

13 August 13th

13.1 Goals for the day

Goals from Last Time

1. Test \LaTeX installation
2. Continue Readings
3. Continue analysing Gauley errors?

Additional Goals

4. Look at the code
5. Look into potential Tremont errors?
6. Talk to Cedric about errors

13.2 Progress/Notes

13.2.1 Test \LaTeX installation

\LaTeX installation works?

Compiles, works just fine. Was a slight adjustment from the windows MikTeX GUI, but largely the same. Developed the format used here.

13.2.2 Readings

13.2.3 Gauley Errors

Cedric says that the Gauley errors AREN'T ERRORS. It's actually just a regular part of the code output. Also the “#warning:iterations terminated because half-step sizes are very small” isn't a problem. Copasetic.

13.2.4 Code Examination

It seems like I should examing roc.appr.r first. any *appr* is for approximating the codon bias without knowing the expression levels, which is my goal.

What does roc.appr.r do?

1. Get initial ϕ value from data/ex.test.rda
2. Change renew.iter variable from 100 to 3 in .CF.AC, which is in data/control.r
3. Calls cubappr to do the actual approximations. Times this process. cubappr is exported to mycubappr in mycubappr.r

- (a) Setup model and check data, including data structures and storage
- (b) Initialize ϕ using `my.pInit` in `my.init_param.r`. Uses `EMCluster` iff `estimate.bias.Phi` (false by default)
- (c) Initialize β , a vector of amino acids
- (d) Runs MCMC, starting at line 124
 - i. `my.drawBConditionalAll` Updates β . Seems quite important. uses VGAM?
Further study
 - ii. Other parameters: `nu.Phi` and `sigma.Phi`
 - iii.

13.2.5 Tremont Errors?

Likely because I was using the `cubfits-0.1.0`, the version on CRAN, while I should be using `cubfits-0.1.1`, the version in `cubfits-master.tar.gz` Moving to Gauley?

13.3 Future Goals

1. Continue analyzing `cubfits-master/R` files, especially `mycubappr.r`
 - (a) Especially study the MCMC from `mycubappr.r` 124-196
 - (b) `nu.Phi` and `sigma.Phi`?
2. Read REU Paper on Nonsense Model