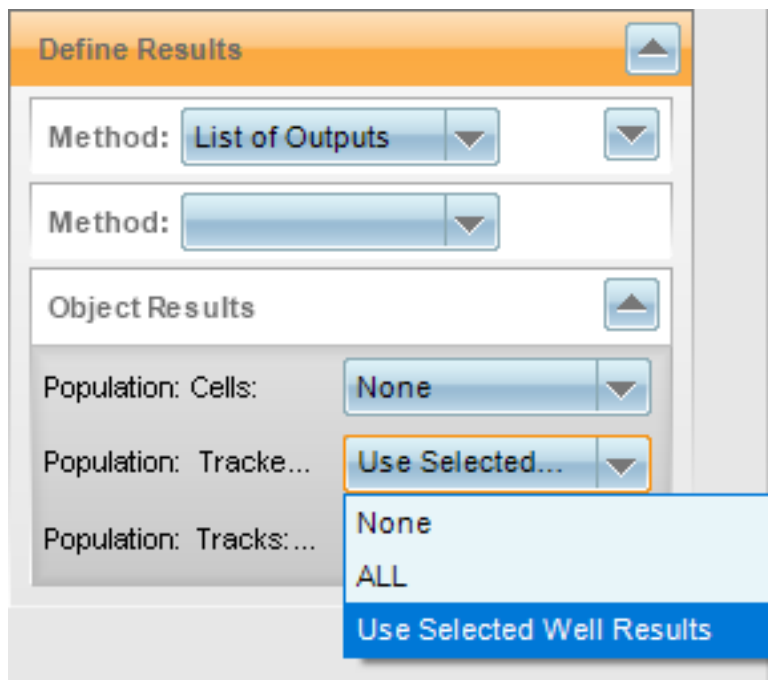




1, 在Define results的最后有一个object results的选项, 设置好分析程序后, 如果需要导出单细胞结果的txt结果, 则需要在此进行选择。可以选择不同的population, 然后下拉菜单中选择all则输出所有计算过的结果, 选择use selected well results则输出和孔数据有相关性的结果。请优先选择use selected well results以避免大量冗余数据量。

### Object Results

你也可以通过点击help搜索 object results看到相关解释

The **Object Results** section allows you to save also **results** per **object** to the database. This can be configured individually for each population (local or global). After the evaluation they are available for inspection on the **Evaluation** screen and/or can be exported for use in statistical programs via **Settings – Export Data**. See also section [“Export Data”](#).

#### First level parameter

##### Population

- **None:** **Object results** are not saved in the database to save disk space (default setting).
- **ALL:** All **object** properties which are explicitly calculated by building blocks (e.g. *Calculate Intensity Properties*, *Calculate Morphology Properties*) are saved in the database. I.e. all columns of the **object** tables displayed by the **Define Results** building block are saved.

##### Notice for “ALL”

This option generates large and space-consuming tables in the database. Only choose this option if you really need it.

- **Use Selected Well Results:** Only those cell properties which have corresponding well **results** defined are saved in the database. E.g. if the “mean nuclear area” is defined as a well result, the “nuclear area” is also saved for each individual cell.

##### Notice for “Use Selected Well Results”:

- Properties referenced in formula outputs are not saved in the database.
- The **object** table displayed by the **Define Results** building block shows all properties (not only the selected ones).

**Graph**

Population: Well Results

Readout Parameter: Well Results

X-Axis: Index

Logarithmic: ☐

Plot-Type: Bars

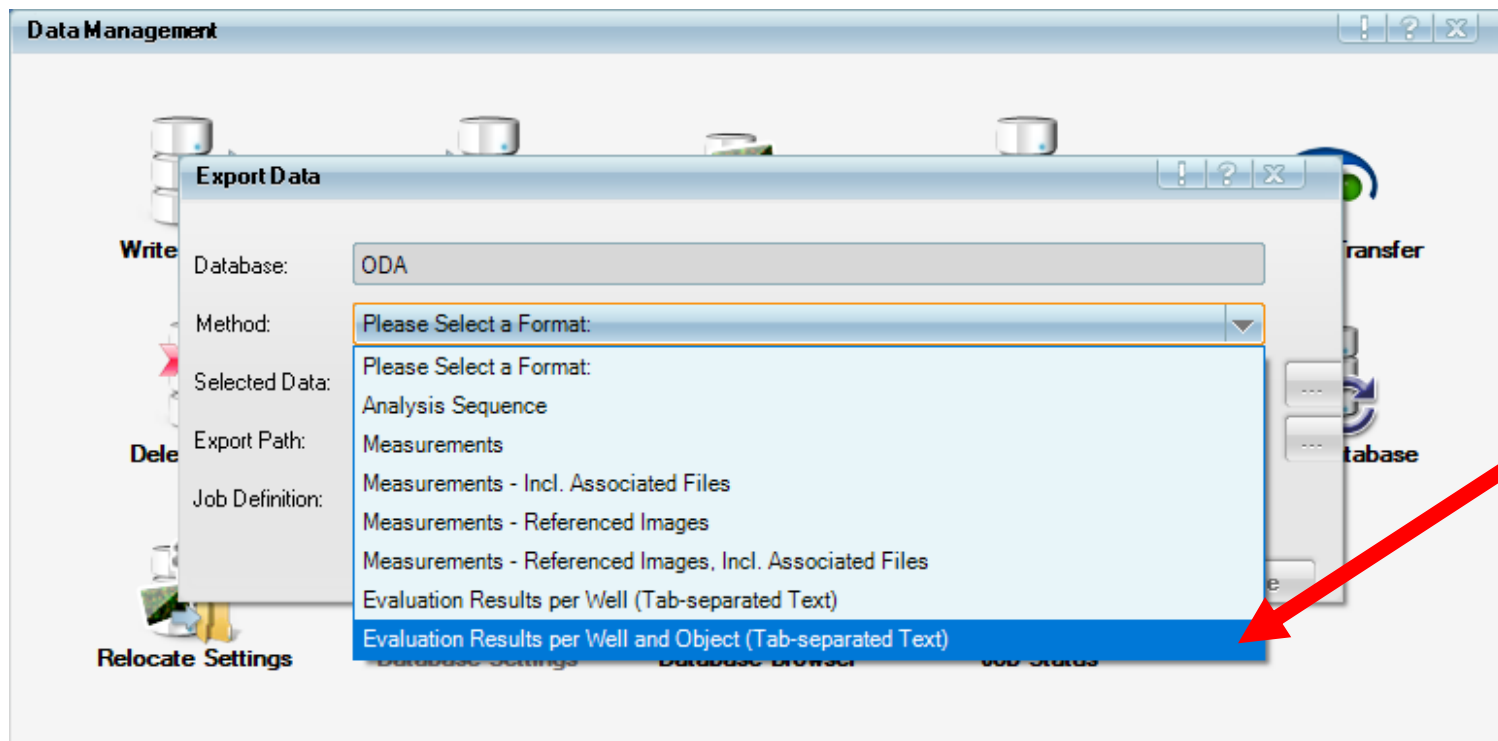
Grouping: none

Grouping Style: One Graph

Data Points: 9 / 9 displayed

Display all

2, 按照步骤1设置好 define results, 选择好需要输出的孔数据和单细胞数据后, 右边选择想要计算的孔和视野, 到 evaluation处点击start, 计算完毕后可以在graph的 population中选择孔数据或单细胞数据的结果进行查看。



3, 数据导出的时候, setting-data management-export data, 此时选择最后一个选项, 则可以导出单细胞数据和孔数据的txt文件, 该txt可以拷贝到个人电脑后右键open with Excel。