

Interactive comment on “Flux balance modelling to predict bacterial survival during pulsed activity events” by Nicholas A. Jose et al.

Anonymous Referee #1

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The authors have responded adequately to all my questions and comments. I have one last remark, though: The SBML file generates a couple of warnings when loaded with R package sybilSBML. More specifically, two reactions appear to miss an identifier (no_id) and another reaction with identifier rxn00062 occurs twice. In addition, the number of genes and metabolites displayed in R and Matlab (with iNJ1153.mat) is not the same. I hope the authors can look into these issues before publication.

Interactive comment on Biogeosciences Discuss., <https://doi.org/10.5194/bg-2017-403>, 2017.

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Discussion paper

