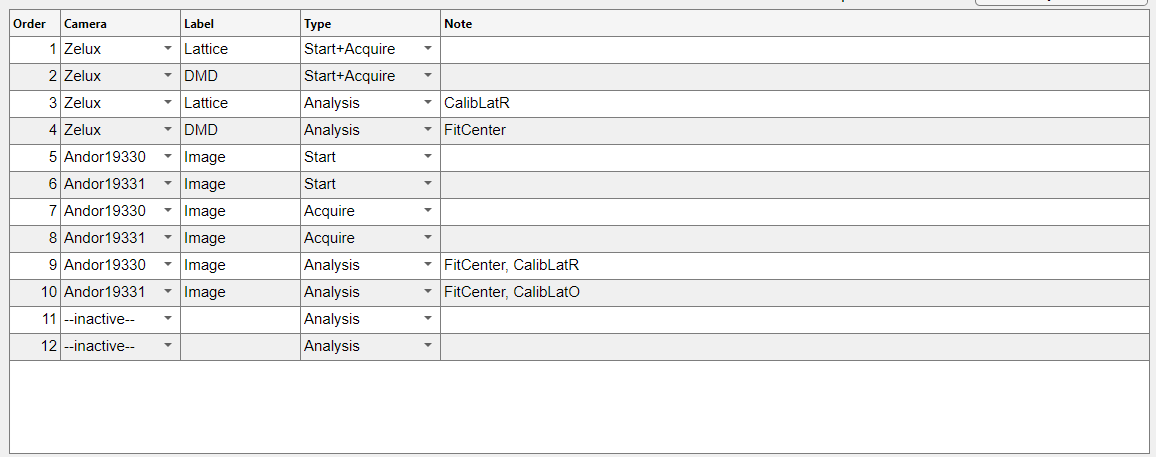
Instruction for Setting Up SequenceTable

**Basic information**

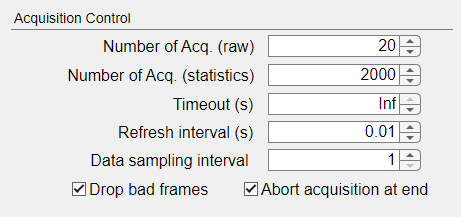
The acquisition and analysis pipeline are mostly defined as a *SequenceTable*.

An example of the table:



Additional to the SeuqenceTable, there are a few parameters to control the acquisition behaviors.

An example of the parameters:



**Configure SequenceTable**

A *step* is a single row in the SequenceTable. To define a step, one need to specify 4 varaibles:

* **Camera:** categorical variable, the camera/projector to use for this step, can be selected from cameras {Andor19330, Andor19331, Zelux} or projectors {DMD}
* **Label:** text string, the label of the image
  + The purpose is to distinguish two images taken by the same camera
  + Can be set to anything except for blank string ""
* **Type:** categorical variable, type of operations, can be set to {Start, Start+Acquire, Acquire, Analysis, Project}
* **Note:** text string, optional, additional input parameters for the step, separated by comma, leaving empty means default settings

To define **Type**:

* "Start" mode starts the acquisition
  + It either triggers the camera internally, or set the camera to be ready for external trigger
* "Acquire" mode consists of two processes: acquisition and preprocess
  + In acquisition, it acquires a raw image from specified camera with specified label
  + Immediately after acquisition it pre-processes the raw image
* "Start+Acquire" mode does both "Start" and "Acquire"
* “Analysis” mode performs a series of analysis processes
* "Project" mode only works for projectors

**Note** specifies the parameters for the step, which will be different depending on the **Type:**

* "Start", the note parameters control the behavior of starting acquisition.
  + Available parameters are defined in class Camera.startAcqusition under /core/camera/Camera.m
* "Acquire"/"Start+Acquire", because it consists of more than one process, the parameters require an identifier to specify which process they are controlling
  + For “Acquire”, available identifiers are {Acquire, Preprocess}
    - “Acquire”, available parameters are defined in class Camera.acquire under /core/camera/Camera.m
    - “Preprocess”, available parameters are defined in class Preprocessor.process under /core/preprocess/Preprocessor.m
  + For “Start+Acquire”, available identifiers are {Start, Acquire, Preprocess}
  + An example note is: “Start, verbose=1, Acquire, min\_wait=1, Preprocess, verbose=1*”*
* "Analysis", the parameters need to specify which analysis process to make as identifier and followed by the parameters
  + The available analysis processes are defined in class AnalysisRegistry under /core/analysis/AnalysisRegistry.m
* "Project", the parameters control the projected pattern

**Configure Acquisition Parameters**

* **Number of Acq. (images):** control the storage space for raw images, the raw images generated by this number of full sequence run will be kept in memory.
* **Number of Acq. (Statistics)**: control the storage space for analysis results, because the analysis usually gives only a few numbers, this number can be much larger than "Number of Acq. (images)" while still taking much smaller storage space.
* **Timeout (seconds)**: control the maximum wait time for acquiring a single image from any camera. If wait time exceeds the max but there is still no new image available, the program will throw a warning and return an empty image.
* **Refresh interval (seconds)**: control the refresh interval for acquiring a single image from any camera. The camera will check if a new image is available every this interval and return the acquired image when it sees the new data.
* **Data sampling interval**: integer, control the saving interval of data acquisition. The app will save the data every this interval. When set to 1, it saves all the data.