

PLEASE NOTE:

For most science cases, I assume the main flowchart to be needed. This section of the code is mainly designed for high resolution data or RV fitting to a single spectral line. For single line fitting, a bootstrap algorithm instead of MCMC fitting is used to can optimally normalise data, whereas I have not done this for the MCMC for various reasons (mainly for accuracy & processing time).

Speak to me directly (email, conferences) if you have specific questions or want to know more.

