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Wed, 07/22/2020 - 15:40 #1

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Hi.

I was wondering if it was possible to export .star file from cisTEM refinement package without generating a particle stack. My particle stack is very large and I only have 1 class in the refinement package.

Thanks,

Qianglin

Wed, 07/22/2020 - 21:11 #2

timgrant

Hi Qianglin,

Hi Qianglin,

There is currently no way to do this i'm afraid. If disk space is the issue you can always use /dev/null as the output file. This will stop the file actually being written to disk, but it will still be somewhat slow as it still does the reading / image processing.

Also, I just want to make sure that you know that cisTEM defaults to protein being black, and Relion defaults to it being white - one of the things that the export does is invert the stack, and normalise it according to the Relion requirements.

Thanks,

Tim

Wed, 07/22/2020 - 21:48 (Reply to #2)

Hi Tim,
Hi Tim,
Thank you very much for the information. It is very helpful.
Best,
Qianglin

Sat, 03/11/2023 - 16:03 #4

Daniel Asarnow

A little late, but a

A little late, but a workaround is to export particles with a 2 pixel box.

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 $\textbf{Source URL:} \underline{\textbf{https://cistem.org/export-star-file-cistem-refinement-package-without-generating-particle-stack}$