

***Interactive comment on “Observing and modelling phytoplankton community structure in the North Sea: can ERSEM-type models simulate biodiversity?” by David A. Ford et al.***

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# Comment to Ford et al. "Observing and modelling phytoplankton community structure in the North Sea: can ERSEM-type models simulate biodiversity?", Biogeoscience-discussion, 2016

Interactive  
comment

This work addresses an important aspect of marine ecosystem modelling and spends a commendable effort towards the assessment of the North Sea phytoplankton community structure.

However, with respect to the speculation on the reason of differences in community structure in the two models already discussed by the referees, I'd encourage the authors to take into consideration the insights of Sinha et al. 2010, demonstrating the sensitivity of the modelled ecosystem food-web structure to the physical driver alone. These findings are potentially in contrast to the hypothesis that the causes for differences in community structure are mainly in the parametrisation of the biogeochemical model, which therefore requires a much more solid base (e.g. simulations using identical physical driver and external forcings).

Furthermore, it should be noted that one of the model versions used in this work (referred to as PML-ERSEM) is based on a parametrisation published in 2004. Since then, other versions and parametrisations of this ERSEM flavour have emerged (e.g.

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the recent works of Butenschön et al. 2016, Ciavatta et al. 2016 and de Mora et al. 2016) that show a distinctively different community structure than the parametrisation used in this work (even if for different configurations, see figures 2 and 3 in de Mora et al. 2016 and figure 10 in Butenschön et al. 2016). While I wouldn't expect the authors to repeat their work with a different parametrisation, the fact that the used parametrisation (which is the established one still in use in the operational suite of the UKMO) has evolved since its publication clearly deserves mentioning as these new parametrisations are openly available in the published, peer-reviewed scientific literature.

#### References:

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