1. Locate your “init.m” file.
   1. Open Mathematica, run the command “$UserBaseDirectory”. To run a command in Mathematica, press shift-enter, or use the enter key on the numeric keypad.
   2. Usually, the $UserBaseDirectory is C:\Users\YOUR-USERNAME\AppData\Roaming\Mathematica\
   3. Go to the directory indicated by $UserBaseDirectory. Enter the “kernel” and open “init.m” with text editor.
2. Modify “init.m”
   1. Make sure you defined “DropBox”

DropBox = "C:\\Users\\YOUR-USERNAME \\Dropbox";

* 1. Add this line at the bottom:

$Path = Append[$Path, DropBox<>"\\MathematicaConfig\\YG"]

* 1. Save the file and restart Mathematica.

1. Load the package when you need it:
   1. Use this command in your notebook:

Get["ProteinStructure.wl"]