Supplementary File V: Options of cryosieve-csrefine

The program cryosieve-csrefine is designed to automatically and sequentially execute a series of operations in CryoSPARC, namely import particle stack, ab-initio, homogenous refinement or non-uniform refinement jobs.

--i

The --i option specifies the input star file(s) or txt file(s). This parameter supports multiple input files, such as --i a.star b.star c.star. Wild-cards can also be used, for example, --i output/_iter?.star will include output/_iter0.star, output/_iter1.star up to output/_iter9.star obtained in this protocol. Additionally, the input file can be txt file containing paths of star files, with each file listed on a separate line.

--directory

The --directory option specifies the directory of particles. If this option is not provided, the current directory is assumed as the default location.

--0

The $--\circ$ option is used to specify the output summary file path. When running the program, it will generate a summary report for each particle stack in CSV format. This report includes information such as resolutions and B-factors estimated by CryoSPARC.

--sym

The --sym parameter specifies the molecular symmetry.

--ref

The --sym parameter specifies the path of initial reference model. If this option is not provided, the program will utilize CryoSPARC's ab-initio job to obtain an initial model.

--repeat

The --repeat option specifies the number of trials to be performed for each particle stack. It is important to note that CryoSPARC's estimation of resolution and B-factor includes some randomness. This option can be utilized if the user wishes to reduce noise in the results.

--user, --project, --workspace and --lane

These options specify the user (e-mail address), project, workspace and lane in CryoSPARC.

--nu

If the flag **--nu** is enabled, **cryosieve-csrefine** will perform non-uniform refinement instead of homogeneous refinement.

--resplit

If the flag --resplit is enabled, cryosieve-csrefine will activates the force re-do GS split option in CryoSPARC's homogeneous or non-uniform refinement. It is crucial to be aware of the risks associated with this option. For more details, please refer to Supplementary File IV.

--workers

By default, cryosieve-csrefine refines all star files in parallel, but the option --workers allows you to limit the number of jobs executed simultaneously.