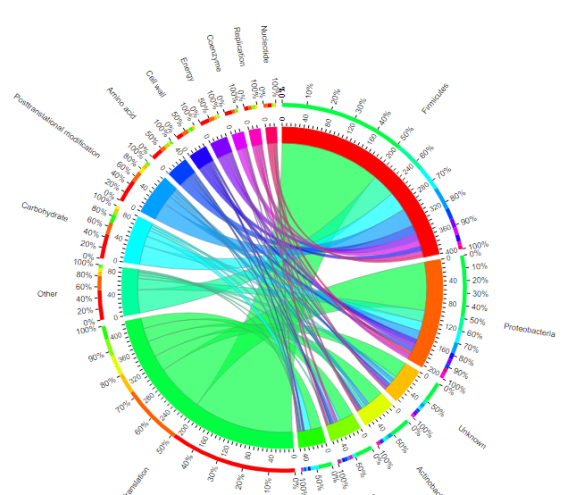# Metaproteomics Data Processing

To process your data from the MPA click on Tab 3: Metaproteomics Data Processing.

Upload the Metaproteins output file from the MPA to Chordomics via the sidebar panel on the left.

Once a dataset has been uploaded, pressing the Prepare Data button begins the processing.

The Progress window will post updates of what is happening in the processing pipeline.

Note large datasets will take a long time to process.

Once the data is ready a Download button will appear. Use this to save the processed file.

The processed data can then be uploaded to Tab 3: Processed File Upload for visualisation.