are sorted from the strongest).
- low values for the variables *Metridinidae*, *Aetideidae* and *Acartiidae* (variables are sorted from the weakest).

The group 3 (characterized by a negative coordinate on the axis) is sharing:

- high values for the variables *Metridinidae*, *Acartiidae*, *Scolecitrichidae* and *Euphausiidae* (variables are sorted from the strongest).
- low values for the variables *Doliolidae*, *Sagittidae*, *Lysiosquillidae* and *Sphaeronectidae* (variables are sorted from the weakest).

The group in which the individuals $X07_X05_A3_18S_AMB$, $X017_X07_Y9a_18S_AMB$ and $X07_X05_A1_18S_AMB$ stand (characterized by a negative coordinate on the axis) is sharing:

• high values for the variables Typhloscolecidae, Centropagidae, Cerianthidae, Subeucalanidae, Synaptidae, Doliolidae, Sapphirinidae, Nausithoidae, Hormathiidae and Creseidae (variables are sorted from the strongest).

8. Description of the dimension 11

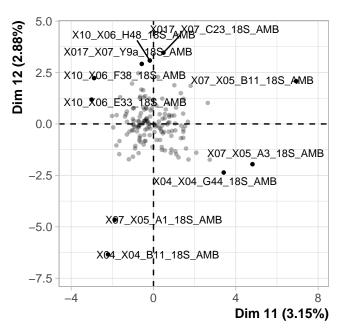


Figure 8.1 - Individuals factor map (PCA) The labeled individuals are those with the higher contribution to the plane construction.

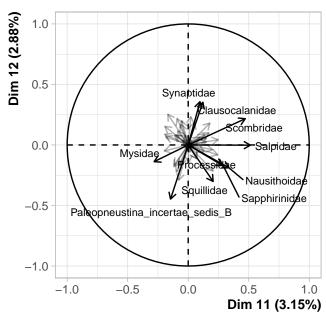


Figure 8.2 - Variables factor map (PCA) The labeled variables are those the best shown on the plane.

The dimension 11 opposes individuals such as X07_X05_B11_18S_AMB, X07_X05_A3_18S_AMB and X04_X04_G44_18S_AMB (to the right of the graph, characterized by a strongly positive coordinate on the axis) to individuals such as X04_X04_B11_18S_AMB, X07_X05_A1_18S_AMB and X10_X06_E33_18S_AMB (to the left of the graph, characterized by a strongly negative coordinate on the axis).

The group in which the individuals $X07_X05_B11_18S_AMB$, $X07_X05_A3_18S_AMB$ and $X04_X04_G44_18S_AMB$ stand (characterized by a positive coordinate on the axis) is sharing:

- high values for the variables Sapphirinidae, Squillidae, Palaemonidae, Salpidae, Processidae, Nausithoidae, Scombridae and Typhloscolecidae (variables are sorted from the strongest).
- low values for the variable Oikopleuridae.

The group in which the individuals $X04_X04_B11_18S_AMB$, $X07_X05_A1_18S_AMB$ and $X10_X06_E33_18S_AMB$ stand (characterized by a negative coordinate on the axis) is sharing:

- high values for the variables *Geryoniidae*, *Mysidae*, *Doliolidae* and *Hormathiidae* (variables are sorted from the strongest).
- low values for the variables Clausocalanidae, Corycaeidae, Heterorhabdidae, Calanidae, Metridinidae, Sapphirinidae, Pontellidae, Synaptidae and Scolecitrichidae (variables are sorted from the weakest).

9. Classification

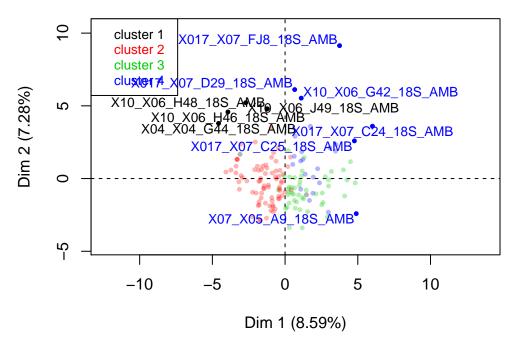


Figure 9 - Ascending Hierarchical Classification of the individuals. The classification made on individuals reveals 4 clusters.

The **cluster 1** is made of individuals such as $X04_X04_G44_18S_AMB$, $X10_X06_H46_18S_AMB$, $X10_X06_H48_18S_AMB$ and $X10_X06_J49_18S_AMB$. This group is characterized by :

• variables whose values do not differ significantly from the mean.

The **cluster 2** is made of individuals sharing:

• variables whose values do not differ significantly from the mean.

The **cluster 3** is made of individuals sharing:

• variables whose values do not differ significantly from the mean.

The cluster 4 is made of individuals such as $X017_X07_C24_18S_AMB$, $X017_X07_C25_18S_AMB$, $X017_X07_D29_18S_AMB$, $X017_X07_FJ8_18S_AMB$, $X07_X05_A9_18S_AMB$ and $X10_X06_G42_18S_AMB$. This group is characterized by :

•	variables whose	values do not d	liffer significantly from th	e mean.

Annexes