

Studies, Clips and Results

A one-page CoroEcho 3D portal reference for contributors. Use the separate Acquisition and Export Guide before scanning.

1 Organize one study

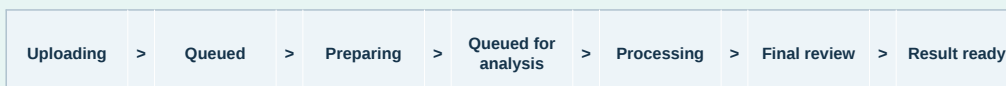
- A **study** groups one examination from one research participant.
- A **clip** is one moving RawDicom acquisition. Put complementary views in the same study.
- Give the study and every clip a distinct, non-identifying nickname.
- Institution is required. Optional participant/operator codes must never be hospital identifiers.

2 Upload the clips

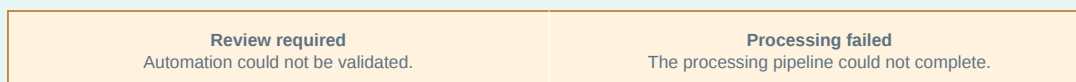
- Add every extensionless RawDicom file to its study.
- Check the view and cycle category for each clip.
- Confirm that files, filenames, and visible annotations are de-identified.
- Keep the page open until the combined transfer reaches 100% and the clip is **Queued**.

3 Follow each clip

Normal progression through My studies:



Alternative participant-visible outcomes:



4 Wait for review

- You may close the page once every clip has reached **Queued**.
- The project team reviews every processed outcome before anything is released.
- If another acquisition or upload is needed, the investigator will contact you.
- A notification email contains a secure portal link, never the result file.

5 Retrieve the outcome

- Return to **My studies**; complementary clips remain grouped by study.
- A released preview, clip, or download appears on the corresponding clip.
- **Review required** or **Processing failed** may remain visible while the team decides whether to contact you or retry.
- Experimental outputs must not guide diagnosis or patient management.

De-identified submissions only. Never place a patient name, hospital identifier, date of birth, or other direct identifier in nicknames, codes, filenames, or visible annotations.