MPAPASS Software Collection

In 1 collection

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DISCLAIMER

This protocol summarizes key steps for a specific type of method, which is one of a collection of methods and assays used for EV analysis in the NCI Translational Nanobiology Section at the time of submission of this protocol. Appropriate use of this protocol requires careful, cohesive integration with other methods for EV production, isolation, and characterization.

ABSTRACT

This is collection contains the protocols required for each step in the mpapass software pipeline for performing stitched multiplex analysis. This is one of a number of protocols in the pipeline for using the mpapass software package and is applicable to the latest release of the software.

BEFORE START

MPAPASS software can be found at https://nano.ccr.cancer.gov/mpapass.

BEFORE START INSTRUCTIONS

MPAPASS software can be found at https://nano.ccr.cancer.gov/mpapass.

Protocol status: Working

We use this protocol and it’s working

Created: Oct 06, 2020
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### FILES

**Protocol**

- **NAME**: MPAPASS - Gating flow cytometry multiplex data
- **VERSION**: 1
- **CREATED BY**: Jennifer Jones

**Protocol**

- **NAME**: MPAPASS - Creating an MPAPASS database
- **VERSION**: 1
- **CREATED BY**: Jennifer Jones