Data_analysis

September 17, 2019

1 Introduction

In this notebook I will apply the data from the experiments to the theory to try and predict the change in temperature sensitivity of respiration as interaction change. To do this I will need to consider:

- 1. how to parameterise the basic growth model (i.e. dynamics in the absence of interactions)
- 2. how to parameterise the interactions between species
- 3. What we can predict exactly at the community level

2 Growth model data

So for each species we have: * r - Growth Rate * R - Respiration Rate

as well as their temperature dependence's (roughly). In this section I derive these parameters from the experimental data.

2.1 Growth - *r*

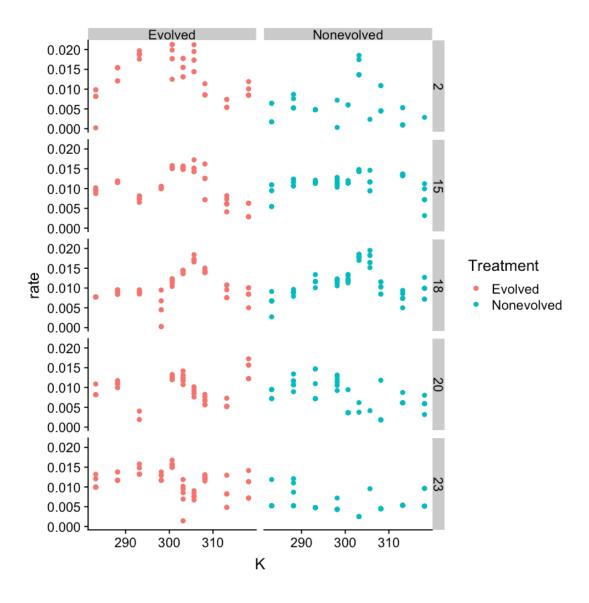
In order to parameterise the growth of species and its temperature dependence we need an r_0 value and a E_r value.

Though the temperature dependence of growth was not directly measured we can estimate E_r by using the optical density (OD) from the respiration experiments. Here the OD is known at two timepoints for each species/treatment combination across a range of temperatures. Assuming we are in exponential growth we can estimate the growth rate over this period using the formula:

$$r = \frac{\log\left(\frac{C(t)}{C(0)}\right)}{t}$$

First we read in the data and calulate the growth rates:

```
In [149]: OTU_OD <- read_csv("../data/Francisca/respirationandODData/DatosOD_5OTUs_evNev.csv",</pre>
                   col_types = cols(
                   Replicate = col_double(),
                   OTU = col_double(),
                   OD = col_double(),
                   T = col_double(),
                   t = col_character(),
                   OD_TO = col_double(),
                   Tf = col_double(),
                   u = col_double(),
                   Treatment = col_character()
                 ))
#get growth (need to correct for the negative growth rates)
OTU_growth <- OTU_OD %>%
    mutate(r = (log(OD) - log(OD_TO))/Tf , K = T + 273.15 ) %>%
    filter(r > quantile(r,probs = c(0.1))[1]) %>%
    mutate(rate = r - min(r)) \%>\%
    filter(rate > 0)
OTU_growth %>%
ggplot(aes(x = K, y = rate, colour = Treatment))+
geom_point()+
facet_grid(OTU~Treatment)
```



as we can see the TPCs are not great. However as we just need a general idea as to the temperature sensitivty we can use the approach used by Dell et al. and just fit a boltzmann to the rising portion. Though a bit rough we can do this by taking only the points that are at temperatures lower than the maximum growth rate, and taking the slope of a linear fit between $log(r) \sim \frac{1}{kT}$.