

Annotation report

The Annotation report provides a table summarizing the cell-level attributes of a single cell counts data node.

To run Annotation report:

- Click a Single cell counts data node
- Click the **Annotation/Metadata** section of the toolbox
- Click **Annotation report**

The task will run and generate a task report.

The Annotation report includes two tables (Figure 1) - the top table summarizes the categorical attributes, giving the number of levels of each attribute, and the bottom table summarizes the numeric attributes, providing some basic summary statistics about the distribution of each attribute.

Categorical attributes

Attribute ↕	# categories ↕
Cell symbol	2218
Graph-based	9
Rows per page	25 ▾
⏪ ⏴ (1 of 1) ⏵ ⏩	
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Numeric attributes

Attribute ↕	Min ↕	Max ↕	Mean ↕	Median ↕	Q1 ↕	Q3 ↕	Missing ↕	Zeroes ↕
Expressed genes	13.00	6,487.00	1,623.48	1,448.00	1,035.50	2,008.50	0	0
Mitochondrial reads percent	0	99.70	10.04	7.69	4.07	10.65	0	1.00
Total count	500.00	56,877.50	5,751.38	4,650.00	3,130.00	7,190.25	0	0
Rows per page	25 ▾	⏪ ⏴ (1 of 1) ⏵ ⏩						Download

Figure 2. Annotation report tables for categorical and numeric attributes

To download a text-file version of one of the tables, click **Download** in lower right-hand corner of the table.

Additional Assistance

If you need additional assistance, please visit [our support page](#) to submit a help ticket or find phone numbers for regional support.



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