

Subject: Re: Your submission to bioRxiv (Bayesian Flows)
From: Michael Betancourt <betan@symplectomorphic.com>
Date: 6/27/24, 11:44 AM
To: "Wolkovich, Elizabeth" <e.wolkovich@ubc.ca>

[CAUTION: Non-UBC Email]

Two options:

1) You create an account and submit, <https://info.arxiv.org/help/submit/index.html>. Using a UBC email address would be helpful.

Submission can be done with LaTeX source (you upload the source which the server compiles it into the PDF that is posted) or the PDF itself (not as encouraged as uploading the source but may be necessary if you're using Sweve or the like). You can also upload a gzip folder of auxiliary code.

Once you submit you get a paper password that you would share with me so that I can link my account as a coauthor. This will essentially endorse the paper so it shouldn't have any trouble being uploaded.

2) You send me all of the source and I upload. I'm familiar with the process but would have to wrangle the source and code into the right formats which might take a bit.

Either way let me know!

On Jun 26, 2024, at 7:09PM, Wolkovich, Elizabeth <e.wolkovich@ubc.ca> wrote:

Hi Mike,

That sounds great. Given I have only succeeded in 1/3 tries at getting a paper through BioRxiv, I would be happy to be sponsored for arXiv. But, do whatever is easiest for you!

Thanks again,
L

From: Michael Betancourt <betan@symplectomorphic.com>
Sent: Tuesday, June 18, 2024 1:07 PM
To: Will <will.pearse@gmail.com>
Cc: Wolkovich, Elizabeth <e.wolkovich@ubc.ca>; Davies, Jonathan <j.davies@ubc.ca>
Subject: Re: Your submission to bioRxiv (Bayesian Flows)

[CAUTION: Non-UBC Email]

I should be able to put this up on the arXiv if that's of interest. Lizzie I should also be able to sponsor you if you wanted to upload it yourself.

On Jun 18, 2024, at 6:37 AM, Will <will.pearse@gmail.com> wrote:

What?! This is nuts!
I guess ecorvoRxiv?
Will
Sent in haste from a tiny screen.

On Tue, 18 Jun 2024, 11:36 Elizabeth M Wolkovich, <wolkovic@mail.ubc.ca> wrote:

Denied! I even tried claiming it was a research article with new data/results (I mean, some of the example code is very new) but somehow they still rejected me. If folks have suggestions around this, I can try to post there -- or other recommended archives (as in, you know who runs them and the long-term plans for them. i.e., they are not like FigShare -- owned by SpringerNature and have never committed to keeping the data open forever).

L

----- Forwarded Message -----

Subject: Your submission to bioRxiv
Date: Mon, 17 Jun 2024 15:51:23 +0000
From: biorexiv@cshlbp.org
To: e.wolkovich@ubc.ca, e.wolkovich@ubc.ca

[*CAUTION:* Non-UBC Email]


MS ID#: BIORXIV/2024/599096
MS TITLE: A four-step Bayesian workflow for improving ecological science

Dear E M Wolkovich;

Thank you for submitting your manuscript to bioRxiv. We regret to inform you that your manuscript is inappropriate for bioRxiv. All submissions are screened to ensure they match bioRxiv posting criteria and scope. bioRxiv is intended for complete research papers with new data/analysis and your submission does not fall into that article type.

Please be assured that this conclusion simply refers to the manuscript's appropriateness for bioRxiv and is not a judgment on the merits of the

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work described.

Thank you so much for your interest in bioRxiv.

Kind regards,
The bioRxiv team