

Transferring Data to SRA

1. Transfer your dataset from DME to the `dme-sra-submit-aws` S3 bucket, as described in [Downloading to an AWS S3 Bucket via the GUI](#). (For credentials to this bucket, contact ncidatavault@mail.nih.gov.) AWS keeps this data in this bucket for two weeks.
2. In SRA (<https://submit.ncbi.nlm.nih.gov/subs/sra>) after login, click **New submission**. The bucket policy is already set for NCBI to access the dataset.
3. In the SRA **Files** tab, specify `dme-sra-submit-aws` (the name of the S3 bucket) as the source of your files.

The screenshot displays the NIH National Library of Medicine Submission Portal. At the top, the NIH logo and "National Library of Medicine" text are visible. Below this, the "Submission Portal" header is shown with a "My submissions" tab. The main section is titled "Sequence Read Archive (SRA) submission:" and includes a text input field. At the bottom, a progress bar indicates the status of the submission, with a yellow box highlighting "4 FILES".