

Date: 05 Aug 2022
To: "Brian Joseph Bohman" bohm0072@umn.edu
From: "Pirjo Peltonen-Sainio" pirjo.peltonen-sainio@luke.fi
Subject: EURAGR11860

Ref.: Ms. No. EURAGR11860
 Quantifying critical N dilution curves across G × E × M effects for potato using a partially-pooled Bayesian hierarchical method
 European Journal of Agronomy

Dear Dr. Bohman,

I can now inform you that the Editorial Board has evaluated the manuscript EURAGR11860: Quantifying critical N dilution curves across G × E × M effects for potato using a partially-pooled Bayesian hierarchical method.

The Editor has advised that the manuscript will be reconsidered for publication after major revision.

Please submit your revision at latest by
 03 Nov 2022

The comments below should be taken into account when revising the manuscript. Along with your revised manuscript, you will need to supply a covering letter in which you list all the changes you have made to the manuscript, and in which you detail your responses to all the comments passed by the reviewer(s) and the Editor. Should you disagree with any comment(s), please explain why.

To submit a revision, please visit <https://www.editorialmanager.com/euragr/> and log in as an Author. You will see a menu item called "Submission Needing Revision". The revised manuscript and covering letter can be submitted there.

To speed up the production process, I would like to ask you to upload all source files separately. Figures should be uploaded per number. For example, figures 1, 2a+b and 3 should be uploaded as 3 separate files.

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Kind regards,

Pirjo Peltonen-Sainio, Ph.D.
 Editor In Chief
 European Journal of Agronomy

Reviewers' comments:

Reviewer #1: This paper presents and discuss a new method for estimating Critical Nitrogen Dilution Curve for potatoes. Authors have pooled a large data set from different countries, Argentina, Canada and Belgium with different cultivars, and by using a Bayesian statistical approach they analysed the uncertainty of CNDC parameter determination and tried to infer on variation across Genotype-Environment-Management conditions. The manuscript is very well written and very well organized. As said by authors the use of Bayesian method for CNDC uncertainty analysis is not fully original, but they used this method very accurately and they proposed a new approach of partial pooling for a better analysis of G-E-M effects. So this manuscript is a very original and relevant contribution to crop N diagnosis problem. As I am not an expert in statistics, I cannot provide any comments on this part of the work. I guess that reviewing this manuscript by an expert in statistics and more particularly in Bayesian approach would be important. So our reviewing focus on agronomic and physiological interpretation.

The problem of "N dilution" process in crop having a strong "reserve" compartment as grain or tuber has been perfectly identified and discussed in introduction by authors. The "N dilution" model, originally developed on forage crop... only concern with plants in vegetative growth: producing only leaves and stems.... It was the reason why CNDC were limited in theory to flowering stage or just to early reproductive development when "grain biomass" was not too high... For taking into account grain or tuber growth... it should be necessary to have a two step N dilution (i) during vegetative growth ... with a given allometry coefficient "b1" reflecting the biomass allocation to "metabolic" and "structural" compartment...; and (ii) during grain or tuber filling reflecting C-N remobilisation from vegetative part and filling grain and tuber...with a more or less different value of allometry "b2" depending on the C-N ratio of grain or tuber accumulated. So "b2" being >> "b1" for grain and tuber accumulating preferentially starch (as for potatoes) or more or less = 0 < b1 for grain accumulating preferentially proteins such as grain legumes... So it is clear that representing a single CNDC with a constant allometry "b"... while there is in fact a "break" in dilution process from "b1" to "b2"... is a problem... This problem is not so important statistically if b1 and b2 are not very different... but becomes very important when they are very different as for potatoes. So the best way for showing this "break" in CNDC and to determine when this break occurs would be to represent CNDC in Log-Log term... So I suggest authors to illustrate that and to determine at which extent variation in the "break time" on the "crop biomass axis... would depend on G-E-M ? I think Giletto et al. (see their paper in EJA) have already well analysed this problem in Argentina? So it should be easy to this group of authors to deep this question. They can also refers to analogous work made on maize and wheat (ZHAO, B., ATA UL KARIM, S., T., LEMAIRE G., DUAN, A., LIU, Z., GUO, Y., QIN, A., NING, D., LIU, Z., 2021. Exploring the source-sink relationship to quantify ear nitrogen accumulation in summer maize and winter wheat using critical nitrogen dilution curve. Field Crop Research, 274. <https://doi.org/10.1016/j.fcr.2021.108332>). By this way their comparison with "other crops" should be more complete because in their manuscript they compare CNDC of potatoes including tuber filling process with other crops where CNDC was limited to "vegetative period"!!! So for potatoes, if G-E-M interaction has an impact on the onset of the change from b1 to b2 (as the start of tuber development)...as a consequence it should have an impact on the "average" CNDC fitted with a constant "b"!!! So it should be important to verify this hypothesis: has G-E-M an effect of both b1 or b2 separately...? or has G-E-M has only an effect of the onset of change from b1 to b2 ? That would be a more fundamental question?

So my conclusion is that this excellent manuscript should be accepted for publication... But I suggest authors to improve its scientific value by adding some informations on "b1" and "b2" for being able to discuss more strongly the hypothesis above.

Details:

Line 67: No, NNI is very sensible to any fertilization management...as it detect any effect on plant N nutrition status....

Line 76: No, b is the ration between relative rate of %N decline (d(%N)/(%N)dt and the relative rate of biomass accumulation dW/Wdt.... that is different of the rate of %N decline (d(%N)/dt

Line 92: add "Acceleration" of dilution....

Line 189: "reduce" is repeated two time....

Reviewer #2: This study advances on the use of Bayesian hierarchical frameworks to develop critical N dilution curves introducing a partial pooling approach through random components. This could represent a useful alternative for comparing CNDs across $G \times E \times M$ conditions. Moreover, it could be further extended for developing critical N dilution curves of potato but also potentially of other crops. I found it interesting to read and review, which makes me think it would be very relevant for EJA journal. I also found the paper excessively (and unnecessary) long in several sections, so my first main suggestion is to reduce the length of the manuscript reorganizing paragraphs and ideas. I identified below several sections and paragraphs where this could be done. Similarly, I would suggest reducing the number of figures considering the complexity of the methodology and number of panels. I believe the paper will have more impact if ideas (including figures) are more succinct. My second main suggestion is on one of the methodologies used and (at some point) recommended to evaluate uncertainty in the CND. This should be addressed before publication. See below specific comments related to this in LI423 & LI524.

Specific comments:

LI28: "was attributed to variation" is a statement that cannot be confirmed with this analysis due to the lack of factorial combinations of the G (maturity classes) x M (plant density) at each site (i.e., E). It would be better to claim "was hypothesized".
 LI59-62: no need to go back to the rate-response approach as the paper is not about it, could be removed.
 LI82-114: I think this section can be largely summarized in a single paragraph. This paper is more about the methodology of fitting and quantifying uncertainty in CND, so only a brief overview of the dilution theory + use in potato is needed.
 LI117-118: this is a good example of the type of "expensive" writing used along the paper that makes it hard to read... "Previous development of CNDs for potato has been conducted using a non-uniform set of statistical methods and with limited quantification of uncertainty in either the range of plausible %Nc values or the fitted parameter values themselves" can be rephrased with the same meaning by "Previous CNDs for potato have been developed with different statistical methods and limited quantification of their uncertainty." Simplifying sentences would not only reduce the length of the paper but also increase the impact of each message. I would recommend considering this point when re-organizing ideas.
 LI126-127 & 133-134: These three paragraphs can be combined into one, no need for break lines
 LI141: "linear plateau was designed to discriminate against" Not clear. In the paragraph above, it was claimed that the linear plateau cannot address these exact two points?
 LI166-192: Should be combined into a single paragraph.
 Tables 2, 3 and 4 can be combined into one. Table 1 can be combined within Table 2-3-4 and Table 5. Will then need only two tables.
 LI273: Is confusing referring here as "experimental" data, considering the first set of sites were classified as "Experimental" vs "Prev. published". Can consider the use of terms here. Section 2.1.2. I suggest using the Tables to report detailed information of these experiments, considering detailed information has been already reported in previous publications. Can use text to report only data that is not in Table, such as location site.
 LI336-347: Is this paragraph needed? Most of this has been mentioned in the introduction. Same with figure 1, I think it is not a critical figure of the manuscript and could be removed?
 LI391: What about replicates? How they were treated in the model?
 LI395: Convergence checks? I assume they were conducted, please add.
 LI397: "biologically or physically impossible predictions" not sure what that means.
 LI401: What about priors for the random effects variances? it looks to me that the priors from Table 6 are very informative and could constrain parameters to a short range of variation, but according to the results, there seems less restriction on priors for the random effects? This can be tested through a sensitivity analysis with less informative prior distributions.
 LI405: This is great addition and authors should be congratulated for this.
 LI423: I am not sure I understood this, but this might not be correct. Did you fit a new model (and frequentist?) to the data estimated by the parameters of the 90% credible limits of the Bayesian curves? There is circularity in this approach, and not sure why it was done.
 LI436: The definition of delta%Nc is not clear. This is a very complex and long sentence, but critical to understand the paper. What is "the difference between the 0.50 quantile for %Nc and the various methods to quantify uncertainty (i.e., 90% credible region for %Nc, CND_{Cup} & CND_{Clo}, and estimates of credible region for %Nc using 90% credible interval for parameters a and b)?"
 LI448-455: Another case of "expensive" writing.
 LI479: Figure 1 again? Check numbering of all figures.
 LI525: Figure 4: why 15 individual draws are represented in red? Please clarify this analysis.
 LI542: Why is quite uninformative? I do not agree with this analysis of comparing methods to measure uncertainty of the CND because there are basically different things being compared. If there is interest in quantifying uncertainty of the %Nc, credible intervals for the %Nc should be analysed. If there is interest in quantifying uncertainty in the parameters of the CND, credible intervals for a and b parameters. This is the advantage of the Bayesian framework through the obtained posterior distributions.
 LI612: If there were significant differences, wasn't it also biased? Can claim that was less biased.

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MethodsX (optional)

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