

Interactive comment on “Variability in copepod trophic levels and in feeding selectivity based on stable isotope analysis in Gwangyang Bay off the southern coast of Korea” by Mianrun Chen et al.

Anonymous Referee #3

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Review of “Variability in copepod trophic levels and in feeding selectivity based on stable isotope analysis in Gwangyang Bay off the southern coast of Korea”. Authors: Mianrun Chen, Dongyoung Kim, Hongbin Liu, and Chang-Keun Kang. Submitted for review for journal Biogeosciences.

The authors use an approach that combines Generalized additive models (GAMs) and multiple regressions using bulk carbon and nitrogen isotopes to address trophic relationships among zooplankton taxa and POM, along a salinity gradient in Gwangyang Bay, off the southern coast of Korea. They find significant spatial variability, somewhat coherent patterns among copepods and microplankton, and seasonal variability in both

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d13C and d15N. Using an abundance-weighted regression approach, the authors estimate bulk values for each copepod genera or taxonomic grouping, and from these infer particle selectivity, diet and trophic level (TL), for each genera with significant regressions.

General comments:

I find a problem in the way the authors estimated the weight differences between cyclopoids and calanoids randomly, as well as assuming that the weight of all calanoid genera was the same. In particular, because the authors have the taxonomic information already, I suggest they do a literature review and obtain the average weight values for each of the copepod genera/species used in the study, and apply these to the bulk regressions. I believe this is especially important as the authors are trying to extrapolate significantly more results than what they measured (i.e. genera-specific isotope values from a mixed community), that the approach be as precise as possible.

In general I appreciate the effort to expand upon simple d13C and d15N bulk measurements for more detailed information on a community. However, in the case of copepods, if the authors do/did intend to investigate these relationships in detail, why not simply measure the values of individual genera? They state that too much material is required, but methodological advances these days mean that an individual *Calanus* female can indeed be analyzed ($\sim 60 \mu\text{gC}$, $10 \mu\text{gN}$), as $5 \mu\text{gN}$ is typically the lower limit of standard bulk analyses (and low-N methods methods have been developed to go down to $\sim 1 \mu\text{gN}$). Cyclopoids would require greater number, but following the authors assumptions of $0.1 < x < 1$, that would be about 10 individuals. When certain problems arise, such as *Paracalanus* and *Sinocalanus* having lower d13C values than any measured prey, it would seem the authors acknowledge them, but then continue their analyses, e.g. calculate a TL (presumably based on prey that has been shown to not be consistent with their isotope values) in the same way as for the other genera. One gets the sense that by plugging it into GAMs and regression models, the error sources and magnitudes are lost. I would like to see a quantitative test of the biases inherent in this

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Bayesian model, and how confident the authors can be that this approach is recovering the actual copepod diets. Given this approach and the number of assumptions that lie within, uncertainty relating to the model (as well as replication, independently) should be presented, discussed, and assessed explicitly with the other sources of uncertainty. This should be done with both the particle feeders and the carnivorous species, and the effects of including or excluding different species types should also be assessed.

Finally, consistent with the point I discuss above, the authors mention a ‘simple energy flow’ in the abstract and discussion. But I wonder if this methodological approach allows for more complex flows. The actual isotopic values were not measured, but inferred from mass balance of dominant genera, and Bayesian approaches, and the violation of the underlying assumptions was not determined. How would a more complex picture emerge? In fact, the problem of Paracalanus and Sinocalanus having lower $\delta^{13}\text{C}$ values could hint at more complexity, yet it is assumed perhaps that this is due to unmeasured food sources and then ignored.

I think if the authors address the issues posed above (and specifics below), the MS is suitable for publication.

Specific comments:

Abstract. P1 – 10. The word ‘trophism’ is introduced yet does not technically mean what the authors define it as (food resources and trophic levels), and is not used within the field’s jargon as such either. I would prefer ‘trophic structure’ or ‘trophic interactions’, or ‘trophic preferences’.

‘Temperature-related’ seasonal variations – The effect of temperature from season was not separated in this study, it should be simply ‘seasonal variations’.

Introduction

P2-0. “With broad feeding spectra and flexible feeding strategies, the bulk copepod assemblage is omnivorous depending on dominant species or group”. Omnivo-

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rous or what? Consider changing to something like 'displays varying degrees of herbivory/omnivory/carnivory, depending on dominant. . .'

P2-5. "In turn, TLs of a diverse. . .". I assume the authors here refer to the average trophic position of the assemblage, and thus should be 'TL' (singular). "Because copepods play a fundamental role in feeding on phytoplankton as primary consumers". Consider re-phrasing as 'Because copepods rely significantly on phytoplankton as prey', otherwise the expectation of this phrase is that the second half will refer to the top-down effect of copepods on phytoplankton, and not the bottom-up effect of phytoplankton on copepods. 'feeding on phytoplankton as primary consumers, so the seasonal and spatial'. Delete 'so'.

P2-15. "Therefore, the assessment of the trophic position (. . .) of copepods within a complex planktonic food web is critical in predicting the ecological relationships between predator and prey". This phrase seems redundant, isn't the study about assessing these ecological relationships? I don't understand the prediction part.

P3. 0. "In contrast, the $\delta^{15}\text{N}$ values of primary producers increase from being nutrient-sufficient (high fractionation) to nutrient-limiting (low fractionation) and are especially high in anthropogenic wastewater nitrogen inputs". Would the later simple swamp the fractionation effect? The literature on $\delta^{15}\text{N}$ of different nutrients in the ocean (nitrate, ammonia, urea) shows ranges that are much larger than fractionation factors, e.g. these vary by about 20‰ compared to 3.4‰ of fractionation. Can you comment on how much you expect the source to vary along the river gradient?

Materials and methods

P4-25. Could you mention the average volume filtered per tow, as the net was equipped with a flow meter?

P5-5. "water samples were transported to the laboratory as soon as possible". Please give a time estimate.

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P6-5. The analytical precision of 0.2‰ and 0.3‰ for d13C and d15N, respectively, seems a bit high. Could you estimate what is the lowest change in TL that you can estimate based on this instrument error?

P6.15. The weight difference between cyclopoids and calanoids was generated randomly. I don't understand why the information from the species identification was not used for this purpose. What is the error associated with this type of computation? I would really suggest the authors do a literature search of the mean weights of the difference species and genera enumerated in their samples, and use this information to estimate both cyclopoid/calanoid weights, and the weights of the different calanoid genera. If the composition has already been estimated, it makes no sense to make these assumptions that only introduce greater error into an already indirect way of estimating species stable isotope composition.

P7-20. 'fractionation factors used in the model estimation were calculated from TLs'. I don't understand this statement, it sounds like 3‰ and 0.5 ‰ were assumed (logically) and not calculated. Please clarify.

Results

The authors discuss their seasonal results in the context of 'temperature'. I would prefer to see this discussed as 'seasonal', since temperature variability within a season was not tested and hence the driver of the observed effects cannot be unequivocally stated to be temperature. Rather, they are probably a combined effect of the changes that co-occur with each season and should be stated as such.

P8-10. 'Despite insignificant spatial variability, higher Chl a concentrations generally occurred in the middle of the bay'. This is not obvious from the values in the table. Please explain in more detail or remove.

P8-25. Please give a mean value for copepod d13C as done for the groups above (nanoplankton and microplankton)

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P9-0. “Overall, seasonal succession of winter-spring, spring-summer, and summer-autumn were apparent for all plankton groups”. Not clear what this means. There appears to be significant overlap in values for the nanoplankton, and no clear increasing progression from winter to autumn, as increases/decreases seem to interchange.

P9-5. It isn't clear to me how the coefficients of variation are calculated. The range of d15N values encountered is less than that for d13C, although the spatial progressions are less monotonic. Please clarify in the methods how this is calculated. P9-10. The result for the microplankton is inconsistent with the figure. In the figure, the highest value for d15N is 10‰ at the bay in spring. There is no 16.2 value.

P9-15. “Copepod d15N . . . being much more consistent with the pattern of microplankton than that of nanoplankton”. This seems true for the summer d15N values, and quite the opposite for the winter values. Regardless, there is such high variability that it is hard to tease out any clear pattern of spatial/seasonal co-variability.

P9-20. The GAM result is very interesting. Perhaps it reflects the food-web processes that affect d15N disproportionately and were not included in the GAM?

P10-20. It is not clear to me how the trophic levels of brackish copepods can be calculated, when their 13C values do not support the sampled nanoplankton and/or microplankton as their food source. I also don't understand how later in figure 6 they show up enriched, but in figure 4 they are depleted with respect to this food source. The differences between these two figures should be stated clearly as they show different results.

P10-25. “The enrichment values for nanoplankton feeding on marine and brackish water calanoids. . .”. This phrase says that nanoplankton are feeding on copepods. That's not right, it should say something like ‘enrichment values for marine and brackish water. . . feeding on nanoplankton’.

P11-5. I disagree with the statement (based on the figure) that “the proportions of the

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two size fractions of POM averaged from all four seasons contributing to copepod diets at different stations were also distinctly different except for station 8 (Fig. 8)". It seems that the error bars overlap at station 1 (hence not different), and stations 6 and 7. I might be missing something but then it should be clarified. P11-10. Does 'spring data available' mean 'only spring shown'?

The authors discuss size-selective feeding of calanoids in the context of 'filtering efficiency', yet they are not true filter feeders, they are suspension feeders that trap and handle particles (Paffenhofer et al, 1982, Mar Bio 67:2), which has different implications for particle handling. This is an important distinction that should be observed throughout the MS.

Discussion P13-0. It seems to me that the sewage explanation deserves a bit more attention. If the authors can't rule it out it means that this could contribute substantially and swamp the other subtle processes discussed in the ^{15}N -enriched ammonia section.

P13-5. "Furthermore, the fractionation effect of phytoplankton will be reduced when phytoplankton became nitrogen-limited and take up nitrogen with little fractionation". I am unsure that this effect could be significant in a coastal areas such as this one. Moreover, if phytoplankton reduce their fractionation, it would mean that their ^{15}N will tend to be higher (as they choose the lighter ^{14}N), and thus doesn't explain this decreasing trend.

P13-10. I would like to see table with the GAM results. It would be nice to have these presented first in the results, and later discussed. It would also be interesting to see the different variables tested and the ones found to be significant within this table.

P13-20. But see Gutierrez-Rodriguez et al (2014, L&O, vol:59, i5) on negligible trophic enrichment of heterotrophic protists.

P14-0. "Because of different feeding behaviors and fractionation effects of copepods,

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the variability of trophic positions of copepod assemblage depends on the overall composition of species and is determined by dominant species.” Change to “. . .the variability of the average community trophic position depends on the overall composition of species and is determined by the dominant species.”

I am somewhat confused about the discussion of trophic levels of the copepods *Paracalanus* and *Sinocalanus*. The authors state that their $\delta^{13}C$ values are lower than all measured food sources, which would imply that their food source has not been adequately measured. How then are these organisms included in the trophic level (TL) component of the paper? A bit of clarification on this topic would really help the reader.

P17-10. This paragraph explaining the Bayesian mixing model methods/results should be moved to the results section.

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