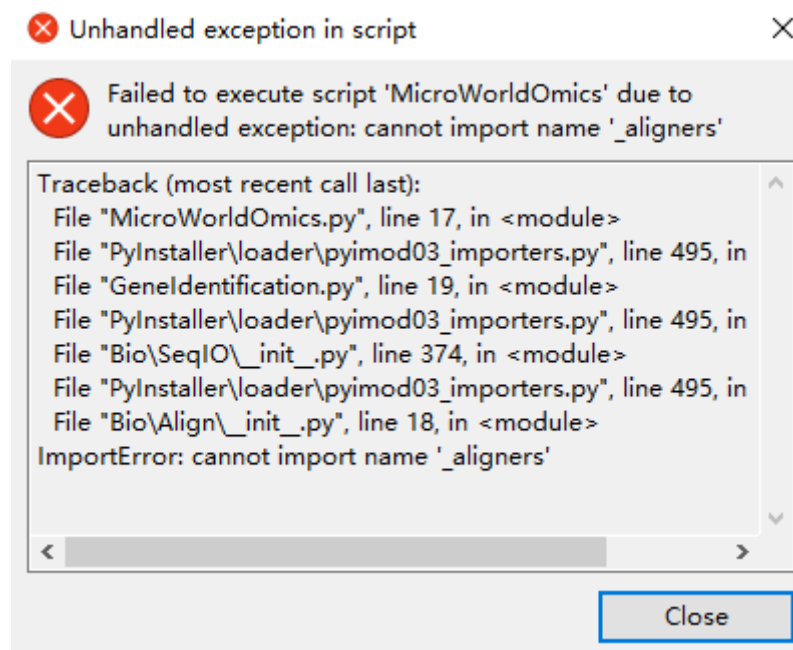


MicroWorldOmics error

(Windows)

Install error:

If similar errors occur during startup after installation, it is necessary to ensure that the installation path is consistent with the path of the module package



Module error:

1. Spieceasi Error in if (max < min) stop ("Did you flip min and max?"): argument is of length zero

Response: Would it be possible to share your data so I can recreate the issue? I'd like to have a more informative error here and better paths to resolution. Anecdotally, I've

seen that too-sparse data can lead to instabilities in the covariance matrix (i.e. - likely columns with constant values have 0 variance, which results in NAs) - but this is after the clr transformation is applied by spiec.easi, so that's difficult for users to catch on their own.

2. If "No such file" appears

Response: Upload after re-determining the path

3. In ShinyBactDating, if appears

Molecular dating results

Error: 'data' must be 2-dimensional (e.g. data frame or matrix)

Response: Please ignore and upload the data directly

4. In ShinyTimeSeries, if appears

Time prediction table

Error: 'data' must be 2-dimensional (e.g. data frame or matrix)

Response: Please ignore and upload the data directly

5. In ShinyTimeSeries, if appears

BAE table

Error: 'data' must be 2-dimensional (e.g. data frame or matrix)

Response: Please ignore and upload the data directly