Hi Nosa,

A quick recap of some of the expectations:

1. Convert the N, S, P, K, Mg, Ca, and Na to mg/kg by multiplying by 100 before the analysis.

2. Use the nutrient yield values by multiplying the concentration of each element by the tissue biomass yield

3. Plot two graphs: one showing the relative change to the control and the other with the error bars (I think we agreed on using the standard errors for this one).

4. Kindly skim through the r script (seldon\_processing...). It might help save some time and give you a broader overview of similar analyses. The 'rough\_nutrient\_script' is what I used to generate the relative change plots I shared with you.

Kindly let me know if you have any questions.

Best regards,

Eneji