Code Readme

Rob Goodsell

2024-02-27

Scripts

This document contains a summary of the scripts contained within the 'code' folder. As many of the files are large we recommend using the 'data.table' R package for data wrangling.

assemble_OTU_data.R - contains a short example of assembling OTU clusters data with the corresponding taxonomies and metadata, as well as some simple filtering steps to remove extraction controls and biological spike ins.