

## Arvados - Story #18100

### Clinical entry of samples

09/03/2021 07:48 PM - Peter Amstutz

<b>Status:</b> New	<b>Start date:</b>
<b>Priority:</b> Normal	<b>Due date:</b>
<b>Assigned To:</b>	<b>% Done:</b> 0%
<b>Category:</b> Workbench2	<b>Estimated time:</b> 0.00 hour
<b>Target version:</b>	
<b>Description</b>	
The clinical user process:	
<ol style="list-style-type: none"><li>1. Patient data is imported from Redcap ahead of time (separate batch process)</li><li>2. Clinician decides a biopsy will be sent for sequencing</li><li>3. Navigate to patient / biopsy</li><li>4. Add each biopsy sample (DNA tumor / DNA normal / RNA) with state AT_SEQUENCING</li></ol>	

### History

#1 - 09/03/2021 08:24 PM - Peter Amstutz

- Description updated

#2 - 09/03/2021 08:59 PM - Peter Amstutz

- Category set to Workbench2

#3 - 09/07/2021 02:57 PM - Peter Amstutz

- Description updated

#4 - 09/07/2021 06:32 PM - Peter Amstutz

- Description updated

#5 - 09/14/2021 04:04 PM - Peter Amstutz

- Description updated

#6 - 09/14/2021 04:06 PM - Peter Amstutz

- Description updated

#7 - 09/14/2021 04:06 PM - Peter Amstutz

- Subject changed from Clinical data entry to Clinical entry of samples