**TMT location:**

1. Log in to TMT (<http://10.39.196.170/tmt/Home.do>).
2. Select Test cases tab.
3. Expand caTissue product from the tree view.
4. Expand Mater List-v2.0 version
5. Expand Admin Component
6. Expand Collection Protocol Area
7. Select Test case ID 111 with short title UPG\_Edit\_Existing\_Coll\_Prot

**Short Title :**

Collection Protocol: UPG\_Edit\_Existing\_Coll\_Prot (111\_UPG\_Edit\_Existing\_Coll\_Prot)

**Purpose:**

Test to ensure that the existing collection protocol and its details can be edited with upgraded application.

**Pre-requisites:**

1) Deploy caTissue v1.2 with the imported oracle dump located at [\\ps6086\DatabaseDumps2\caTissue\Oracle\_v12](file:///\\ps6086\DatabaseDumps2\caTissue\Oracle_v12).

2) Once the application is up and running upgrade this to caTissue v2.0 and re-start the server.

**Procedure:**

1) Login as admin@admin.com (Login123).

2) Navigate to Administrative Data >> Collection Protocol >> Edit page (Refer the expected output)

*2) Collection Protocol page should be displayed with* ***Collection Protocol Search*** *with the attribute as “Title”; Condition Starts with and Value “”*

3) Enter “**Site**” in the Value text box and click on Search (Refer the expected output)

*3)Edit Collection Protocol page should be displayed with* **“***User can not edit existing Events and Specimen Requirements except for the Label Format” message in red color. The page should display the Collection Protocol Details in Tree Format to the left and Collection Protocol Details for Siteman CP to the right..The details should be as follows*

|  |  |
| --- | --- |
| **Attribute** | **Value** |
| *Principal Investigator* | *Admin,Admin* |
| *Protocol coordinators* |  |
| *Title* | *Siteman CP* |
| *Short Title* | *Siteman CP* |
| *IRB ID* |  |
| *Start Date* | *08-26-2010* |
| *Consent Waived* | *No* |
| *Number of Participants Anticipated* | *0* |
| *Description URL* |  |
| *Activity Status* | *Active* |
| *Clinical Diagnosis* |  |
| *Store all aliquot(s) in same container?* | *No* |

*Collection Protocol Details to the left:*

*Siteman CP*

*Event Point: 0.0 pre*

*Specimen Details: Single Parent Specimen with two derivatives. Each derivative with 10 aliquots each.*

4) Edit the Collection Protocol Details as follows and then click on Save Collection protocol button (Refer the expected output)

|  |  |
| --- | --- |
| ***Attribute*** | ***Value*** |
| *Principal Investigator* | *Test,Scientist* |
| *Protocol coordinators* | *Admin,Admin* |
| *Title* | *Siteman CP* |
| *Short Title* | *Site\_CP* |
| *IRB ID* | *09809\_00902* |
| *Start Date* | *08-26-2010* |
| *Consent Waived* | *No* |
| *Number of Participants Anticipated* | *1200* |
| *Description URL* |  |
| *Activity Status* | *Active* |
| *Clinical Diagnosis* | *Nodular melanoma, Amelanotic melanoma and Superficial spreading melanoma* |
| *Store all aliquot(s) in same container?* | *Yes* |

*4) “Collection Protocol successfully updated.” Message should be displayed and the CP should be saved with the edited details*

5) Click on **Consent tab** from the Collection Protocol Details page on right hand side (Refer the expected Output)

*5) The* ***Consent Tab*** *should be selected with no defined consent tiers added.*

6) Enter the following in  Unsigned Form URL : [www.myconsent.com](http://www.myconsent.com). Click on Add More and enter the following consent tier in the text box one by one

i) Can we publish the PHI related information? .

ii) Allowed to distribute the specimen?

iii) Allowed to contact for future research?

Once all the consents are added click on Save Collection Protocol Button (Refer the expected output)

6) *“Collection Protocol successfully updated.” Message should be displayed and the CP should be saved with the newly added consent. The page should navigate back to Collection Protocol details page.*

7) Click on **Privilege tab**.(Refer the expected Output)

7)*Privilege tab>>Privilege Details should by default show “Siteman Cancer”All Default Users and All Default Privileges selected.*

8) Click on LHS Collection Protocol Details Tree View and click on **0.0 pre event point.** (Refer the expected output)

*8) The RHS of the page should display the “****Event Details****”with the following details*

|  |  |
| --- | --- |
| ***Attribute*** | ***Value*** |
| *Study Calendar Event Point* | *0.0* |
| *Collection Point Label* | *Pre* |
| *Clinical Diagnosis* | *Not Specified* |
| *Clinical Status* | *Not Specified* |

9) Click on parent specimen Specimen\_E1\_S0 and click on Add More for Derive Specimen(s). Select “Molecular Class” ,”DNA type”, ”Virtual Location” ,Quantity 10 µg and 0.02 concentration. Under the “Aliquot(s)” section enter the aliquot count as 4 Quantity per aliquot as 0.1 and from the storage location select “Auto”.

Click on Save Specimen Requirements (Refer the expected Output)

9) Event Details page should be displayed

10) Click on Save Collection Protocol Button. (Refer the expected Output )

*10) “Collection Protocol successfully updated.” Message should be displayed. The page should navigate back to Collection Protocol details page.*

**Verification Logic:**

1. Navigate to Collection Protocol--🡪Edit page. Search for the created collection protocol with short title ***Siteman\_CP***
2. Once the collection protocol opens in edit mode.Verify the CP details for the protocol.
3. Verify details such as study calendar event point, clinical diagnosis, and clinical status are saved correctly. (The details should be as per the event details table)
4. Verify all the specimen requirement details such as Specimen Class, Specimen type, Pathological Status are saved correctly. (The details
5. Verify that all the respective audit events are captured appropriately.