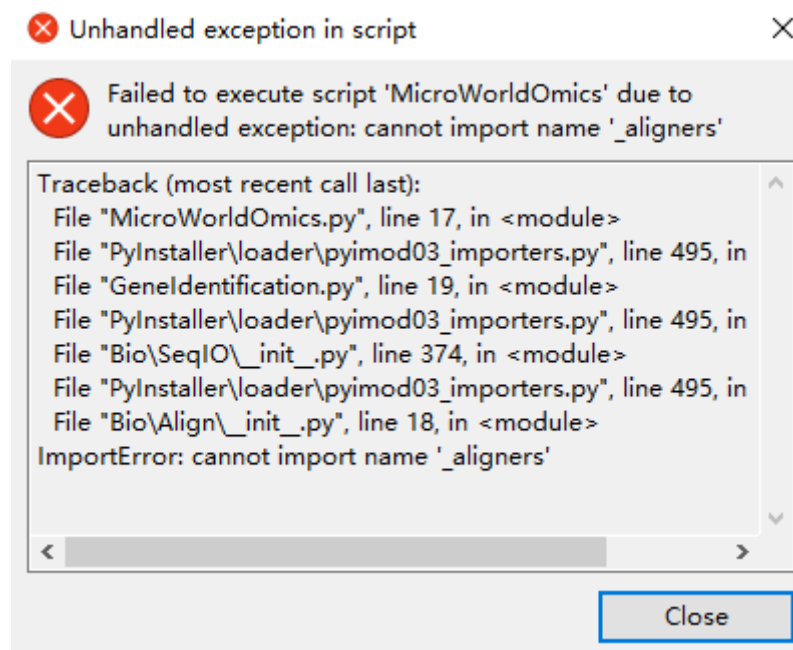# MicroWorldOmics error

(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

6. Windows disable the firewall:

全部　　应用　　文档　　网页　　更多 ▾

最佳匹配

控制面板
系统

搜索网页

Open 'Control Panel', and then click 'System and Safety':

系统和安全
查看你的计算机状态
通过文件历史记录保存你的文件备份副本
备份和还原(Windows 7)

Click 'Defender wall'

Windows Defender 防火墙
检查防火墙状态 │ 允许应用通过 Windows 防火墙

Open the new window under the red line:

Keep the panel as shown below: