

# Adding Data Findings to a Characterization

To add findings to a characterization, you can add data directly to Data and Conditions by creating columns and adding data manually or you can import UTF-8 .csv (comma-separated value) files. Information can include laboratory conditions, pH, or temperature. You can add as many files as you wish.

1. Add data values to Data and Conditions.
  - a. To import a file of data values:
    - i. Organize your data so that each column name is unique. The following table provides an example.

| column_type:condition  | condition   | datum               |
|------------------------|-------------|---------------------|
| column_name:Experiment | Formulation | (other):Size by DLS |
| value_type:            | score       | mean                |
| value_unit:            |             | nm                  |
| constant_value:0       |             |                     |
| 1                      | 5           | 2                   |
| 1                      | 10          | 4                   |
| 1                      | 15          | 6                   |
| 2                      | 5           | 8                   |
| 2                      | 10          | 10                  |
| 2                      | 15          | 12                  |

To add a value that is not currently in the system, such as a column name, insert "(other):" before the value.

The "value\_type:", "value\_unit:", and "constant\_value:" rows are optional. In those rows, an empty cell is acceptable.

- ii. Save the spreadsheet of data values to a UTF-8 csv file (not just csv).
- iii. Click **Import csv** and follow the prompts to add the data file to the Findings Info.



- iv. The columns and data are added to Data and Conditions.
- b. To add the data values manually:
  - i. Specify the number of **columns** and **rows** for the matrix, and click **Update**.

Finding Info

Data and Conditions 2 columns 3 rows **Update**

- ii. Add the data values to the rows.

**i** Whether you imported or added information manually, you can preface each data value with one of the following: Maintain the default, equal to (=), greater than (>), less than (<), or approximate (~).

**Constant Value**

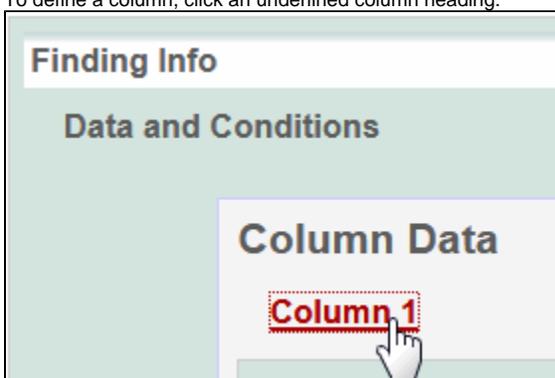
*For boolean column value type, please enter 1 for true, 0 for false*

**Remove** **Save** **Cancel**

|   |   |   |               |
|---|---|---|---------------|
| <b>&gt;</b> <input type="text" value="23.0"/> | <b>&gt;</b> <input type="text" value="23.0"/> | <b>&gt;</b> <input type="text" value="23.0"/> | <b>Delete</b> |
| <b>~</b> <input type="text" value="12.0"/>    | <b>~</b> <input type="text" value="12.0"/>    | <b>=</b> <input type="text" value="12.0"/>    | <b>Delete</b> |

**Files** **Add**

- To define a column, click an underlined column heading.



The Column Definition panel appears.

- Select a **Column Type**, Datum or Condition.
- Select a **Column Name** or select **other** and add a new one.

**i** Column Notes

You can add up to three cell viability Column Names, including **cell viability**, **cell viability B**, and **cell viability C**. You can further identify the column with the Column Value Type.

- For Column Type, **Datum**, the following characterization(s) display customized **Column Name** options.

| Characterization Type | Column Type and Column Name Option(s)   |
|-----------------------|---|
| Physico-Chemical      | <ul style="list-style-type: none"> <li>• <b>Molecular</b> – Molecular Weight</li> <li>• <b>Purity</b> – % purity for sample</li> <li>• <b>Relaxivity</b> – R1, R2, T1, T2</li> <li>• <b>Size</b> – PD1, Peak N, RMS size, Z Average</li> <li>• <b>Surface</b> – charge, zeta potential</li> </ul> |
| In Vitro              | <b>Enzyme Induction</b> – % of Control  |
| In Vivo               | Click <b>Other</b> to name the column yourself.   |

- For Column Type, **Condition**, all characterizations provide the **Column Name** options in the left column of the following table. The Column Name auto-populates the **Condition Property** options in the right column.

| Column Type, Condition Auto-populates<br>Column Name | Column Name Auto-populates<br>Condition Property   |
|--|--|
| Centrifugation                                       | N/A  |
| Culture Media  | media type, serum percentage   |
| Electromagnetic Radiation                            | bandwidth, frequency, time, wavelength   |
| Freeze Thaw  | N/A  |
| Long Term Storage                                    | lyophilized, time  |
| Lyophilization                                       | time   |
| pH   | N/A  |
| Sample concentration                                 | N/A  |
| Short Term Storage                                   | lyophilized, time  |
| Solvent Media  | ion concentration, ionic strength, molecular formula, osmolality, serum percentage, with serum |
| Sonication   | number of pulses, pulse duration   |
| Temperature  | N/A  |

5. To further identify a column, select a **Column Value Type**.



Once the column information is saved, the Column Type is shown in parentheses after the Column Name, such as **cell viability (mean)**.

6. Select a **Column Value Unit**, or select **other** and add one.

7. If you want the same value to fill all rows in a column, add a **Constant Value**.



**For Column Value Type, boolean**

For Column Value Type, **Boolean**, enter a Constant Value of 1 for true and 0 for false.

8. Click **Save**, and the column(s) are updated.



If needed, click **Set Column Order** to change the order of the column headings in the matrix.

9. Click **Save** in the Finding section.